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Helen E. Scharfman  
Paul S. Buckmaster *Editors*

# Issues in Clinical Epileptology: A View from the Bench

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# Issues in Clinical Epileptology: A View from the Bench

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## Foreword

This book is a tribute to Phil Schwartzkroin, who, in addition to over four decades of committed original research into fundamental mechanisms of epilepsy, has been a consummate editor, contributing to archival knowledge as well as the synthesis of new information. Phil served as editor-in-chief of *Epilepsia*, the journal of the International League against Epilepsy (ILAE), editor of the *Encyclopedia of Basic Epilepsy Research* [1], and editor of definitive textbooks on animal models of epilepsy [2, 3], brain development and epilepsy [4, 5], and brain plasticity and epilepsy [6]. This book is a fitting acknowledgment of Phil Schwartzkroin's career achievements, as an edited volume that addresses many of the most pressing research issues concerning neuronal mechanisms underlying epilepsy that were, and continue to be, his passion.

Epilepsy is among the most common serious neurological diseases. According to a study by the World Health Organization, epilepsy accounts for 1% of the global burden of disease [7]. This is equivalent to breast cancer in women and lung cancer in men. Among primary disorders of the brain, epilepsy ranks with depression and other affective disorders, Alzheimer's disease and other dementias, and substance abuse [7]. Public attention on epilepsy, however, and the resultant amount of resources devoted to research on epilepsy, is but a small fraction of that for these other medical conditions. The fact that epilepsy has been a stigmatized disease in most cultures since antiquity might be one reason why it has remained in the shadows, but interest in epilepsy also suffers because it is a complicated multifactorial condition with such diverse manifestations that clinical research alone to elucidate comprehensive underlying fundamental neuronal mechanisms is essentially not possible.

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### Historical Perspective

Epilepsy is an ancient disease, being both common and easy to recognize in antiquity. There is a long history about epilepsy being attributed to many prevailing causes, but it was not until the late nineteenth and early twentieth century that modern concepts of epilepsy as a disease of the brain attributed to excessive neuronal activity was first formulated. Before then, both the clinical phenomenology and some of the brain pathology associated with

epilepsy were described, but the advent of electrophysiological recordings from the human brain by Berger and colleagues initiated a new way of studying the disease [8]. It quickly became apparent that during seizures there was a dramatic change in the electroencephalogram (EEG) and that in many patients there was also an alteration in the EEG even between seizures when they were behaviorally “normal” [9, 10]. The interictal EEG could show “spikes” (fast, sharp transients) or spike and slow wave discharges. As the field of basic neurophysiology began to develop around that time, investigators were able to demonstrate, predominantly in animal models of seizures, that some neurons in the cortex fired abnormally during seizures and also during the interictal EEG spikes. From the early 1950s through the early 1970s, single unit studies predominantly were carried out in animal models of either acutely provoked seizures in neocortex [11–13] or hippocampus [14, 15]. Although most of our current concepts about the origin and spread of seizures were developed from this work, opportunities to investigate functional mechanisms at the cellular and subcellular level were limited.

In the early 1970s, remarkable methodological advances occurred in the capacity to understand fundamental physiological and pharmacological properties of mammalian brain function: the development of the brain slice and dissociated cell culture. At that time, Per Andersen in Oslo devised the ability to maintain a slice of mammalian hippocampus in a dish for many hours and to record from the cells with extracellular and intracellular microelectrodes [16]. Within a year or two, Phil was working in Anderson’s laboratory to extend his studies on the mechanisms responsible for epileptic seizures and along the way to investigate many other important physiological functions of mammalian cortical neurons. He brought this preparation back to Stanford to collaborate with David Prince and others, and eventually in his own laboratory he continued to make significant contributions to our understanding of mechanisms underlying the development and spread of epileptic seizures.

Over the ensuing four decades, much has been learned about the electrophysiological substrates of epileptic seizure activity from studies with simplified slice preparations and from additional studies utilizing cell cultures of mammalian neocortex and hippocampus. However, two main conceptual problems persisted: the recognition that seizures artificially provoked in an *in vitro* preparation, or even those induced in a normal animal brain, are not epilepsy, and the mechanisms by which a normal brain can become chronically epileptic were not understood.

A person with epilepsy has an enduring epileptogenic abnormality responsible for the generation of spontaneous seizures, which continues to be present during the interictal state [17]. Although models of chronic focal epilepsy were created in rats and primates in the 1950s and 1960s with topical application of metals such as cobalt, iron, and alumina [2], scars were produced by these metals and made microelectrode recordings at the site of application – the area of most interest – difficult. In the 1970s, kindling became a popular animal model to study mechanisms of chronic epilepsy at the cellular level [18]; however, kindling is a model of secondary epileptogenesis and, as usually performed, kindling results in stimulation-induced seizures, not spontaneously generated seizures. Intensive investigations into mechanisms

of chronic epilepsy were facilitated by the introduction of the status epilepticus models of mesial temporal lobe epilepsy (MTLE) with hippocampal sclerosis, the most common, and most pharmacoresistant, form of human epilepsy [19]. Status epilepticus induced by kainic acid, pilocarpine, or electrical stimulation causes a pattern of hippocampal cell loss and neuronal reorganization resembling human hippocampal sclerosis, and eventual spontaneous limbic seizures [20–22]. Opportunities for invasive studies of MTLE in the epilepsy surgery setting made parallel reiterative multidisciplinary animal/human investigations possible [23]. Epileptogenesis, the process by which a normal brain is converted to one that is capable of generating spontaneous seizures, is of increasing interest to epileptologists and, as yet, can only be pursued in animal models. It is now understood that these enduring changes occur in many brain networks, not just in the areas in which the seizures appear to originate, but also in areas into which seizures propagate, and even in more remote brain regions [24].

Epilepsy is a diverse disease, and the extent to which research results obtained from animal models of a few types of epilepsy, such as MTLE, apply to other types of epilepsy remains to be determined. Future research will require the use of a wide variety of animal models of human epilepsy, such as post-traumatic epilepsy, febrile convulsions, neonatal hypoxia, infantile spasms, and genetically engineered models of genetic epilepsies and genetic diseases associated with epilepsy, such as tuberous sclerosis [25, 26]. Creation and validation of experimental animal models of the diverse forms of human epilepsy are now a high priority in order to search for targets not only for antiseizure interventions, but also for antiepileptogenic interventions that can prevent or cure epilepsy [25, 26]. Phil continued to contribute importantly to resolving many of these questions in his later work. The following are brief discussions of the questions he chose for this volume, intended to stimulate the field of epilepsy research today:

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## **The Role of Animal Models**

The ideal approach to the study of human epilepsy is to investigate patients with epilepsy; however, ethical concerns, technical constraints, and cost dictate that most of the critical questions concerning fundamental neuronal mechanisms of epilepsy still need to be resolved with experimental animal models. Although attempts are being made to create models of entire epilepsy syndromes and diseases, epilepsy can also be broken down into its component parts, e.g., epileptogenesis, ictogenesis, seizure maintenance, seizure termination, postictal disturbances, and interictal disturbances, each of which may be modeled individually [27].

*Is there more to learn about human epilepsy by studying acute seizures in animals?* Acute seizures in a normal brain are not the same as chronic epilepsy. The pathophysiological and anatomical substrates of the enduring epileptogenic abnormalities underlying different types of human epilepsy need to be elucidated. Does this mean that we have already learned all we can from studying acute seizures?



*Do people with acquired epilepsy have a genetic susceptibility?* Although an increasing number of epilepsy genes are being identified as responsible for single-gene epilepsy conditions, these diseases are rare [28]. The genetic bases of inherited diseases *associated with* epilepsy, such as tuberous sclerosis, are being elucidated. Both of these directions provide opportunities to create animal models of specific epilepsy syndromes and diseases, using genetic engineering. More importantly, however, it is now apparent that most genetic epilepsies, formally referred to as idiopathic epilepsies, can result from multiple different genetic mutations, and these represent susceptibility genes rather than epilepsy genes. The distinction between genetic and acquired epilepsies is not absolute, just as the distinction in the 1989 classification of the epilepsies between idiopathic and symptomatic disorders is a false dichotomy [29]. It is likely that some acquired disturbances are necessary for the manifestation of epilepsies primarily due to genetic abnormalities, and that genetic predispositions, susceptibility genes, influence the manifestation of epilepsies with acquired etiologies. Consequently, just as realization that acute seizures induced into a normal animal brain is not the same as epilepsy caused by an enduring epileptogenic abnormality was a paradigm shift, it must now also be realized that artificial introduction of an enduring epileptogenic abnormality into a normal animal brain is not the same as introduction of this abnormality into a brain genetically predisposed to generate specific types of epileptic abnormalities. In order to create more appropriate animal models of human epilepsy, more information is needed regarding specific susceptibility genes.

*How relevant are animal models to human epileptic phenomena and how can they be validated?* There are many different types of human epilepsy [30], and it is unreasonable to assume that any animal model will completely reproduce all aspects of a human epilepsy disease or syndrome. Rather, models will likely reproduce component parts of human epilepsies, and studies need to be designed to take advantage of the likely similarities while accounting for the differences between any given animal model and the type of human epilepsy that is being modeled. For the rare epilepsies caused by a single gene mutation, these mutations can be introduced into animals to investigate the pathophysiological consequences of their abnormal protein products, even if the phenotype does not resemble the human condition. Reiterative patient/animal investigations utilize clinical data to identify relevant questions, make use of relevant animal models to pursue investigations that are not ethically or financially feasible in patients, and then validate results in the clinical population. This has been a valuable paradigm, particularly where invasive EEG recordings can be carried out in an epilepsy surgery setting and tissue is then available for analysis.

*Comorbidity:* Patients with epilepsy have a high incidence of comorbid conditions that can contribute significantly to disability. Many disturbances, such as depression, anxiety, attention deficit hyperactivity disorder, and autism, have a bidirectional relationship with epilepsy, suggesting shared mechanisms [31]. Because these conditions can precede epilepsy, it is not clear which condition is the comorbid one. Animal models of these conditions in association with epilepsy are necessary to begin investigations into fundamental neuronal mechanisms of epilepsy comorbidity.

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## Epileptic Activity

Questions persist concerning how to model and investigate the various types of epileptiform activity encountered in patients. In this book, experts in the field address several of the most pressing questions:

*Human focal epilepsy is not focal; how can studies in animal models recreate the epileptic network necessary for the manifestation of human epilepsy?* There probably is no such thing as a single discretely localized epileptic focus in chronic human epilepsy. Epilepsy manifests as a result of disturbances in distributed networks. The ILAE states: “*Focal epileptic seizures are conceptualized as originating within networks limited to one hemisphere. They may be discretely localized or more widely distributed. Focal seizures may originate in subcortical structures. For each seizure type, ictal onset is consistent from one seizure to another, with preferential propagation patterns that can involve the contralateral hemisphere. In some cases, however, there is more than one network, and more than one seizure type, but each individual seizure type has a consistent site of onset*” [30].

*What is generalized epilepsy?* The distinction between generalized and focal epilepsies in the 1989 ILAE classification of the epilepsies is a false dichotomy [28]. No epilepsy condition, or epileptic seizure, is truly generalized. The ILAE states: “*Generalized epileptic seizures are conceptualized as originating at some point within, and rapidly engaging, bilaterally distributed networks. Such bilateral networks can include cortical and subcortical structures, but do not necessarily include the entire cortex. Although individual seizure onsets can appear localized, location and lateralization are not consistent from one seizure to another. Generalized seizures can be asymmetric*” [30].

*What are interictal EEG spikes and what is their significance?* Some interictal EEG spikes may represent exactly the same underlying neuronal mechanisms as an ictal event, as, for instance, is the case with typical absence seizures; the so-called interictal events are too brief to be associated with obvious clinical behavior. In this situation, even with focal seizures, careful investigations can demonstrate behavioral disturbances during the so-called interictal spike [32]. Similarly, generalized paroxysmal fast activity (GPFA) without behavioral correlates, which can be seen in some patients with severe epilepsy, most likely represents the same underlying mechanisms as some low-voltage fast ictal discharges. These, therefore, are fragments of seizures and the terms “interictal spike” or “interictal GPFA” would be oxymorons. There are, however, different types of interictal spikes and not all represent fragments of ictal events. Some may, in fact, reflect seizure-suppressing mechanisms [33].

*What are the limitations of studying epileptic phenomena in slice preparations?* Epileptic seizures are defined clinically as behavioral events with an electrographic correlate [17]. Electrographic changes that occur in the slice preparation, therefore, cannot be called epileptic seizures, although they may reproduce certain neuronal events similar to those which would underlie behavioral seizures in the intact animal. Disturbances related to ictogenesis at

the molecular, cellular, and perhaps microcircuit levels can be studied in slice preparation; however, disconnections from important influences of distant brain areas make it difficult to draw definitive conclusions concerning seizure generation at the level of whole-brain networks.

*How is epilepsy mediated by non-neuronal influences?* Not all epileptogenic, or homeostatic, mechanisms involve neurons. Glia play an important role in modulating neuronal activity, and other non-neuronal influences, such as hormonal changes, inflammatory and immune-mediated processes, and external toxic substances need to be considered.

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## Synaptic Plasticity

Concepts of epileptogenesis derived from an understanding of the development of human MTLE with hippocampal sclerosis, and reproduced in the animal laboratory, indicate that an initial epileptogenic insult causes cell loss, which is then followed by synaptic reorganization of surviving neuronal elements. Aberrant excitatory and inhibitory connections ultimately lead to epileptiform hypersynchronization. Epileptogenesis can occur in experimental animals, however, in the absence of obvious cell loss or synaptic reorganization, for instance with classical amygdala kindling, and epilepsy also occurs in patients who have no evidence of cell loss. Cell loss and synaptic reorganization may not, therefore, be a universal mechanism essential for epileptogenesis.

*Changes during epileptogenesis can be protective:* Neuronal plasticity occurring in response to injury can be responsible for the development of epilepsy, but homeostatic plastic changes also occur, resulting in protective seizure-suppressing influences. Investigations to identify pathophysiologic disturbances following an epileptogenic insult must clearly distinguish epileptogenic from homeostatic protective processes. These homeostatic changes could also be responsible for the appearance of interictal behavioral disturbances.

*What is the significance of cell death in acquired epileptogenesis?* Cell death is clearly not necessary for all forms of epilepsy, but when it occurs, it can be a cause of the epilepsy, or an affect of epileptic seizures.

*Inhibition is not necessarily decreased in human epilepsy, and increased inhibition may be necessary for hypersynchronization:* The old concept that epilepsy is due to an increase in excitation and a decrease in inhibition is clearly an oversimplification. In MTLE with hippocampal sclerosis and animal models of this condition, there is an increase in inhibition as well as in excitation [27]. Whereas some increased inhibition may have a protective effect, inhibition is also necessary for hypersynchronization, which is a component of most epileptic seizures. It is the types and location of aberrant excitatory and inhibitory synaptic reorganization that determine the epileptogenic process.

*Features of epilepsy in the pediatric population differ considerably from those in patients with more mature brains:* Synaptic plasticity leading to epileptogenesis is different in the developing brain than in the mature brain, and epileptic seizures can alter the synaptic plasticity necessary for normal brain development [4].

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*Research on patients with epilepsy is revealing increasing numbers of genetic aberrations, as well as disturbances in important protein products such as ion channels and neurotransmitter receptors; how do these defects explain the development and maintenance of epileptic phenomena?* Plastic changes underlying epileptogenesis involve alterations in expression of genes whose protein products are ion channels, neurotransmitter receptors, and other membrane and intracellular structures that determine excitability. The location of these changes on the cell, and their influence on neuronal interconnections, also determine propensity for hypersynchronization. Although characterization of epileptogenic disturbances at the molecular and cellular levels do not reveal how epilepsy arises at the systems level, this research can help to identify novel targets for antiseizure and antiepileptogenic drugs designed to prevent and cure epilepsy, as well as control ictal events.

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## Conclusions

The enduring legacy of Phil Schwartzkroin is impossible to summarize here, but reflected well by the discussions in this volume, written by his colleagues, who have watched his contributions evolve over time. These discussions show how complex epilepsy is, that there is much to do to resolve the questions that are associated with epilepsy, and the approaches that have allowed us to make the most advances; using the best neuroscience and clinical epileptology together – an approach Phil mastered, and would want us all to continue.

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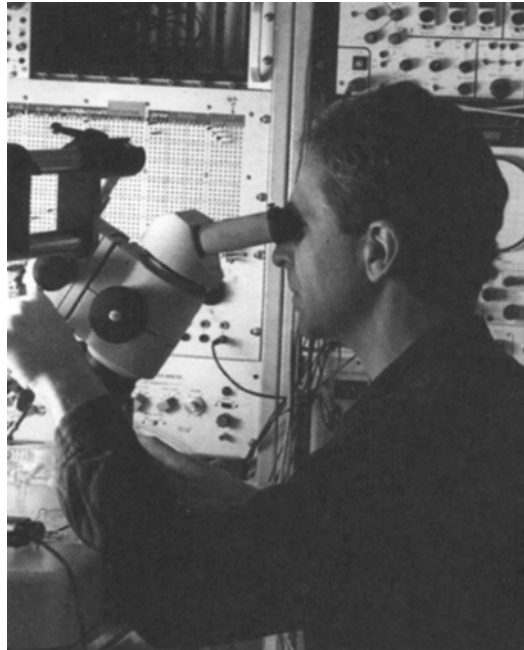
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## Preface



Philip A. Schwartzkroin recently retired after an outstanding, influential career in neuroscience and epilepsy research. He influenced the work of many neuroscientists either by the techniques he developed or his pioneering discoveries of neuronal mechanisms underlying excitability and microcircuitry in health and disease. He personally touched the lives of all those with whom he collaborated and mentored. This volume is dedicated to Phil by many of the numerous colleagues and trainees who respect him both personally and scientifically. As one might expect from Phil's 'hands-on' approach to his work, Phil had a direct influence editing the volume and designing its unusual format, in which key questions in epilepsy research are addressed from both basic science and clinical perspectives. Phil's goal for this volume was to allow experts in the field the opportunity to address critical questions in ways that would stimulate a broad readership. Instead of leaving readers with the sense that they have all the answers, his goal was to encourage them to think about how to address the important questions in epilepsy research today.



Phil began neuroscience research in high school. Through a summer fellowship from the National Science Foundation he investigated mouse behavior at the Jackson Laboratories, where he continued working during several subsequent summer breaks from college. Phil was a National Merit Scholar and graduated *magna cum laude* with highest honors in Psychology from Harvard University. As an undergraduate, Phil investigated neocortical sensory processing in the laboratory of Charles Gross. Even in this early stage of his career Phil was exceptional, publishing articles as first author while still in high school and college.

After graduating from Harvard, Phil moved to Stanford University where he earned a Ph.D. in Neurological Sciences, working in the laboratory of Kao Liang Chow, who had trained with Karl Lashley. Chow served as a role model for Phil, because he was a basic scientist in a clinical department and focused on understanding the relationship between brain structure and function in health and disease. In the coming years, Phil would also serve as a role model in much the same way – for Phil’s own trainees.

Phil’s dissertation addressed the effects of vestibular stimulation on single cells in cat visual cortex and superior colliculus. Following his dissertation, Phil started to address questions related to epilepsy, which became the major research focus of his career. He started as an Epilepsy Foundation trainee with David Prince for one year before becoming a postdoctoral fellow in Per Anderson’s laboratory at the University of Oslo in Norway. Phil returned to the Department of Neurology at Stanford one year later and brought with him a brain slice recording chamber, which had recently been developed. The experiments Phil conducted using brain slices, put him at the forefront of controversy, because there was skepticism that brain slices would be a useful experimental preparation. Nevertheless, he persevered and played a critical role in the ultimate acceptance of the approach. He also was a pioneer; he was the first in the USA (after Yamamoto in Japan) to develop the slice preparation for intracellular recording. He demonstrated the utility of brain slices for studying normal synaptic transmission and synaptic plasticity – and was the first to demonstrate long-term potentiation in the slice preparation. He showed how brain slices could be used to study epileptiform activity, paving the way for decades of epilepsy research based on the slice preparation.

In 1975, Phil was appointed Assistant Professor of Neurology at Stanford, where he was first in the world to carry out intracellular studies in slices of surgically resected human epileptic neocortex and hippocampus. He began what would become a standard structural and functional approach in his laboratory, and characterized numerous cell types in the hippocampus with correlative cellular electrophysiology and intracellular staining techniques. His initial studies began with CA1 pyramidal cells and were followed by some of the most difficult recordings at that time, of GABAergic interneurons.

Phil moved to the Department of Neurological Surgery at the University of Washington in 1978, where he continued to use brain slices to make fundamental discoveries in hippocampal anatomy and physiology. For example, he pioneered the use of the slice preparation for studying brain development. With that approach, he was first to demonstrate depolarizing IPSPs in immature hippocampus. He also was at the forefront of the most challenging electrophysiological techniques, such as the use of simultaneous intracellular

recording from two monosynaptically-coupled neurons. Phil was also a leader in applying the slice preparation to questions related to animal models of epilepsy. His laboratory was one of the first to characterize, using both morphology and electrophysiology, transgenic mouse models of epilepsy. Some of this work, such as the studies of the Kv1.1 knockout mouse, were major advances in epilepsy research. In addition, Phil addressed other areas of epilepsy research, including cortical dysplasia. He was an early contributor to studies on the basic mechanisms of the ketogenic diet, and was first to demonstrate that furosemide, a chloride co-transporter antagonist, was anti-epileptic. During this time Phil's productivity was exceptional. For example, over a one-year period in 1988 he was senior author of 14 research articles, six of which appeared in *The Journal of Neuroscience*. Phil earned many awards, including fellowships from the Guggenheim and Klingenstein Foundations, two Jacob Javits Awards from the NIH, and he was one of the first recipients of the American Epilepsy Society/Milken Family Medical Foundation Research Award. In 2001, Phil moved to the Department of Neurological Surgery at the University of California at Davis where he held the Bronte Endowed Chair in Epilepsy Research. He became an emeritus professor in 2013.

In addition to his outstanding contributions to research, Phil was a dedicated member of the epilepsy research community. He served on NIH study sections and on scientific advisory boards for the Epilepsy Foundation and Citizens United for Research in Epilepsy. He led some of the first efforts to address translation, organizing seven workshops and six books that brought together basic and clinical epilepsy researchers. These workshops, and the books that resulted from them, remain some of the most influential in the field. He served as one of the first chairs of Investigators' Workshops for the American Epilepsy Society and was the first basic scientist president of the American Epilepsy Society. He chaired the International League Against Epilepsy (ILAE) Commission on Neurobiology and organized an ILAE Workshop on the Neurobiology of Epilepsy. After these accomplishments, Phil served as co-editor-in-chief of *Epilepsia*, where he strengthened the impact and reputation of the journal.

In addition to his achievements in research and service to the epilepsy research community, Phil trained many students and postdoctoral fellows. Many of these individuals ultimately became independent neuroscientists themselves, including numerous leaders in epilepsy research today. Phil provided his trainees with a great degree of independence. But when help was needed, he was an efficient, "hands-on" trouble-shooter who quickly solved technical problems. Phil trained largely by example. He demonstrated a strong work ethic and began days in the lab at an extremely early hour. During meetings in his office, Phil demonstrated impressive collegiality with colleagues, both near and far, often phoning them in the middle of conversations if he wanted to address a question. He could pick up the phone and call almost anyone in the field, often the original source of information on a given topic. Phil's trainees benefited greatly from exposure to some of these investigators when they visited the laboratory or attended meetings organized by Phil. Despite his considerable accomplishments, Phil was modest and easy to work with, characteristics that helped shape the laboratory environment into one that was truly enjoyable. Writing was an area where Phil's training method was more direct, but just as constructive. Phil routinely transformed

manuscripts – often long-hand – with extensive editorial remarks that illustrated how to clearly convey ideas. Remarkably, Phil could do so rapidly and effectively, which left trainees wondering if they could ever master scientific writing and editing as well. In addition, he made it clear that excellent scientific writing was extremely important.

On May 3–5, 2013 a workshop entitled “Issues in Clinical Epileptology: A View from the Bench” was held in honor of Phil. The workshop was supported by several organizations, including the American Epilepsy Society and CURE. It was not possible for all of Phil’s colleagues and trainees to attend, but the group that was able to come considered it an excellent meeting – as well as a great opportunity to honor Phil (see review by C. Stafstrom, *Epilepsy Curr.*, 2013). In considering the type of book that would complement this ‘festschrift,’ Phil provided a great deal of input, as mentioned above. His contribution to this volume shows that – despite his retirement – his influence will be present for years to come.



Issues in Clinical Epileptology: A view from the Bench. A Festschrift in Honor of Philip Schwartzkroin. Pajaro Dunes Resort, Watsonville, California. May 3–5, 2013. First row: Paul Buckmaster, Jong Rho, Jurgen Wenzel, Phil Schwartzkroin, Gerry Chase, Helen Scharfman, Laura Reece, Jean-Claude Lacaille, Mike Haglund, Scott Baraban. Second row: Mareike Wenzel, Catherine Woolley, Carol Robbins, Alan Mueller, Dennis Kunkel, Dennis Turner. Third row: Elsa Rosignol, Daryl Hochman, Robert Fisher. Fourth row: Sloka Iyengar, Jerome (Pete) Engel Jr., James Trimmer, Carl Stafstrom, Damir Janigro, Robert Hunt. Fifth row: Aristeia Galanopoulou, Tracy Dixon-Salazar, Solomon (Nico) Moshé, David Prince, Massimo Avoli, Jeffrey Noebels, Robert Wong, Michael Gutnick, Leena Knight. Back row: Satoshi Fujita, Aylin Reid, Charles Behr, Ben Strowbridge, Robert Berman.

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**Part I**

**Seizures, Epileptiform Activities,  
and Regional Localization**

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# How Can We Identify Ictal and Interictal Abnormal Activity?

1

Robert S. Fisher, Helen E. Scharfman,  
and Marco deCurtis

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## Abstract

The International League Against Epilepsy (ILAE) defined a seizure as “a transient occurrence of signs and/or symptoms due to abnormal excessive or synchronous neuronal activity in the brain.” This definition has been used since the era of Hughlings Jackson, and does not take into account subsequent advances made in epilepsy and neuroscience research. The clinical diagnosis of a seizure is empirical, based upon constellations of certain signs and symptoms, while simultaneously ruling out a list of potential imitators of seizures. Seizures should be delimited in time, but the borders of ictal (during a seizure), interictal (between seizures) and postictal (after a seizure) often are indistinct. EEG recording is potentially very helpful for confirmation, classification and localization. About a half-dozen common EEG patterns are encountered during seizures. Clinicians rely on researchers to answer such questions as why seizures start, spread and stop, whether seizures involve increased synchrony, the extent to which extra-cortical structures are involved, and how to identify the seizure network and at what points interventions are likely to be helpful. Basic scientists have different challenges in use of the word ‘seizure,’ such as distinguishing seizures from normal behavior, which would seem easy but can be very difficult because some rodents have EEG activity during

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normal behavior that resembles spike-wave discharge or bursts of rhythmic spiking. It is also important to define when a seizure begins and stops so that seizures can be quantified accurately for pre-clinical studies. When asking what causes seizures, the transition to a seizure and differentiating the pre-ictal, ictal and post-ictal state is also important because what occurs before a seizure could be causal and may warrant further investigation for that reason. These and other issues are discussed by three epilepsy researchers with clinical and basic science expertise.

### Keywords

Convulsion • Convulsive • Electroencephalogram • Epilepsy • Epileptic • Focal seizure • Epileptiform • Seizure-like • Spike-wave discharge • Theta • Sharp wave • Behavioral arrest • Interictal spike • Ictal • Pre-ictal • Transition to seizure

## 1.1 Introduction

Seizures are common and important neurological symptoms that may require treatment. Seizures can signal underlying disease. In addition, many research laboratories study mechanisms of seizures. Therefore, a commonly accepted definition of “seizure” is needed for both clinical and research purposes. Some events may obviously be seizures, but others might comprise imitators of seizures [62], epileptiform non-seizure events, or variants of normal laboratory animal behavior.

### 1.1.1 Clinical Perspective

#### 1.1.1.1 Definition of a Seizure

Webster says that a definition should capture the “essence” of an entity. What then is the essence of a seizure? Table 1.1 highlights definitions from various authorities, dating back to Johns Hughlings Jackson in 1870 [58].

Terms that recur in the various definitions include excessive, disorderly discharge, synchronous, self-limited, abnormal, paroxysmal, neurons, central nervous system (CNS) and cortex. Corresponding symptoms are listed as alteration or loss of consciousness, involuntary movements, sensory, psychic or autonomic disturbances and other clinical manifestations. These terms cover a lot of territory. Delineating

the possible clinical manifestations of seizures is beyond the scope of this chapter, but an overview may be found in [73]. In 2005, a task force of the International League Against Epilepsy [37] provided a parsimonious definition of a seizure as “a transient occurrence of signs and symptoms due to abnormal or synchronous neuronal activity in the brain.”

In clinical practice, a clinician rarely sees the abnormal electrical discharge, with the exception of successful video-EEG monitoring, so this discharge is inferred on the basis of a typical constellation of clinical symptoms. Application of the definition also requires ruling out other conditions. For example, abnormal and synchronous firing of thalamic neurons in a patient with Parkinson’s disease [17] represents a transient symptom correlated to tremor, but it is not a seizure. Therefore, a definition of seizures must include an implied qualifier: “and not due to other known conditions producing a similar picture.”

Some writers use the modifying term “epileptic seizures” to distinguish them from common usage of terms such as heart seizures, psychogenic seizures or other non-epileptic paroxysmal events. However, not all seizures imply epilepsy, particularly for single seizures with low likelihood of recurrence or for provoked seizures. Hence, the phrase “epileptic seizures” tends to be either misleading or redundant.

**Table 1.1** Prior Definitions of Seizure

References	Definitions – Note that several say “epilepsy” in place of “seizure”
Jackson [58]	Epilepsy is a symptom... an occasional, an excessive and a disorderly discharge of nerve tissue (in the highest centers)
Penfield and Jasper [75]	An epileptic seizure is a state produced by an abnormal excessive neural discharge within the central nervous system
Aird et al. [3]	Epilepsy may be defined as a paroxysmal disturbance of central nervous system (CNS) function, which is recurrent, stereotyped in character, and associated with excessive neuronal discharge that is synchronous and self-limited
Engel [34]	Epileptic seizures are the clinical manifestations (symptoms and signs) of excessive and/or hypersynchronous, usually self-limited, abnormal activity of neurons in the cerebral cortex... An epileptic seizure may consist of impaired higher mental function or altered consciousness, involuntary movements or cessation of movement, sensory or psychic experiences, or autonomic disturbances
Hauser and Hesdorffer [53]	A seizure can be defined as a paroxysmal disorder of the central nervous system characterized by abnormal cerebral neuronal discharge with or without loss of consciousness
Hopkins et al. [55]	An epileptic seizure is a clinical manifestation presumed to result from an abnormal and excessive discharge of a set of neurons in the brain. The clinical manifestation consists of sudden and transitory abnormal phenomena, which may include alterations of consciousness, motor, sensory, autonomic, or psychic events, perceived by the patient or an observer
Adams et al. [2]	Epilepsy may be defined as an intermittent derangement of the nervous system due presumably to a sudden, excessive, disorderly discharge of cerebral neurons

The seizure definition of excessive neuronal discharges derived from Hughlings Jackson’s time, is 144 years old, when awareness of brain electrical activity was new. This mindset has led generations of clinicians and researchers to think of a seizure as an electrical disorder. Abnormal electrical discharges are just one manifestation of seizures, not necessarily more important than metabolic, blood flow, receptor, gene activation, network connectivity and many other changes that are intrinsic to seizures. A contemporary definition of seizures would likely be less electrocentric and focus more on excessive and sustained activation of specific brain networks. The research community should be challenged to invent a better definition for seizures.

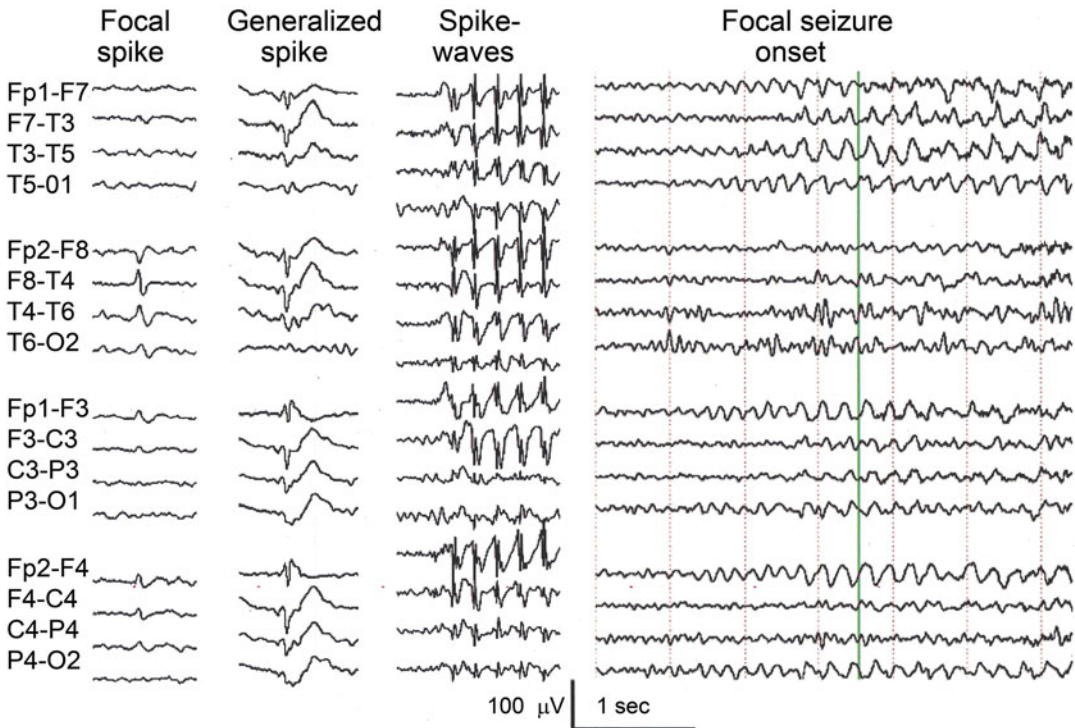
### 1.1.1.2 EEG Manifestations of Seizures

Clinicians rely heavily on electroencephalographic patterns to identify, classify, quantify and localize seizures [7]. Figure 1.1 illustrates common epileptiform EEG patterns. The term epileptiform is used to connote EEG patterns believed to be associated with a relatively high risk for having seizures. Gloor [43] defined spikes as

potentials that stand above the background, have a “pointy” shape, duration between 30 and 70–80 ms, asymmetric rise and fall, and followed by a slow wave. The potential should have a sensible field, meaning that it should be reflected in physically adjacent electrodes and perhaps in synaptically linked regions such as the contralateral hemisphere. “Sharp waves” have durations of 70–200 ms. The distinction between spikes and sharp waves is arbitrary in the clinical arena and is discussed further below (see also [28]).

Spikes may be focal or apparently generalized across widespread regions of brain bilaterally. Rhythmic recurrence of spikes followed by slow waves is referred to as spike-waves. Focal spikes tend to be associated with focal seizures with or without secondary generalization. In contrast, generalized spikes tend to be associated with seizures that are nonfocal at their onset. Generalized spike-waves are associated with absence (previously called petit mal) seizures.

The right panel of Fig. 1.1 illustrates the onset of a focal seizure in the top four channels, which are in the left temporal region. The local rhythm can be seen evolving in amplitude,

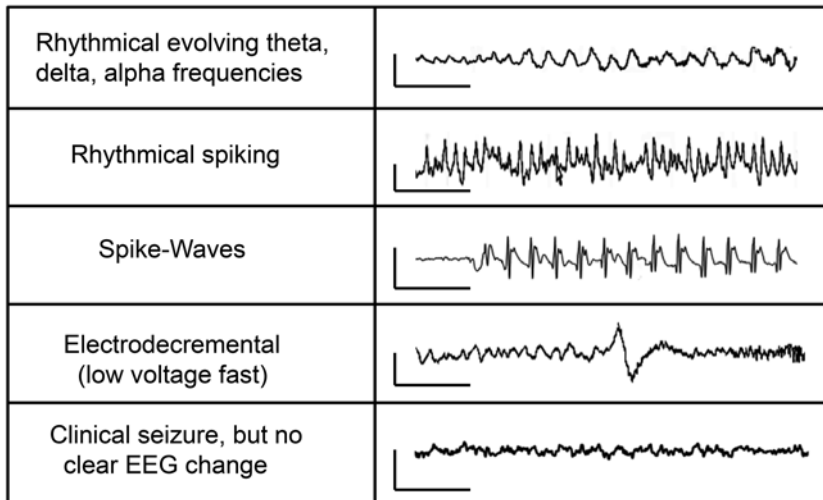


**Fig. 1.1** Common epileptiform EEG patterns. Common patterns are shown for individuals with focal spikes, generalized spikes, spike-waves, and a seizure with focal onset (From Fisher, unpublished)

frequency and degree of sharpness. Other channels also reflect the seizure activity, but it is best formed and earliest in the top four channels. Where the potential becomes sharp, there is a phase reversal (down in one channel and up in the next channel) between the top and the second from the top channel. Polarity conventions of the EEG indicate that the electrode common to both these channels is the site of maximum negativity compared to neighbors on either side. Active (discharging) seizure foci are extracellularly negative, since positive ions flow from the extracellular space into the neuron during excitation. Therefore, the phase reversal of a spike or seizure onset can be used to approximately localize the region of seizure origin.

The EEG recorded from the human scalp at the start of the seizure can take at least five different forms, as illustrated in Fig. 1.2. One pattern is rhythmically evolving frequencies in the theta (4–7/s), delta (0–3/s) or alpha (8–12/s) bands.

The rhythmical activity can have varying degrees of sharpness, but spikes and sharp waves are not required to be part of the rhythmical pattern of a focal seizure. An evolution of frequency and amplitude over time is needed to distinguish a seizure from many other normal and abnormal rhythmical events encountered in the EEG. The second pattern of seizure origin is rhythmical spiking. This may be most commonly seen with seizures in hippocampus and neighboring structures. Spike-wave patterns typically occur during generalized absence seizures, but presence of spike-waves cannot be equated with absence epilepsy. Spike-waves also can appear focally during focal seizures or during the course of generalized tonic-clonic seizures. Neocortical seizures often manifest with an electrodecremental pattern, referring to a general flattening of brain rhythms at the start of a seizure. Electrodecremental patterns are commonly seen with tonic, atonic and sometimes tonic-clonic



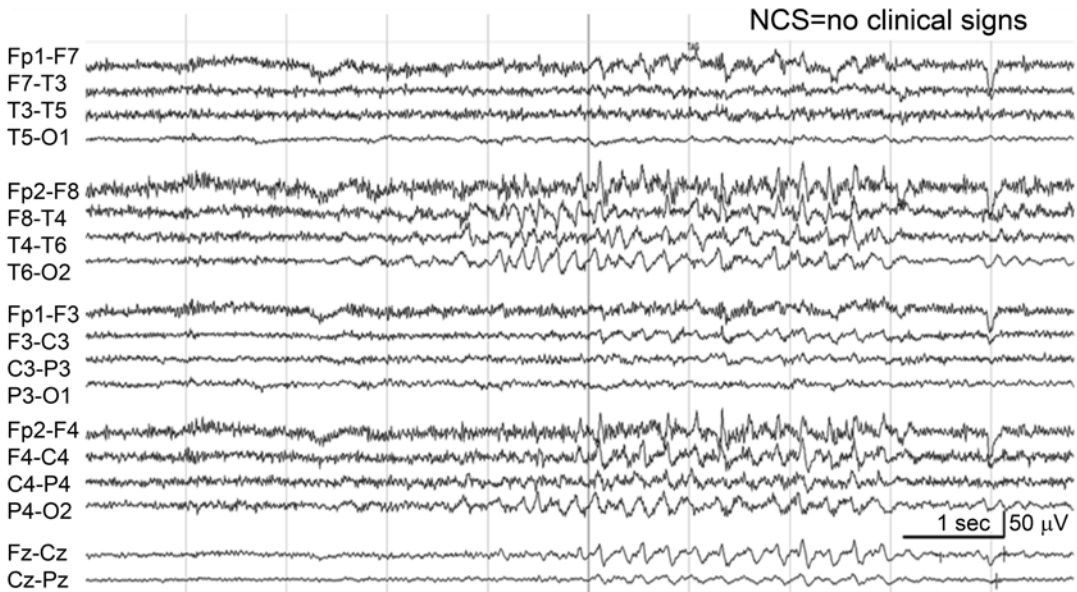
**Fig. 1.2** Common EEG patterns at the start of seizures in patients with epilepsy (From Fisher, unpublished)

seizures [35]. The apparent disappearance of EEG activity is a consequence of the typical 1–70 Hz bandpass filter used to review EEG. In fact, a very low frequency potential heralds the start of such seizures [57, 92] but is largely filtered out by the low frequency filters commonly utilized during scalp EEG revision. Careful examination of the electrodecremental region shows presence of low voltage, high frequency activity [29, 38]. Considerable study has demonstrated importance of frequencies in the beta (13–30 Hz), gamma range (30–100 Hz), ripple (100–250 Hz) and fast ripple (250–1000 Hz) range. Activity in the fast ripple or higher ranges is sometimes referred to as high-frequency oscillations (HFO's) [32, 98, 100]. HFO's can be useful markers for the region of seizure onset. Epilepsy surgery is more successful when regions generating high frequencies are resected [41]. The fifth electrographic pattern of a seizure onset is no change in the scalp EEG. The presumption here is one of sampling error. Two-thirds of cortex is enfolded in sulci and dipole discharges in sulci do not always project to scalp EEG electrodes. Seizures can originate in mesial temporal, orbitofrontal or inter-hemispheric regions far from scalp electrodes. Negative EEG findings therefore do not rule out underlying focal seizures. The EEG

must be correlated with the clinical picture. Of note here is that seizures that begin in the brainstem in experimental animals often lead to convulsions before the forebrain EEG shows any change from normal [42] (personal observations, HES).

### 1.1.1.3 Ambiguities in EEG Manifestations of Seizures

Electroencephalographers sometimes disagree about whether a particular pattern is epileptiform and representative of associated seizures. Figure 1.3 shows an evolving event over the right mid-temporal region lasting for about 5 s. The EEG technician noted no clinical signs. Such events might be considered too brief to represent a seizure: duration of at least 10 s has occasionally been applied operationally [1], but there is no official minimum time to define a seizure. In animal research, 2–3 s is often used as a minimum time for an electrographic seizure but the length of time that is sufficient to define a seizure is extremely variable [26, 31]. However, discharges accompanied by clinical seizures qualify as electrographic seizures regardless of their duration. In the extreme, a single generalized spike associated with a myoclonic jerk could be considered to be a very brief seizure.



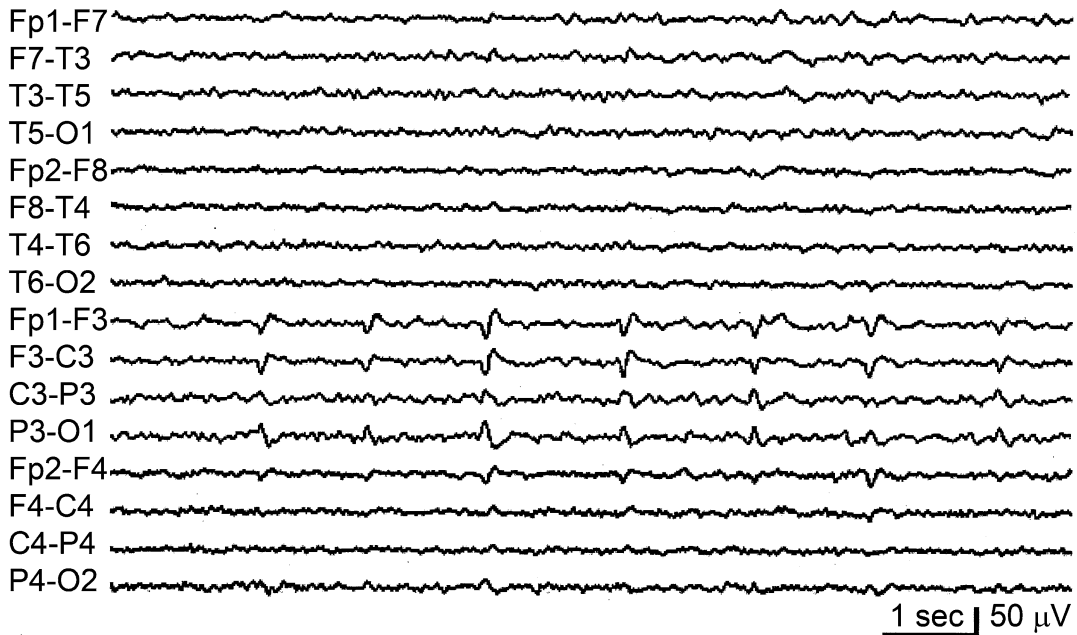
**Fig. 1.3** Is this a seizure? Rhythmic brief epileptiform activity, illustrating the ambiguity involved in deciding whether an EEG event corresponds to interictal activity or a seizure (From Fisher, unpublished)

Epileptiform EEG activity has been categorized as ictal, meaning during a seizure, postictal, meaning after a seizure and interictal, meaning between seizures. While ingrained in common usage, these terms may be more confusing than helpful [36]. What sense does it make to designate an interictal spike in cases where there have not been two seizures? Where does the behavioral and EEG pattern of an ictal event merge into the postictal behavioral confusion and EEG slowing? Is postictal slowing always a consequence of the seizure [33]? Delineations between ictal and postictal may not be obvious. Are periodic lateralized epileptiform discharges (PLEDs, Fig. 1.4) interictal, ictal or either depending upon circumstances [76]? When is a burst of generalized spike-waves interictal and when is it ictal? Behavioral manifestations, such as unresponsiveness and automatisms, tend to occur in direct proportion to the duration of spike-wave discharges [77]. Whether a person is noted to have clinical signs such as limited responsiveness depends upon how carefully they are tested. Meticulous studies [4] show that responsive latency and task accuracy declines even during a period of so-called interictal spikes. Research in animals

suggests the same is true for rodents [54], although the assumptions in these studies – that blocking interictal spikes improves behavior and therefore interictal spikes cause behavioral impairment – may not be true. Instead, blocking interictal spikes may only be helpful because of a reduction of other brain abnormalities, not necessarily the spikes *per se*. Clinically, interictal spikes tend to correspond to the zone of origin of a seizure, but not always. Figure 1.5 illustrates interictal spikes from the right temporal region and electrographic seizure onset from the left temporal region in the same patient.

#### 1.1.1.4 Clinical Conclusions

The commonly employed definition of a seizure as a transient occurrence of signs and symptoms due to abnormal or synchronous neuronal activity in the brain is almost a century and a half old, and it does not capture the essential nature of seizures as depicted by modern neuroscience. Seizures are diagnosed clinically, taking into account numerous entities that can imitate seizures, such as syncope, transient ischemic attacks, sleep disorders, confusional migraine, tremor, dystonia, fluctuating delirium and psychological episodes. The scalp EEG is



**Fig. 1.4** Periodic lateralized epileptiform discharges (PLEDs) – are they ictal or interictal? PLEDs over the left central (C3) region are shown. Some electroencephalog-

raphers consider this pattern to be interictal and others ictal, while still others believe it depends upon particular circumstances (From Fisher, unpublished)

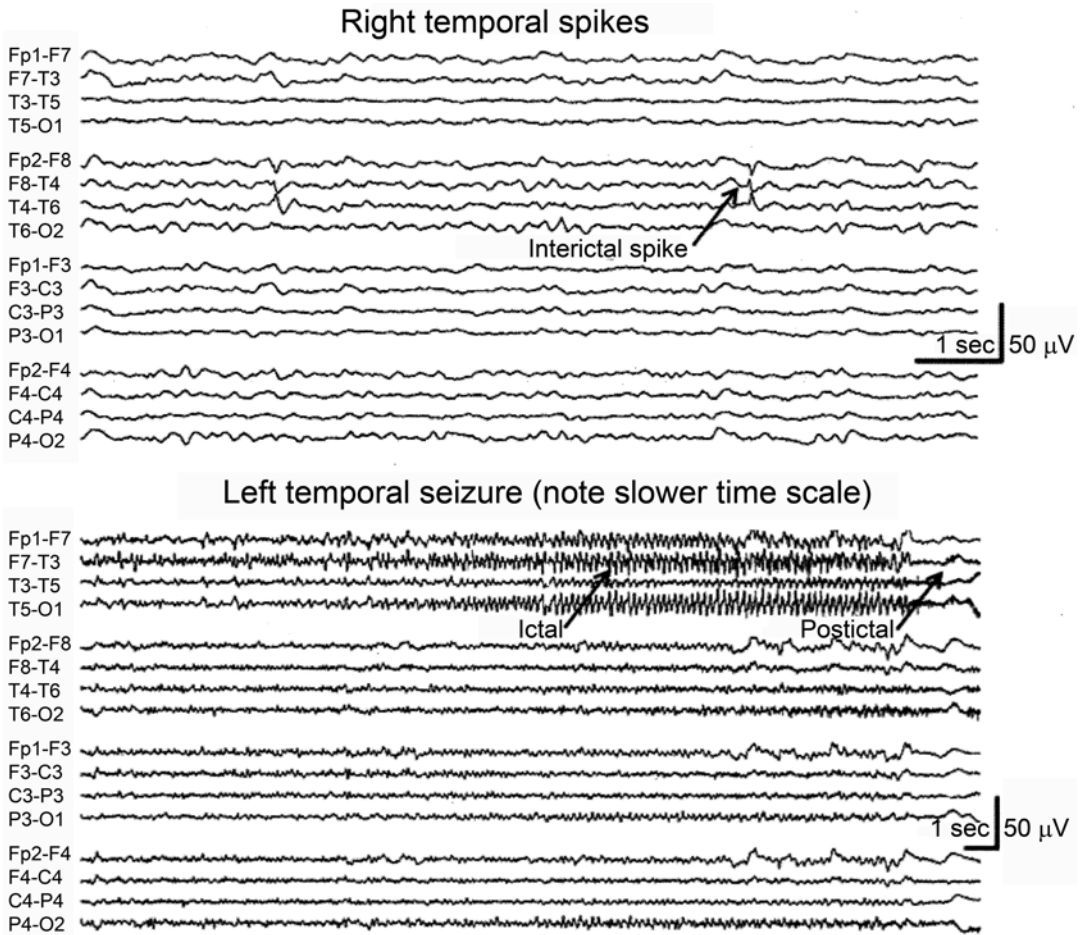
a helpful adjunct to diagnosis of seizure disorders, but it is not clear that an EEG pattern should be intrinsic to a definition of seizures. There is no unifying form; instead at least five different EEG patterns can accompany seizures. EEG correlates of high risk for seizures are categorized as ictal (during a seizure), postictal (after seizure) or interictal (between seizures). These distinctions often are unclear and arbitrary, in that the interictal-ictal boundaries are blurred for many seizures. Even so-called interictal spikes can affect behavior.

We need a better understanding of what constitutes the pathophysiological and behavioral essence of a seizure. Numerous questions arise for basic researchers. Need a seizure always involve an excessive discharge and increased synchrony? Have neurons been given excessive primacy in seizures over glia? Do seizures emerge only in cortex or can they develop in subcortical structures as well? Does it make sense to talk about where seizures start, given the involvement of widespread networks? What brain networks are involved in seizures of different types and

which behaviors correlate with seizures in these networks? These questions will only be answered with a collaboration between basic researchers and clinicians.

## 1.2 Defining Seizure Correlates with Intracranial Electrodes in Patients

The advent of intracranial recordings (with grid and strip electrode arrays) and intracerebral recordings (with depth electrodes) during pre-surgical evaluation in patients with partial epilepsies resistant to pharmacological treatment changed our view of the electrographic correlate of a seizure. During pre-surgical intracranial monitoring, seizures are recorded with electrodes positioned close to the generators of ictal epileptiform discharges. In particular, depth stereo-EEG electrode implants aim at the epileptogenic area. This is done by accurately planning electrode insertion on the basis of the analysis of the

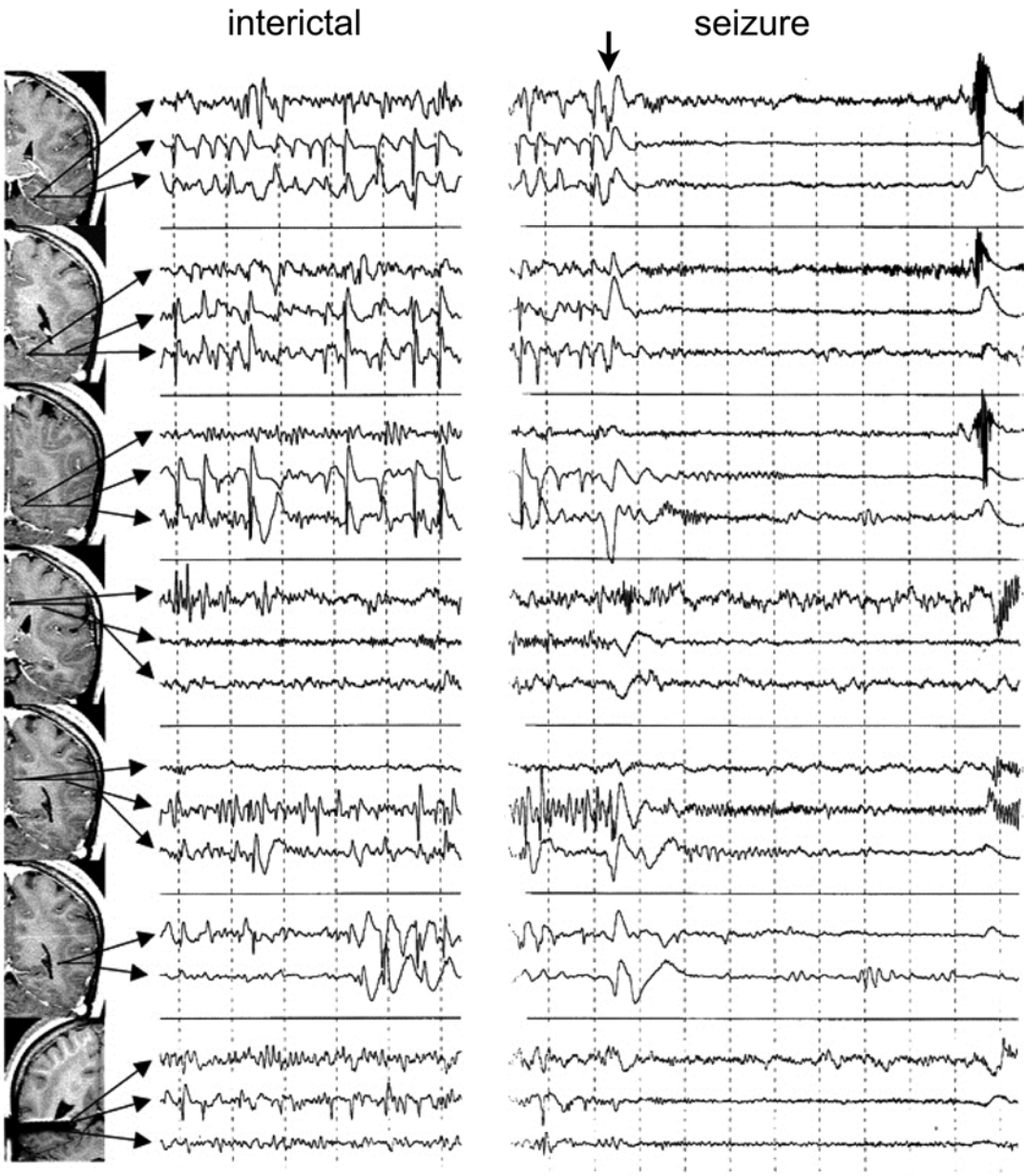


**Fig. 1.5** Interictal-ictal disparity with spikes in the right hemisphere and seizures on the left. Interictal-ictal disparity in the same patient as Fig. 1.5, with interictal spikes

over the right temporal region, but seizure onset from the left temporal region. Note different time scales for each segment (From Fisher, unpublished)

sequence of localizing clinical features observed during seizures recorded by video monitoring with scalp EEG performed as part of the pre-surgical examination [23, 85]. Intracerebral recordings are finalized to identify the cortical networks activated during a seizure that should be surgically removed to cure the patient. The areas involved in seizure generation are defined as the seizure-onset zone and the epileptogenic zone, which includes the regions of onset and propagation of the ictal epileptiform discharge. Intracranial recordings contribute to outline a larger area, defined as irritative zone, that generates abnormal interictal events/potentials, but is not directly recruited during a seizure discharges.

A large number of pre-surgical studies focused on the functional interactions between the epileptogenic and the irritative zones have been reported in the last 20 years. These studies demonstrate that (i) the irritative area is not coincident and it is usually larger than the epileptogenic/seizure onset zone, (ii) interictal discharges do not show a coherent relationship with seizure discharges, in terms of location and activation patterns, (iii) the rate of interictal discharges can either increase or decrease just ahead of a seizure and (iv) in most cases the electrographic pattern of seizure onset is completely different from the activity recorded during interictal discharges (for review see [28, 29]; Fig. 1.6).



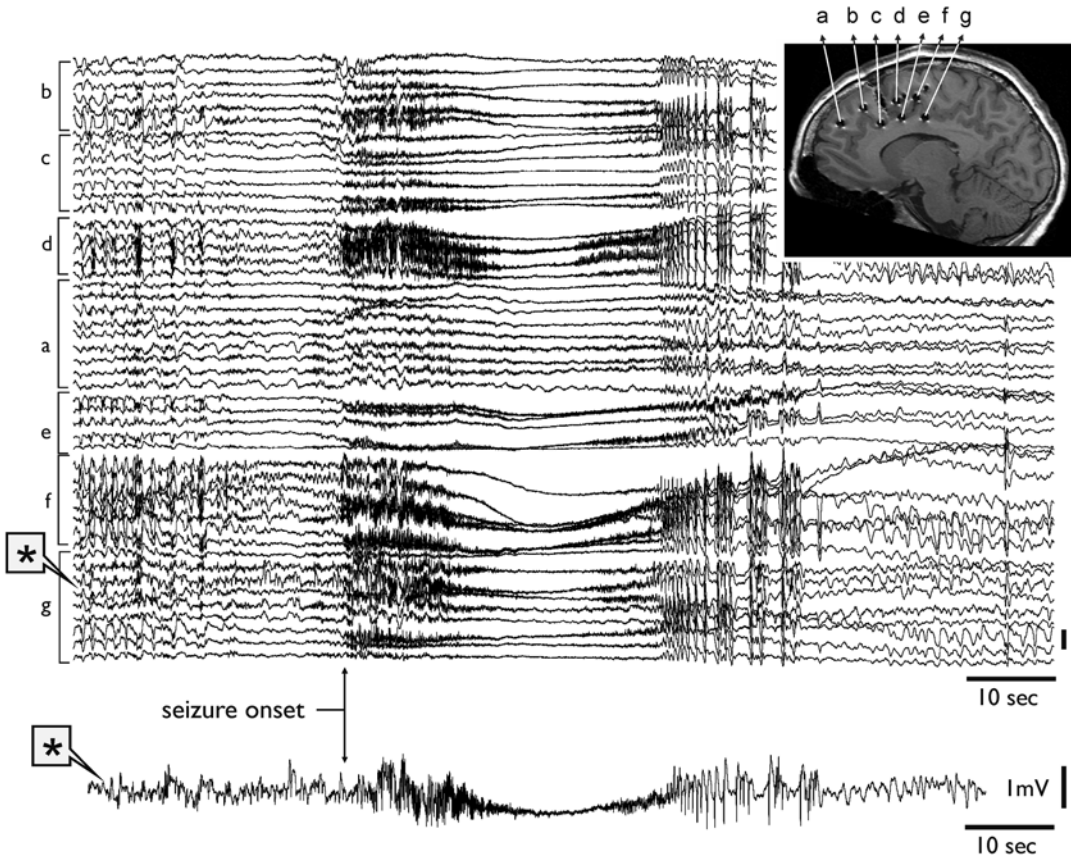
**Fig. 1.6** Recordings with intracerebral stereo-EEG electrodes in a patient with focal epilepsy secondary to focal cortical dysplasia. *Far left:* The position of the recording electrodes is illustrated. *Left:* Interictal discharges recorded with intracerebral stereo-EEG electrodes in a patient with focal epilepsy secondary to focal

cortical dysplasia. *Right:* Seizure onset is marked by the arrow. The *slow spikes* that precede the ictal low-voltage fast activity are different in location and morphology from the interictal spikes (Courtesy of Francione, Tassi and LoRusso of *Claudio Munari* Epilepsy Surgery Center, Niguarda Hospital, Milano)

Intracranial pre-surgical studies revealed that the most consistent pattern observed at the onset of a seizure is characterized by fast activity of low amplitude in the *beta-gamma* range ([5, 38, 48]; for review see [29]) that can be preceded by large

amplitude spike potentials. The latter events have often be defined as pre-ictal spikes, but their consistent and reproducible occurrence at the very onset of a seizure include them by definition as integral part of a seizure. Experimental studies in animal





**Fig. 1.7** Intracerebral recording of a focal seizure with stereo-EEG electrodes (as shown in the *upper right inset*) in a patient with cryptogenic focal epilepsy during pre-surgical evaluation. Multi-contact electrodes are identified by letters. The EEG marked by an asterisk is expanded

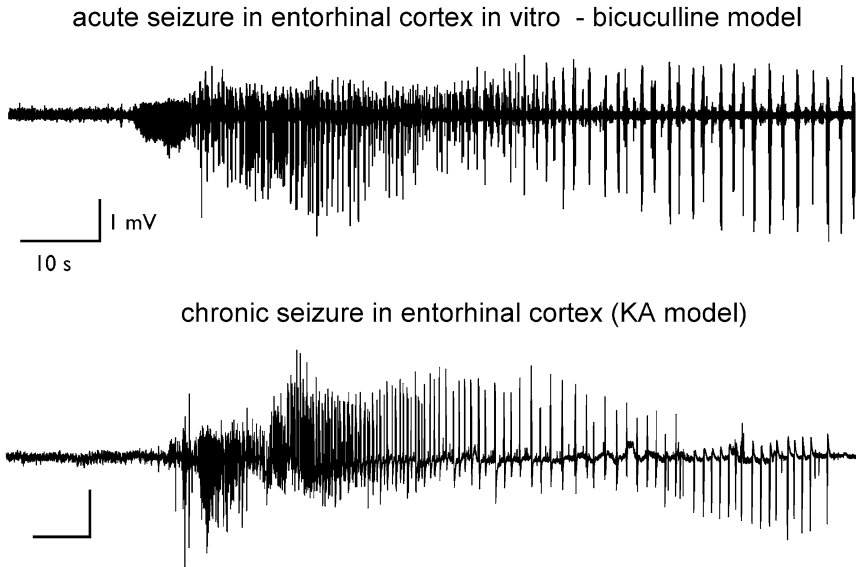
at the *bottom*. When the seizure begins (*seizure onset, arrow*) there is a reduction of background activity, appearance of fast activity, and subsequently there is a very slow potential (From Gnatkovsky, Francione, Tassi and de Curtis, unpublished)

models and in human post-surgical tissue and intracranial stereo-EEG observations demonstrated that these (pre)ictal population spikes are distinct from interictal potentials [21, 44, 56] and are possibly generated by network mechanisms that are different from those sustaining interictal potentials.

More recent studies demonstrated that the low-voltage pattern associated to the initiation of a seizure correlates with the abolition and possibly the desynchronization of background activity. The substitution of background activity with low-voltage fast activity is the intracranial correlate of the electrodecremental pattern defined as EEG “flattening”, a phenomenon that is commonly pursued to localize the seizure onset area on the scalp EEG (as discussed above). Low-voltage fast activity is also associated with the appearance

of large amplitude, very slow potentials lasting several seconds that can be identified on intracranial recordings when low EEG frequencies are not filtered out [9, 57]. These three intracranial electrographic features (fast activity, EEG flattening and very slow potentials) have been proposed as biomarkers of seizure-genesis in the epileptogenic zone [45], since a retrospective evaluation demonstrated that their location on stereo-EEG recordings coincides with the area that has been surgically removed to cure the patient (Fig. 1.7).

The above-mentioned triad of electrographic elements defines seizure networks and the epileptogenic zone in the majority of patients selected for stereo-EEG recordings with intracerebral electrodes. The type of epilepsy referred to surgery could be the reason for the homogeneity of seizure



**Fig. 1.8** Seizures recorded in guinea pig entorhinal cortex. The *upper trace* was recorded in the *in vitro* isolated guinea pig brain after systemic application of 50  $\mu$ M bicuculline. In the *lower panel* a seizure is shown, which was

recorded *in vivo* 3 months after injection of kainic acid in the hippocampus. Both seizures are characterized by fast activity at the onset followed by irregular firing and late periodic bursting (From DeCurtis, unpublished)

pattern reported in the literature. Most of the patients selected for pre-surgical studies, indeed, have pharmacoresistant epilepsies due to either focal cortical dysplasia, low-grade epileptogenic tumors (such as gangliogliomas or dysembryoplastic lesions), or mesial temporal lobe epilepsy with hippocampal sclerosis. Seizures in these types of epilepsy may present with similar EEG features. In mesial temporal lobe epilepsy, seizures that initiate with a hypersynchronous spiking pattern have been reported [8, 93]. Fast activity consistently follows the hypersynchronous discharge, suggesting that this pattern represents a variant of the low-voltage fast activity pattern.

Seizure onset patterns different from low-voltage fast activity have been described during intracranial EEG monitoring, for instance in tuberous sclerosis and in cortical malformations such as polymicrogyria [16, 51, 70, 79]. Whether such patterns are the expression of the epileptogenic network specifically caused by the type of lesion or are due to the failure to implant electrodes precisely in the epileptogenic area, is an open question. Moreover, variable seizure onset patterns have been detected with intracranial and extra-cerebral electrode arrays, such as grid and strips, positioned on the cortical surface in the subdural space. The localizing value of subdural

electrodes has been questioned (e.g., [47, 90]) and, therefore, their ability to define sources and features of ictal patterns is assumed to be less precise than depth electrodes.

Another crucial issue that emerged from intracranial recording studies and can be confirmed by retrospective analysis of earlier reports on seizure patterns, is the demonstration that focal seizures are characterized by a clear sequence of events that starts with a fast activity pattern and ends with highly synchronous, large amplitude bursting. The striking novel finding in this context is the observation that seizures do not initiate with the explosion of sustained, large amplitude, synchronous potentials, as commonly assumed, but feature low amplitude activity and background activity desynchronization that in several occasions last several tenths of seconds. In between seizure onset and seizure termination, a transition from fast, possibly desynchronized activity [59, 82] into an irregular spiking pattern (referred to as “tonic” in several reports) is observed. During the latter phase synchrony of activity builds up and progressively promotes clustering of highly synchronous discharges separated by periods of post-burst depression (Fig. 1.8). The late-seizure bursting (sometimes defined as “clonic phase”)

precedes seizure termination. Interestingly, if seizure onset is restricted to a spatially limited region, seizure termination characterized by synchronous periodic bursting is usually more diffuse and shows the tendency to involve the entire epileptogenic zone. The mechanism for such a widening of the epileptogenic network during the late seizure is still unclear. A synchronizing influence mediated by the involvement of subcortical structures can be proposed. After the end of a focal seizure, post-ictal depression is evident and can be measured as a reduction of background activity in comparison to the pre-ictal condition. These findings can be reproduced in animal models, as discussed in the next section.

In summary, direct evaluation of seizure-generator networks with intracerebral electrodes in focal human epilepsies demonstrates that specific electrographic patterns with a quite reproducible temporal progression define a seizure (typically a focal seizure). De-synchronization of background activity and the appearance of fast low-voltage rhythms characterize seizure initiation and excessive synchronization correlate with termination of the seizure [59]. Post-ictal depression is typical of focal seizures and should always be verified to identify a seizure.

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### 1.3 Seizures, Seizure-Like Events and Afterdischarges in Animal Models

Based on the intracranial human findings observed in focal epilepsies during pre-surgical monitoring, it is mandatory to re-define the term “seizure” in experimental studies of animal models. We will first address *in vivo* studies performed on animal models of seizures or epilepsy, and then discuss *in vitro* studies carried out on preparations featuring complete or partial preservation of brain networks.

Diverse seizure patterns have been illustrated with *in vivo* intracerebral recordings in animal models of epilepsy obtained with different methods and protocols. In several studies, seizure-like patterns were defined only with EEG, i.e., without the aid of video monitoring. This approach is

problematic, because the correspondence of EEG patterns with behavioral symptoms should be verified when seizure events are described. The possibility that the reported EEG potentials are interictal events or even physiological patterns, if not artifacts, should be carefully considered (see Sect. 1.4, below). Incidentally, the lack of a precise definition of a normal EEG in different animal species is a serious limitation to the evaluation of pathological patterns in animal models of seizures and epilepsy. These considerations further support the concept that epileptic phenotypes in animal models should always be carefully analyzed with the aid of video-EEG monitoring, to correlate possible seizure patterns to behavioral/motor changes.

Behavioral seizure correlates are not easy to identify in animals, even when careful electro-behavioral evaluation of the video-EEG is performed, because focal seizures may present with minor symptoms that have little, if any, motor sign. This is a major limitation for seizure identification in animal models: we can only be sure of seizures that correlate with enhanced or decreased motor signs, since other critical non-motor symptoms are difficult to detect. Seizures generated in the hippocampus in animal models (and in patients as well), for instance, can occur during immobility ([8, 15, 80]; see Sect. 1.4, below) and are indistinguishable from normal pauses in behavior unless intracerebral EEG recordings are performed in parallel to video monitoring. In this respect, human EEG studies on the definition of electro-clinical seizure patterns are more standardized and detailed than animal reports. The precise electro-clinical correlation of symptoms during seizures performed in humans demonstrates the finer scientific development of clinical epileptology in comparison to experimental epileptology, and sets an example to improve phenotyping in animal models of epilepsy.

*In vivo* recording of seizures and characterization of seizure patterns have been performed in a relatively small number of studies that describe animal models of epilepsy, largely on temporal lobe epilepsy models developed in rats and mice. Other models in which video-EEG electro-behavioral characterization of focal seizures was

analyzed in detail include post-traumatic epilepsy models [25, 65, 66], models of perinatal anoxia-ischemia [61] and infantile spasms [81]. These reports confirmed that EEG correlates of seizures are largely characterized by fast activity at onset, followed by irregular spiking; and periodic bursting that develops with time during seizures (and usually represents the last pattern before seizure termination: [8, 15, 46, 95]). Post-ictal depression ensues and is infrequently characterized in these models.

Other electrographic potentials that supposedly represent the expression of an epileptic brain have been reported and quantified to support the characterization of epilepsy models. The behavioral correlates of these pathological patterns are often not described (and may not be possible to identify), and in some reports the claim is made that a specific pattern that does not respond to the criteria defined above is regarded as seizure. It is frequently assumed that epileptiform discharges that last longer than 2–3 s can be considered as ictal, as mentioned above [26, 31]. The criterion of duration to discriminate between an interictal and ictal discharge is quite subjective and could be misleading when applied to focal epilepsies. Since a consensus on this issue is still missing, more stringent criteria to define a seizure are required and should be identified.

Seizure patterns comparable to those described *in vivo* in animals (and in human focal epilepsies) can be reproduced in preparations of the entire brain or portions of brain tissue maintained *in vitro* in isolation. Obviously, the absence of the peripheral limbs that expresses motor symptoms prevents any definition of seizure in these experimental conditions. Therefore, the identification of interictal and seizure-like patterns on *in vitro* preparations relies exclusively by electrophysiological recordings, and the identification of stringent criteria for seizure definition is quite critical.

Seizure-like events characterized by fast activity at onset, followed by irregular spiking and terminating with periodic bursting discharges are induced by diverse pharmacological manipulations in adult whole guinea pig brain preparation ([44, 89]; Fig. 1.8), in neonatal en-bloc preparation of cortical areas/systems, such

as the *in toto* hippocampal-parahippocampal structures [30, 64] and in complex tissue slices, in which connectivity between cortical structures is preserved, such as entorhinal-hippocampal slices ([6, 60]; Fig. 1.8).

In several studies performed on slice preparations, prolonged epileptiform events are described, which are characterized either by repeated spikes or by large paroxysmal depolarizing shifts followed by a depolarizing plateau potential on which decrementing discharges occur (see [28]). These types of discharges are often defined as seizure-like, even though their identification as seizures is questionable: similar events, indeed, are never observed during spontaneous seizures recorded *in vivo*, but can be generated by repeated stimulations, as afterdischarges induced by the kindling procedure. In slice studies, the measurement of the duration of “afterdischarges” is usually reported as a criterion to distinguish between interictal and ictal events. This assumption is based on the idea that the mechanisms that generate interictal and ictal events are similar and differ only by the duration and persistence of repetitive spiking or bursting activity. However, this conclusion may not be correct, based on recent findings demonstrating that seizure-like events in complex preparations are initiated with a prominent activation of inhibitory networks, whereas this may not be true for interictal spikes. The analysis of seizure-like discharges in neocortical and hippocampal slices exposed to different pro-epileptic conditions demonstrate that GABAergic networks are active at the very onset of a seizure [30, 39, 40, 67, 99]. These findings were confirmed in the *in vitro* isolated whole guinea pig brain [29, 44]. In this preparation, pre-ictal (ictal) spikes and fast activity that characterize seizure onset correlate with activation of GABAergic interneurons and with a cessation of neuronal firing in principal excitatory cells that last several seconds. In this model, the progression of seizure activity characterized by the transition to the irregular spiking and periodic bursting phases was sustained by ectopic firing of principal cells driven by changes in extracellular potassium induced by inhibitory network activation at seizure onset [88].

In conclusion, the definition of seizure-like events in *in vitro* preparation should be reconsidered and should rely on the reproduction of seizure patterns observed in humans and in chronic animal models of epilepsy. This “reverse translational” approach might help to focus future *in vitro* studies on the mechanisms of seizure generation that more reliably reproduce human focal epilepsy.

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## 1.4 Defining Seizures in Basic Epilepsy Research: Potential Problems Specific to Rats and Mice

Defining seizures in humans requires consideration of several issues, as discussed above. In basic epilepsy research, conducted mainly in rodents (rat or mouse), there are other issues that are important. In order to quantify seizures for preclinical studies, one would want to be precise about seizure onset and seizure termination. However, not only are seizures hard to define, but the exact time of their onset and termination are also problematic. Other issues are also relevant: if there are brief pauses between seizures, when is the pause sufficient to define the events as two separate seizures? Post-ictal depression is often followed by a series of afterdischarges or spikes that become more and more frequent – when does the repetitive spiking become frequent enough to be called the onset of the next seizure? This issue is not only important in establishing seizure frequency, but it also is important when defining status epilepticus (SE). When examined at high temporal resolution, there are often pauses between seizures during SE. Does this mean it is not SE? If there are no convulsions (non-convulsive SE) how does one determine what is and what is not SE? Similar to humans, defining a seizure in rodents is not as easy as one might think.

### 1.4.1 Behavioral State

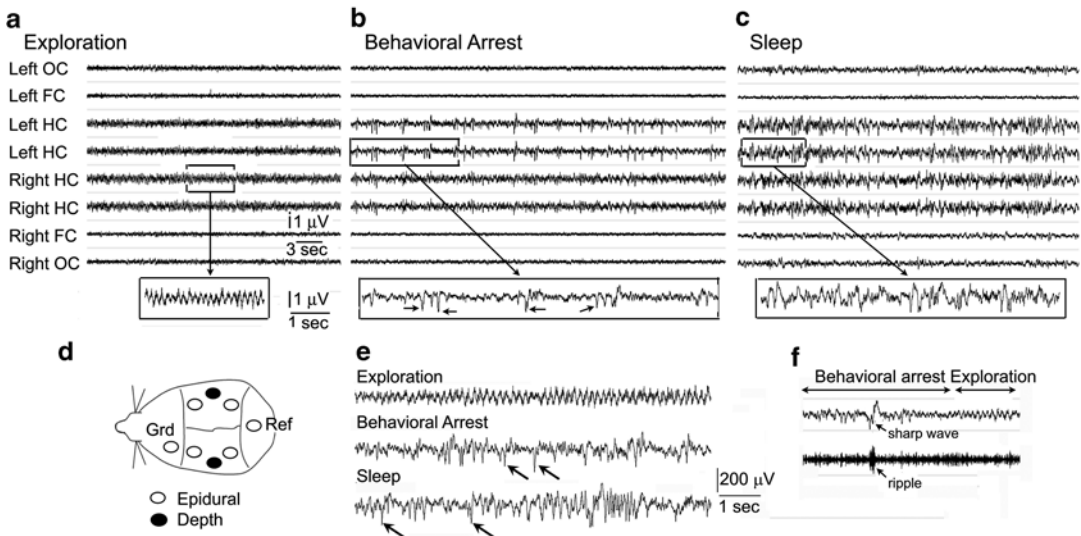
There are several behaviors that make up the vast majority of the lifespan in rats and mice: explora-

tion, sleep, grooming, eating and drinking. In addition, there is a state called “quiet immobility,” “awake rest” or “behavioral arrest” where rodents stop moving, their eyes are open, and they stare blankly into space. Typically the animal is standing at the time, and has just walked across the cage or explored its surroundings. Unlike humans, this behavioral state can be prolonged (over 10 s). It presents problems for the epilepsy researcher because it appears similar to an absence seizure. Therefore, understanding the normal behavioral states of rodents, and their EEG correlates, is important for epilepsy researchers using these species.

#### 1.4.1.1 Hippocampal EEG Associated with Exploration: Theta Rhythm

Associated with exploration, behavioral arrest, and sleep in rodents are distinct EEG rhythms that can be recorded with chronic electrodes implanted in hippocampus [14]. As shown in Fig. 1.9a and originally described by Green and Arduini [50], EEG oscillations at theta frequency (commonly called theta rhythm) are recorded in hippocampus when an animal explores. Theta oscillations vary in frequency but are typically 6–10 Hz in rats and mice [12, 50, 91].

In animal models of epilepsy, theta rhythm is interesting because epileptic animals are less likely to exhibit seizure activity during exploratory behavior, when theta oscillations occur in hippocampus [69]. This “anticonvulsant” nature of exploration and theta rhythm in hippocampus has been attributed to many potential mechanisms but has not been defined conclusively [22]. It is useful to record theta oscillations *in vivo* because large theta rhythm is found in hippocampus but it is much smaller or not observed elsewhere. Therefore, theta oscillations can be used to confirm the recording is in hippocampus. Theta oscillations are also useful to record because their amplitude can be used to define the specific layer within hippocampus where the recording electrode is located. For example, if a stimulating electrode is used to evoke field EPSPs in area CA1 from the Schaffer collateral input, the field EPSP should be recorded in the layer where theta is relatively small, stratum radiatum.



**Fig. 1.9** EEG characteristics in the normal adult rat. (a) Using 8 electrodes (shown in d), awake behaving rats were recorded in their home cage. During exploration, hippocampal electrodes exhibited theta oscillations. The area outlined by the *box* is expanded at the *bottom*. (b) During a spontaneous arrest of behavior, sharp waves (*arrows*) occurred regularly in the hippocampal EEG. (c) During sleep, the hippocampal EEG became active. (d) The recording arrangement included 4 epidural

electrodes and 2 twisted bipolar electrodes in the dorsal hippocampus, one in each hemisphere. *Grd* ground; *Ref* reference. (e) A summary of a-c is shown. In three behavioral states there are large differences in the hippocampal EEG with sharp waves (*arrows*) in behavioral arrest and sleep. (f) During sharp waves, filtering in the ripple band (100–200 Hz) shows that a ripple occurs at the same time as the sharp wave (From LaFrancois and Scharfman, unpublished)

In contrast, where theta is larger, the adjacent stratum lacunosum-moleculare, the field EPSP evoked by the same stimulus would be small or have a positive polarity. Because the entorhinal cortex is a source of theta rhythm (the other major source originates in the septum; [12]), theta oscillations are very large in stratum lacunosum-moleculare and the outer two-thirds of the molecular layer of the dentate gyrus, where the entorhinal cortical projection (the perforant path) to hippocampus terminates.

#### 1.4.1.2 Hippocampal EEG Associated with Behavioral Arrest: Sharp Wave-Ripples

The hippocampal EEG shown in Fig. 1.9b is taken from a rat that explored and then paused – entering a period of behavioral arrest. As described by Buzsaki originally [10, 11], the hippocampal EEG changes dramatically when an animal stops exploring and pauses in a frozen stance, with eyes still open. Theta oscillations

decrease and the EEG becomes irregular. In addition, sharp waves (SPWs) occur intermittently. SPWs are ~100 msec duration spikes that reflect synchronous firing in a subset of area CA3 neurons, which in turn activate area CA1 apical dendrites by the Schaffer collateral axons and the dentate gyrus, most likely by backprojecting axons of CA3 pyramidal cells. Therefore, SPWs can be recorded in many locations within hippocampus [10, 11].

The term SPW is important to discuss in the context of epilepsy, because it is sometimes used interchangeably with the term interictal spikes (IIS). Hippocampal SPWs are distinct from interictal spikes because hippocampal SPWs occur without seizures, i.e., they are not interictal (between ictal events). Hippocampal SPWs are recorded only in hippocampus- if one moves a recording electrode just outside the hippocampus, SPWs are not observed (Pearce and Scharfman, unpublished). IIS in an epileptic rodent can be typically recorded from multiple

cortical electrodes simultaneously at many sites in the brain. However, SPWs can be generated by circuits outside hippocampus, i.e., other types of SPWs besides those generated in area CA3. For example, SPWs are generated in entorhinal cortex and piriform cortex [68]. Notably, the underlying mechanisms for an IIS may or may not be the same mechanisms for a SPW, although they do seem related. For example, GABAergic mechanisms may trigger IIS (as discussed in the previous section); GABAergic network oscillations (ripples) are also involved in SPWs. The classic view of the IIS is that it is generated by a giant paroxysmal depolarization shift (see previous section); a synchronous depolarization in pyramidal cells also drives SPWs. Regardless, if SPWs and IIS are terms that are used synonymously, there may be differences in the underlying cellular processes/mechanisms that are overlooked, so it is important to consider the terms carefully.

When recording electrodes are positioned near the CA1 pyramidal cell layer, fast oscillations called ripples [84] can be detected at about the same time as the SPW (Fig. 1.9f). Therefore, the term “SPW-R” (sharp wave-ripple) is now used instead of the original term, sharp wave. Ripples in the hippocampal EEG correspond to synchronous oscillations of pyramidal cells that are caused by rhythmic IPSPs that are initiated by action potentials in a subset of hippocampal GABAergic interneurons that innervate pyramidal cell somata and initial axon segments. As synchronous release of GABA from these peri-somatic targeting interneurons hyperpolarize pyramidal cell somata that are in close proximity, chloride ions enter the pyramidal cells in a repetitive manner and cause a series of extracellular positivities. The positivities wax and wane as the pyramidal cell IPSPs start and stop, leading to an oscillation [19].

#### 1.4.1.3 The Hippocampal EEG Becomes Active During Sleep

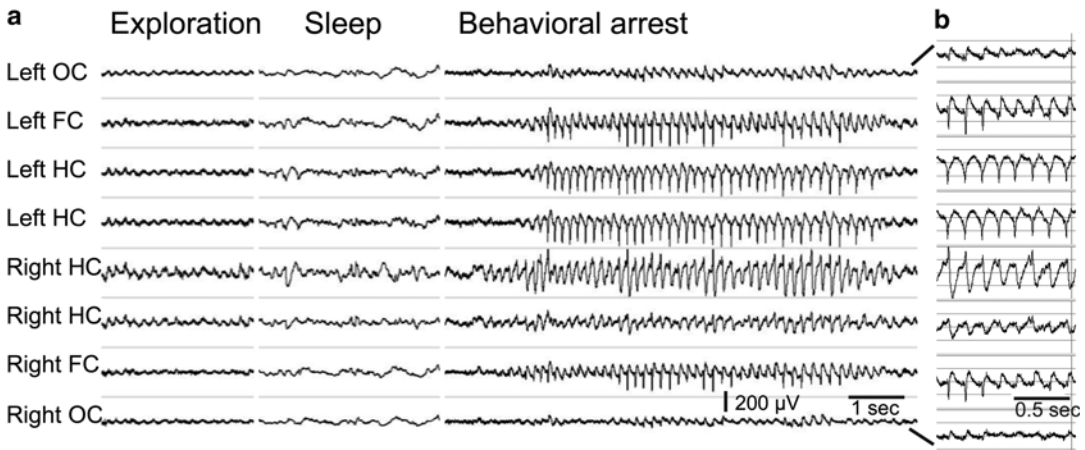
The hippocampal EEG becomes extremely active during sleep in the rodent, and is irregular, called large irregular activity (LIA: Fig. 1.9c). The increase in the hippocampal EEG is often

simplified as a type of disinhibitory state that coincides with a ‘switch’ from sensitivity to sensory input to a state where intrinsic circuitry is active [52]. A similar idea has been proposed for piriform cortex during slow-wave sleep; odor input is reduced in favor of processing between piriform cortex and other forebrain sites [97]. For the epileptologist, it is important to recognize that comparing the hippocampal EEG between animals without considering the behavioral state may make one animal seem normal (if it is exploring) compared to seizure like activity in the other if it is asleep (Fig. 1.9). Compressing the EEG can make this more difficult; for example, if the EEG is compressed it is hard to distinguish a noisy baseline from theta oscillations, so the EEG may look inactive when an animal is exploring. For these reasons, expansion and compression of the EEG should be varied during examination of the EEG for seizures. In addition, the type of electrode and recording system should stay the same for any given set of experiments.

#### 1.4.1.4 When Normal Activity Appears To Be Epileptic

One of the implications of the discussion above for epilepsy research is the possibility that normal EEG activity may be mistaken for seizures. For example, an investigator may think that the animal is freezing because it is having a seizure, but actually exhibiting normal behavioral arrest. This interpretation is based on the limbic seizure stage scale of Racine, who based the scale on behaviors of rats during electrical stimulation of the amygdala during kindling. He suggested that there was initially a period of immobility with small mouth or face movements with small mouth or head movements, and called this a stage 1 or 2 in his scale of limbic seizure severity [78]. The only problem with this idea is that it can be confused with behavioral arrest.

During behavioral arrest, investigators could interpret the irregular activity and repetitive SPW-Rs to be a seizure (Fig. 1.9). Likewise, the transition from behavioral arrest back to exploration may seem like the termination of a seizure, particularly when the EEG is compressed



**Fig. 1.10** Spike-wave discharges recorded from the normal adult hippocampus of the rat. **(a)** A recording from an adult Sprague-Dawley rat shows typical EEG activity during exploration and behavioral arrest. In behavioral arrest, there were spike-wave discharges.

Animals were monitored during the recordings to be sure that artifacts related to grooming or chewing did not occur during spike-wave discharges. **(b)** Recordings in **a** are expanded (From Pearce and Scharfman, unpublished; see also [101])

(Fig. 1.9). In light of these potential problems, describing stage 1 seizures without a hippocampal electrode is problematic. An animal that suddenly stops and appears unresponsive could be interpreted to have a stage 1 seizure when it actually is pausing between episodes of exploration.

Another problem arises in studies of seizure frequency evaluated over time. For example, studies of epileptogenesis often record animals over weeks. There is typically no consideration of behavioral state when the results are quantified. If there is less exploration because an animal is sleeping more, EEG power in the theta band may decrease. EEG power in high frequency bands may increase if there are more SPW-Rs because the animal is pausing more, or sleeping more.

#### 1.4.2 Spike-Wave Discharge

In many strains of rats, the state of behavioral arrest is accompanied by spike-wave discharge in thalamocortical networks [20, 27, 96]. These discharges have been noted in almost every strain of rat, such as Long-Evans [83],

where approximately 90 % of female rats exhibited spike-wave discharges spontaneously by 4 months of age. In Wistar rats, Galewicz [49] reported that 73 % of male rats showed spike-wave discharges by 6 months of age and 93 % of males at 24 months of age. Kelly [63] reported spike-wave discharges in female Fischer 344 rats at 4 and 20 months of age. In rats that are genetic models of absence epilepsy (GAERS, Wag Rij) spike-wave discharges are a characteristic of the strain, and used to gain insight into mechanisms of absence epilepsy [20, 27]. Numerous genetic models of absence epilepsy also exist based on spontaneous mutations in mice (*e.g.*, *lethargic*; [18, 74]). As shown in Fig. 1.10, spike-wave discharges accompany behavioral arrest in naïve Sprague-Dawley rats. These discharges vary according to the sex, age, environment and other factors [13] but are not always observed [96], making control recordings critical to any study of rats in an animal model of epilepsy.

These observations raise several questions: are spike-wave discharges in rodents normal? It has been suggested that they could serve important purposes related to sensory processing [71, 86, 94] or aging and excitability [71].



If this is true in rodents, is human spike-wave discharge normal too? One possibility is that spike-wave discharges and behavioral arrest were present in early stages of evolution and then reduced because behavioral arrests (without complete attention) would be dangerous in the presence of predators – vigilance would be advantageous. In humans, the spike-wave discharges that do arise may be vestiges of rodent circuitry that have not completely been removed by evolution. Photic stimulation can trigger spike-wave discharges in humans [24, 87], and may be a method to trigger these ‘vestigial’ oscillations.

Another implication of the observations in rats in Fig. 1.10 is relevant to the detection of seizures in hippocampal electrodes in rodent studies of epilepsy. In Fig. 1.10, the hippocampal electrode appears to show rhythmic spiking when spike-wave discharges occur in the frontal and occipital leads. The rhythmic spiking in hippocampus could be volume conducted from thalamus, or it could reflect hippocampal neural activity. In light of the fact that the frontal cortical lead shows spike-wave oscillations, one would know that volume conduction in the hippocampal lead is a possibility. However, if there were only an electrode in hippocampus, which is a common recording arrangement in epilepsy research, the rhythmic activity in hippocampal electrodes might be interpreted to be a seizure generated in hippocampus. Because it is accompanied by a frozen, ‘absence’ behavior, it could be concluded that there was a Racine stage 1 seizure. Importantly, some of the normal rodents with spike wave discharges also have head nodding or mastications, which could make an investigator more convinced of seizure activity – because these movements were also noted by Racine in his classification of stage 1–2 behaviors. Importantly, most of the spike-wave discharges occur at approximately 7–9 Hz and are stable (in frequency) within a spike-wave episode or across episodes (Fig. 1.10; [13, 20, 27, 96]). Therefore, rhythms at this frequency (e.g., theta rhythm) that occur in hippocampus can be a signal to investigators to interpret their EEG data cautiously.

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# What Is the Clinical Relevance of *In Vitro* Epileptiform Activity?

# 2

Uwe Heinemann and Kevin J. Staley

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## Abstract

*In vitro* preparations provide an exceptionally rapid, flexible, and accessible approach to long-standing problems in epilepsy research including ictogenesis, epileptogenesis, and drug resistance. Acute slices suffer from a reduction in network connectivity that has traditionally been compensated through the application of acute convulsants. The utility and limitations of this approach have become clear over time and are discussed here. Other approaches such as organotypic slice preparations demonstrate the full spectrum of spontaneous epileptic activity and more closely mimic human responses to anticonvulsants, including the development of drug resistance. Newly developed transgenic and vector expression systems for fluorophores, optogenetics, and orphan receptors are being coupled with advances in imaging and image analysis. These developments have created the capacity to rapidly explore many new avenues of epilepsy research such as vascular, astrocytic and mitochondrial contributions to epileptogenesis. Rigorous study design as well as close collaboration with *in vivo* laboratories and clinical investigators will accelerate the translation of the exciting discoveries that will be revealed by these new techniques.

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## Keywords

Seizure • Epilepsy • Epileptogenesis • *In vitro* models • Translation • Ictal • Interictal

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## 2.1 Current Challenges

From the standpoint of translation, experimental epilepsy research is confronted with two major problems: the first is the discovery of mechanisms underlying drug resistance in epilepsy and development of new agents that would be useful in seizure control in pharmacoresistant patients.

The second is discovering the signaling cascades that lead from an initial brain injury to epilepsy later in life, and conversely, discovering signaling cascades that would prevent this process, that is commonly referred to as epileptogenesis. It is important to note that epileptogenesis occurs in only a small percentage of patients who have suffered brain injuries. For example a recent prognostic study reported that only 8.2 % of patients developed epilepsy after stroke [46]. Thus it is important to identify biomarkers that predict which patients will develop epilepsy. It is also important to analyze how these biomarkers illuminate or participate in the process of epileptogenesis. Research into this issue will provide opportunities to identify mechanisms which protect the brain against seizures. To keep these and other research projects in epileptogenesis from being “lost in translation”, a number of general guidelines should be kept in mind:

- (i) **Spectrum:** Carefully consider the disease and the entire range of observations during the disease in probing for their potential role. This applies, for example, to the observation that astrocyte activation often precedes epilepsy in many animal models. This raises the possibility that one component of acquired epileptogenesis may involve the influence of astrocytes on synaptic and cellular properties of neurons, microglia, NG2 cells and vascular cells. Thus it may be useful to test whether prevention of astrocyte activation has an antiepileptogenic effect.
- (ii) **Statistics:** matching the experiment to the disease. When experimental groups are too small relative to the variance of the parameters that will be studied, the chance of falsely positive results increases. Many of our current animal models of acquired epilepsy have been developed to ensure that a large percentage of animals develop epilepsy. This hinders development of useful biomarkers, because their predictive value depends on the incidence of the disorder. The incidence of epilepsy after brain injury is much lower in humans, so the predictive value of a biomarker needs to be tested in an experimental population with a similar incidence. Finally, animals that do not develop epilepsy after brain injury are useful for more than service as controls – we may be overlooking antiepileptogenic characteristics and processes that may provide additional prognostic, mechanistic, and therapeutic insights into epileptogenesis after brain injury.
- (iii) **Heterogeneity:** To maximize the chance that results will extrapolate to humans, preclinical studies should include more than one species and take intra and interspecies inhomogeneity into account. Most animal studies including in vitro studies are done on animals which are rather young and come from genetically homogenous breeding stocks. Hence the genetic inhomogeneity of human species is not taken into account. A second source of inhomogeneity is the injury itself. Common human brain injury mechanisms include trauma, infection, and both global and focal hypoxic- ischemic insults. The severity and anatomical location of each of these injuries varies profoundly from patient to patient. Understanding which circuit elements are altered after both experimental and clinical brain injury will be a necessary step in evaluating the risk and rate of subsequent epileptogenesis.
- (iv) **Comorbidity:** multiple hits are often a critical factor in human disease, but this is usually not considered in experimental work. One approach to correcting this oversight could entail choosing the right animals for study. An example would be using stroke models of epileptogenesis in rodent models of chronic hypertension or type 2 diabetes.
- (v) **Communication between basic and clinical epileptologists:** “Losses in translation” often arise from the different perspectives of experimentalists and clinicians, combined with the barriers to free communication between these groups. Ideally, interactions between clinical and basic investigators should be sufficiently close that experimentalists can contribute to clinical study focus and design. This requires a centralized infrastructure to provide close scientific collaboration as well as institutional mechanisms

for patient access, patient monitoring, access to biostatistical resources, and guidance for approved use of patients in research. Perhaps the greatest institutional challenge is provision of protected time for interactions between clinical and basic researchers.

- (vi) Pipeline repair: The interruption of classical translational pipelines also leads to losses in translation. Because drug resistant epilepsy is relatively rare, most of the pharmaceutical industry has lost interest in drug development for this type of epilepsy. Filling this gap, including toxicological and pharmacokinetic studies in preclinical and clinical populations, will require the training of clinician scientists who are equipped for drug development in an academic environment. This will require new ways of financing such research, as well as developing processes to provide academic credit for the type of applied research that is essential for the later stages of drug development involving toxicology, compounding, and pharmacokinetics.

## 2.2 How Can *In Vitro* Research Help Meet These Challenges?

The advantages of *in vitro* models have long been recognized in epilepsy research, starting with the pioneering work of P. Schwartzkroin [77]. These advantages include speed, convenience, low cost, the availability of a wide variety of genetically modified animals from which slices can be prepared, and electrophysiological, pharmacological, and optical accessibility. *In vitro* models make it possible to understand pathophysiology at a high level of electrophysiological, molecular and cell biological resolution.

*In vitro* models have a number of drawbacks. *In vitro* preparations usually have no blood brain barrier, and there is no circulation. Rather, drugs are applied in an aqueous solution, and reach their targets by routes that are more relevant to CSF administration than oral or intravenous routes. Brain slice preparation induces massive damage to afferent and efferent circuitries, which

is a particularly significant problem in the investigation of network-level phenomena such as seizures. Experiments are done at non-physiological oxygen and glucose concentrations. Many of the preparations are based on tissue from perinatal animals. For example, organotypic slice cultures, or the intact (whole) *in vitro* hippocampus preparation are best prepared before the 8th postnatal day (P8). Although acute brain slice preparations can be obtained from animals at any age, there is only a relatively brief period of time when the slice is physiologically stable. This time limit can restrict the types of experimental manipulations that can be performed *in vitro*, such as those involving viral or expression of exogenous proteins.

Some scientists and clinicians argue that *in vitro* models are too far removed from human epilepsy, and therefore one should focus on *in vivo* models. However, *in vivo* models have the dual problems of complexity and access, such that it is difficult to identify the pathogenic mechanisms in sufficient detail to initiate pharmacological or genetic interference. Moreover, studying acquired epileptogenesis *in vivo* involves brain injury. Therefore the “3R” strategies of replacement, reduction and refinement (3Rs) in research using animals are relevant.<sup>1</sup> “Replacement” refers to the use of other preparations, such as induced pluripotent stem cells derived from patient fibroblasts. “Refinement” refers to alteration of the experiment to focus the experiment to minimize pain and maximize information return. For example, many conditions leading to epilepsy are associated with activation of astrocytes. Addressing this question specifically might involve replacing a status epilepticus model with a model in which astrocytes are primarily activated [50, 66]. This will – if some investigators are correct – still cause epilepsy but presumably with less damage to the brain. “Reduction” refers to minimization of the number of animals used. Preparation of multiple brain slices per animal can make possible multiple independent tests of the hypothesis for each experimental subject. Using the reactive astrocyte hypothesis as an example, many of the

<sup>1</sup> <http://www.nc3rs.org.uk/>

consequences of astrocyte activation can also be studied *in vitro*. Some questions may not be feasibly studied *in vivo* – for example disturbance of potassium homeostasis [42] and/or glutamate homeostasis [29] may not be detectable with currently available *in vivo* methods. Focusing on the 3 Rs can have beneficial consequences – for example, markers of astrocyte activation might prove to be a biomarker predictive of epileptogenesis.

In the end it is important to recognize “*in vivo veritas*,” i.e., *in vitro* studies should be complemented by *in vivo* studies. For example, *in vitro* studies can be used to rapidly screen drug libraries or target proteins and RNA, and slower, more costly, but more relevant *in vivo* studies can be used to study the most promising lead compounds. Indeed *in vitro* studies often underestimate potential side effects. *In vivo* experiments can determine whether translational relevance is hampered by the unwanted side effects of a given intervention, by toxic effects on organs other than the brain, by long-term loss of efficacy due to development of tolerance, or ineffectiveness due to interference with attention or sleep states of an animal and a patient.

If *in vivo* studies are used to complement *in vitro* work, it is important to optimize the *in vivo* protocols for maximal translational relevance. One improvement in the translational efficacy of *in vivo* studies could be achieved by completely phenotyping animals undergoing epileptogenesis and experimental therapeutic studies. A critical aspect of thorough and unbiased phenotyping of mice or rats includes continuous seizure surveillance using video EEG monitoring [71]. Wherever possible, experimental approaches should be employed that are based on clinical observations.

One chance to strengthen epilepsy research is also to take advantage of pathophysiological discoveries in other disciplines. For example the abnormalities observed in patients with Alzheimer’s disease may not only be relevant for neurodegenerative disease but also for epilepsy because many patients with Alzheimer’s disease may also develop a symptomatic form of epilepsy [88]. Mitochondrial disorders are not only

observed in certain forms of Parkinson’s disease [13] but also in epilepsy [48]. Elements of the inflammatory response are observed in many brain injuries (for example after trauma, stroke and status epilepticus) which may contribute to epileptogenesis [44, 89]. If useful discoveries and approaches in other areas of applied neuroscience are exploited, the translational gap may be more readily overcome.

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### 2.3 Lessons Learned: In Vitro Techniques for Epilepsy Research

**Ictogenesis:** The utility of *in vitro* preparations for epilepsy research was first suggested by a study in which seizure like events were induced by lowering of extracellular  $\text{Cl}^-$  concentration in the perfusate of acute hippocampal slices [93]. It was therefore surprising that  $\text{GABA}_A$  receptor antagonists induced only short interictal-like discharges in the hippocampal slice preparation, because the hippocampus was presumed to be the most epileptogenic region [77]. Similar findings were observed in cortical slice preparations. These data suggested that  $\text{GABA}_A$  receptor blockade was not a sufficient condition for seizure induction *in vitro*, and that other conditions were necessary for ictogenesis, i.e. the induction of seizure-like events. The first seizure-like events recorded *in vitro* were generated by conditions that accompany seizures *in vivo*, such as low concentrations of extracellular  $\text{Mg}^{2+}$  [90] or  $\text{Ca}^{2+}$  [43] or elevated concentrations of extracellular  $\text{K}^+$  [87]. While low  $\text{Ca}^{2+}$  and high  $\text{K}^+$  induced seizure-like events in hippocampal subregions, low  $\text{Mg}^{2+}$  and application of 4-aminopyridine, a potassium channels blocker, initiated seizure-like events more reliably in cortical structures than in hippocampal slices [60], unless juvenile tissue was used [31]. These studies suggested that the hippocampus is not as seizure prone as originally thought from the pathological studies of patients with epilepsy, or that seizure generation involves distributed circuits that are lost after slicing-induced deafferentation. In light of this, it is interesting to note that seizures in patients with temporal lobe



epilepsy often originate outside the hippocampus, for example in the amygdala and the entorhinal cortex [81]. These cortical structures are now considered to be more seizure prone than the hippocampus or other cortical areas. The ionic manipulations that were used to study ictogenesis also provided early insights as to why and where (cortex vs. hippocampus) Mg administration acts to antagonize eclamptic seizures [24].

The next surprising finding from *in vitro* studies of epilepsy was that seizure induction was more easily accomplished in control slices than in slices obtained from animals with epilepsy and from specimens of patients with drug resistant epilepsy [30, 98]. This suggests either that ictogenic processes are active only transiently in epilepsy, or that epilepsy also entails protective mechanisms that are more robustly preserved *in vitro* compared to ictogenic mechanisms. Understanding such protective mechanisms could lead to new antiepileptogenic strategies. A related insight from recordings of tissue from patients with epilepsy was that the transections that accompany slice preparation may be more functionally important in chronically epileptic tissue than normal tissue. One interpretation is that a fundamental and widespread alteration in connectivity occurs in the chronically epileptic brain. On the other hand, many patients with refractory epilepsy know that following a seizure there is usually a seizure-free interval, sometimes of considerable duration. Here, translation is bidirectional: Clinical questions can be “translated” into an experimental approach and experimental observations suggest new possibilities for interfering with epilepsy and epileptogenesis.

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## 2.4 In Vitro Models: The “Nuts and Bolts”

Among the *in vitro* preparations available for studying ictogenesis, perhaps the most versatile is the slice preparation. Slices can be obtained from any mammalian species including humans following neurosurgical interventions. Slices can also be obtained from animals that are epileptic

as a result of trauma, tumors, status epilepticus, inflammation, etc. However, slices have circuits that have been reduced in size by transection of processes, and therefore more intact preparations are sometimes required to gain insight into epilepsy. There are two acute preparations that address the connectivity issue: one is the isolated, intact hippocampus (also referred to as the whole or *in toto* hippocampal preparation [61]). The other is the intact isolated brain preparation [19]. The first preparation is only feasible if animals are used at young ages, and the second preparation is feasible only if guinea pigs are used (unless one uses non-mammalian species). Also, in these preparations, many aspects of epilepsy cannot be readily studied. A preparation that can be used for long-term observations is the organotypic slice culture, in which many different aspects of epileptogenesis can be studied “in a dish.” Slice cultures represent a model of brain trauma by virtue of the trauma involved in slice preparation, and also of developmentally increased seizure susceptibility because as mentioned above, organotypic slices are most reliably prepared from animals in the first postnatal week. In the following section we will discuss each of these preparations in more detail, and discuss some translational aspects of this research.

### 2.4.1 Acute In Vitro Brain Slices

#### 2.4.1.1 Interictal Activity and Seizures

Interictal activity refers to paroxysmal epileptic discharges that are much more brief and occur more much more frequently than seizures. Most epileptiform events in acute slices have these two characteristics. Currently, it is not known whether interictal epileptiform activity is pro-convulsive or epileptogenic. Because EEGs are not performed routinely in brain injured patients, it is not known whether interictal spikes precede seizures after brain injury. In acute *in vitro* preparations, brief recurrent epileptiform events can be readily induced by ionic and pharmacological manipulations. Thus many experimentalists have argued that these interictal-like events observed *in vitro* embody the essential features of ictogenesis.

Indeed preictal spikes often precede seizures in humans, and in *in vivo* models of epilepsy. During the *in vivo* spikes, depolarization shifts of the membrane potential occur that are very similar to the membrane potential changes recorded *in vitro* [45, 78]. On the other hand, questions have been raised about the translational relevance of activity that is induced by acute proconvulsant pharmacological or ionic manipulations *in vitro* [94].

One example is the interictal-like activity induced by low Mg induced in adult rat hippocampal slices. The pharmacological relevance of this activity is modest, because anticonvulsants have limited effects on this activity. Thus this activity cannot be used in drug screening, and pathophysiological studies based on this activity must be interpreted with caution. In fact, seizure-like events were induced when the GABA<sub>B</sub> agonist baclofen was added to the perfusate containing low Mg [83]. One interpretation of the combined effect of baclofen and low Mg is that interictal activity was preventing ictogenesis. Another interpretation is that low Mg induces a state similar to Periodic Lateralize Epileptiform Discharges (PLEDs) rather than interictal spikes, and that seizures can only be observed by reducing the severity of the ictogenic conditions, for example by reducing probability of glutamate release through activation of presynaptic GABA<sub>B</sub> receptors with baclofen. Similar results are obtained when elevated K, which induces spontaneous epileptiform burst discharges in adult hippocampal slices, is combined with strontium, which reduces the rate of glutamate release [84].

Consistent with these observations of the interaction between manipulations that alter excitability, induction of recurrent epileptiform burst discharges by tetanic stimulation makes it more difficult to induce seizure-like activity using elevated K<sup>+</sup> [59]. Barbarosie and Avoli [7] observed that in the presence of the convulsant 4AP, seizure activity could be initiated following transection between CA1 and entorhinal cortex.

These experiments and many others performed over the last two decades, emphasize that the sum of multiple manipulations *in vitro* are not predictable and are often difficult to interpret

with respect to human epilepsy. A particularly problematic correlate is that in slices exposed to convulsant conditions, anticonvulsants, even at anesthetic concentrations, while blocking seizure-like events [14, 57, 99], rarely inhibit the interictal-like epileptiform activity induced by convulsants in acute *in vitro* preparations [94]. This is an important area for future optimization, and is discussed in more detail in the section on organotypic slices.

#### 2.4.1.2 Age, Area, and Astrocytes

Slice preparations can be used to determine age dependence of seizure susceptibility, and to compare different regions of the brain with respect to epileptogenesis. Thus, susceptibility to low Mg or low Ca is much higher in tissue from young animals [31] and treatment with ictogenic agents often results in spreading depression [38]. There are many potential developmental mechanisms to explain these data, including circuit development and or the maturation of astrocytes. At the time when Hablitz and colleagues reported their findings, it had not yet been established that astrocytic properties differ in epileptic vs. control animals [29, 40]. It also was not known that activation of astrocytes can be achieved by many perturbations that are considered minor, such as opening of the blood brain barrier, or exposure to albumin. Conditions leading to chronic astrocyte activation can also acutely reduce seizure threshold. Brain injuries are often accompanied by spreading depression episodes, which in cases of disturbed neurovascular coupling can cause neuronal damage and may therefore exacerbate brain injury in ischemic and hemorrhagic stroke. Spreading depression may also be relevant to epilepsy associated with Alzheimer disease [22, 55, 92].

The regional variation of seizure susceptibility can also be investigated with acute brain slices. Seizures never seem to originate from basal ganglia and cerebellum, perhaps because information is relayed by activation of inhibitory cells through disinhibition of target cells. This brings up the fact that all brain activity involves both excitation and inhibition of both inhibitory neurons and principal cells. Thus pharmacoresistance could involve

failure of GABAergic agents (which may exacerbate inhibition of interneurons) or Na channel blocking agents (which may reduce GABA release from strongly inhibited interneurons). Of course there are many other possible explanations for pharmacoresistance, such as alterations in the expression of Na channels.

#### 2.4.1.3 Channelopathies

A good example of a genetic channelopathy that has been benefited from the *in vitro* approach is the murine Nav1.1 knockout model of epilepsy. These mice exhibit a pattern of seizure activity that is similar to the clinical syndrome with a similar defect in Na channels, Dravet syndrome [74]. One hypothesis for the generation of seizures in these animals is that they arise from preferential expression of Nav1.1 channels in interneurons which – if defective – would result in strongly reduced excitability of interneurons and GABA release [95]. The result could be a pharmacoresistant epilepsy.

Defects in ion channels have long been implicated in the epileptiform discharges induced *in vitro* by low extracellular calcium concentrations [36, 52]. This idea was based on observations from baboons where seizures induced by stroboscopic stimulation were associated with decreases in extracellular Ca concentration to less than 0.2 mM, and where seizures were accompanied by increases in potassium concentration to near 10 mM [69]. Mimicking this condition did not induce seizure-like events in human or animal cortical structures *in vitro*. An exception was area CA1 in rat and mouse hippocampus, where the packing density of neurons is higher, promoting ephaptic interactions that are thought to be enhanced by lowered extracellular calcium. On the other hand, cation channels [37] and more recently certain TRP channels are activated by decreasing either Ca or Mg concentration [91]; this is not only important for spreading depolarization [80] but possibly also for generation of seizures and cell death [62]. The regulation of these excitatory channels by Ca and the activity-dependent decrease in extracellular Ca suggests a new mechanism for seizure spread and for modifying seizure generalization.

Similarly recent work has identified KCNQ channels as potential targets for the treatment of seizures. The first drug introduced for treatment is retigabine which affects KCNQ2,3 and 5 channels but not KCNQ 1 channels (KCNQ1 is expressed in the heart; [96]). The distribution of KCNQ channels in principal cells and in interneurons varies, with KCNQ5 channels being expressed on GABAergic cells as well as principal cells [97], which can explain the finding that retigabine can reduce GABAergic inhibition, which may limit its use as an antiseizure medication. More recently however, agents were identified which only affect KCNQ2 [12]. These are preferentially expressed on glutamatergic cells therefore are more suitable as antiseizure drugs. Thus slices prepared from transgenic animal models can make possible the rapid testing of potential anticonvulsant effects; however, side effects are better assessed *in vivo*.

#### 2.4.1.4 Evoked Seizure-Like Events In Vitro as a Model of Status Epilepticus

*In vitro* models of ictogenesis such as the high K model, the low Mg model, and the 4-AP model are characterized by epileptiform activity that recurs at short intervals without intervening physiological activity. Clinically this pattern of activity is similar to status epilepticus, which is defined as either a seizure lasting for more than a specified time period or seizures that recur without an intervening period of normal consciousness. There are serious clinical implications of the definition, because after 30 min, status epilepticus can become pharmacoresistant (Kapur and MacDonald). Moreover, prolonged experimental status epilepticus can cause considerable neuronal death [33]. The lack of agreement regarding the duration of seizure activity necessary for status epilepticus is related to our lack of knowledge regarding the time course of the damage to neurons.

*In vitro*, it turns out that shortly after their initiation, exposure to low Mg, 4-AP, high K and low Ca induce seizure like events that all respond well to standard AEDs. However, if the activity persists for some time, then seizure like events

gradually shorten, and ultimately short recurrent discharges occur, which are also unresponsive to standard AEDs [15, 23]. The analysis of such events indicates that the transition from long seizure-like events to short discharges is probably due to reduced GABAergic transmission. A variety of pre and postsynaptic processes may underlie this loss of efficacy, such as internalization of GABA receptors, consumption of GABA for synthesis of ATP, and alteration of the anionic transmembrane gradients that subserve GABA<sub>A</sub> receptor-mediated inhibition. The reduction in GABAergic function explains why GABAergic agents that prolong the GABAergic signaling lose efficacy during the course of prolonged seizures. However, in some cases, agents that directly activate GABA receptors are still effective [67]. In other situations GABAergic agents are either minimally ineffective or exacerbate epileptiform activity, a situation that can be improved by agents that improve the transmembrane anionic gradient [26, 27]. This improvement has also been observed in human case studies [47] and is being investigated in human trials. In some circumstances these additional processes that are dependent on the duration of seizure activity prior to drug application have provided the key to resolving seemingly contradictory results [1, 27, 28].

Neurons are depolarized during prolonged seizures, and so are their mitochondria. Brian Meldrum's experimental neuropathological studies of status epilepticus in the baboon focused attention on mitochondrial changes accompanying ictal neuronal cell death [33]. Subsequent studies have provided evidence that at least part of the neuronal damage arising during status epilepticus seems to be due to mitochondrial depolarization and increased production of free radicals [17, 54]. This suggests that some neuroprotection can also be achieved by free radical scavenging [54, 76]. This is an area that can be profitably studied *in vitro*, where microscopic imaging during epileptiform activity is more feasible than *in vivo*. Barbiturates and other anesthetics used to terminate status epilepticus are typically titrated to a burst suppression pattern that is very reminiscent of the periodic population

discharges that are observed in acute brain slice preparations exposed to convulsants with anesthetic concentrations of barbiturates [23]. Indeed, recurrent epileptiform discharges cause considerable cell loss due to mitochondrial depolarization and increased free radical production sensitive to neuroprotection by free radical scavengers such as tocopherol. Other anticonvulsant and neuroprotective strategies such as cooling of patients' brain by a few degrees or anticonvulsants which do not involve GABAergic signaling should continue to be investigated [75]. It will be important to advance these early results *in vitro* and then translate the results of these experimental findings into good clinical studies.

#### 2.4.1.5 Increased Seizure Threshold of Epileptic Tissue *In Vitro*

As mentioned above it is often difficult to induce seizure like events in tissue from animals with epilepsy acquired after drug-induced status epilepticus. This may reflect an endogenous anticonvulsant effect. It has not yet been described for kindled animals, and it depends on the number of seizures an animal has experienced [98]. In chronically epileptic human tissue resected for seizure control, it is even more difficult to evoke seizure like events [30, 39]. In resected hippocampal tissue, seizure-like events can often be induced by elevating potassium concentration in the dentate gyrus and subiculum [30]. In neocortex 4-AP can be employed but it works in only a subset of patient specimens [5]. We have also been able to induce seizure activity with high potassium combined with bicuculline. These observations raise important questions as to the mechanisms underlying relative seizure resistance in epileptic tissue.

Kindling is most effective when a critical interval is included between kindling stimuli. It was first suggested that this may relate to upregulation of opioid receptors [73]. Later it was shown that a single repetitive stimulation of the perforant path from the entorhinal cortex to the dentate gyrus could upregulate the GABA synthesizing enzyme GAD with subsequent co-release of GABA and glutamate from mossy fiber terminals which leads to an elevated seizure threshold [35].

These effects seemed to be transient and fade away with time. Additional evidence for endogenous antiepileptic processes come from slices prepared from kainate treated and also from pilocarpine treated animals, where the convulsant 4-AP was ineffective. This effect was due to up-regulation of the enzyme adenosine deaminase acting on RNA (ADAR2). This causes mRNA editing of AMPA type glutamate receptors as well as Kv1 potassium channels that lose some of their sensitivity to 4-AP [82]. An additional mechanism involves arachidonic acid which is directly blocking K channels [11] and in addition can be metabolized to a number of intrinsic convulsant or proconvulsant derivatives [44]. Activity dependent editing of alpha 3 subunits of glycine receptors has also been described. This editing leads to an increased affinity for glycine and some of its agonists [63]. Although such processes may decrease seizure susceptibility, we need to keep in mind that there is not sufficient circuitry in a slice for seizure generation. Thus we may need to pay more attention to network preservation when studying network phenomena such as seizures and epilepsy *in vitro*. Nevertheless, hypothesis driven searches for other anti-ictogenic mechanisms that are active in epileptic tissue comprise a promising route for discovering new treatments of pharmacoresistant epilepsies.

#### 2.4.1.6 Analysis of Proepileptogenic Factors

Another translational opportunity for epilepsy research is the *in vitro* study of mechanisms of epileptogenesis. Trauma and stroke research led to the important discovery that neuronal circuits reorganize following a brain lesion, and this had important implications for the study of epilepsy. For example, the observation of mossy fiber sprouting, that is sprouting of dentate granule cell axons back into the input layer of the dentate gyrus, has been a central model of the recurrent positive feedback that is a necessary component of any sustained network activity, including seizures [56, 65]. However, some investigators now wonder whether this neurocentric approach to the understanding of epilepsy may have been too

narrow. Many conditions which lead to epilepsy are associated with an open blood brain barrier [72, 86]. The immediate effects of blood brain barrier disturbances include vasogenic edema due to extravasation of albumin and other serum proteins into the brain interstitial space. This increases intracranial pressure, potentially reducing microperfusion. Opening of the blood brain barrier also increases extracellular potassium and reduces extracellular Ca and Mg concentrations, because these are lower in serum than in the brain interstitial space [79]. Activity-dependent increases in blood flow might not occur under these conditions. Thus if seizures emerge, relative metabolic deprivation may ensue. Seizures and metabolic deprivation lead to cell swelling, i.e. cytotoxic edema [21]. When the blood brain barrier is opened, albumin is absorbed into perivascular macrophages and astrocytes, perhaps reflecting an attempt to reduce the extracellular colloid pressure. This process is associated with activation of TGF $\beta$  receptors and subsequent activation of astrocytes, including increased expression of GFAP [40] and down regulation of K<sub>IR</sub> channels. This results in depolarization of astrocytes and changes in the expression of connexins, resulting in reduced astrocytic electrical coupling [16]. Both effects lead to enhanced accumulation of extracellular potassium and perhaps glutamate in the extracellular space. Under these conditions, seizure threshold is strongly reduced and when seizures develop they rapidly progress to spreading depression [55]. Preliminary evidence suggests also that these alterations in astrocyte properties may be associated with increased release of chemokines and cytokines and potentially also with release of gliotransmitters. Importantly these alterations precede appearance of seizures and if stopped may prevent later epileptogenesis. Probing for an open blood brain barrier may be an important biomarker for epileptogenesis following trauma, stroke and encephalitis and some form of tumors. However, not all tumors are associated with an open blood brain barrier and criteria that take into account constraints on the role of the open blood brain barrier in epileptogenesis have still to be evaluated.

### 2.4.2 Isolated Hippocampus and Isolated Brain

The isolated intact hippocampus has recently received considerable attention because a number of questions can be addressed that are of potential clinical relevance. One is the induction of a mirror focus by using the two hippocampi interconnected by commissural fibers [49]. Induction of seizure-like events in one hippocampus induced a seizure focus in the contralateral hippocampus without any additional pharmacological treatment. This is potentially important as it could explain why in some cases seizures do not stop when one hippocampus is removed. On the other hand this is an acute finding that may be more closely related to mechanisms underlying rapid kindling than the development of mirror foci in chronic epilepsy. The finding is limited to young age, as maintenance of the intact hippocampus beyond postnatal day 10 is presently not possible. For such studies in older age it may be more feasible to use preparations from turtles or birds. Another aspect of studies in juvenile intact hippocampus is that the evoked seizure like events seem to be resistant to clinically employed drugs [70], perhaps reflecting immature ion transport mechanisms [27]. This may therefore be a preparation in which new agents can be tested which specifically address seizures in babies and young infants.

Another intact *in vitro* preparation is the intact guinea pig isolated brain preparation [19]. It permits studies on long range interactions within the brain during seizure like events and indicates that seizure generation is based on multisite interaction in wide spread neuronal circuits. However it is apparently difficult to induce epilepsy in guinea pigs and the intact brain is difficult to prepare from aged animals.

### 2.4.3 Use of Human Tissue *In Vitro*

About 30 % of patients with epilepsy do not become seizure free with presently available drugs. Thus there remains a pressing need for models of pharmacoresistance that are correlated

with data from patients. At present, human tissue resected during epilepsy surgery is primarily used for diagnostic purposes. In past years however the neuropathology field has opened itself to molecular biology aspects concerning expression of peptides, transmitter receptors, ion channels, gene regulation and epigenetics [18]. Human tissue samples can to some extent also be used for determination of changes in interneuronal connectivity and in probing for alterations in astrocyte properties [20]. Moreover in human tissue spontaneous events may be detected that might resemble interictal spikes and fast ripple activity [51]. This may permit the study of mechanisms of fast ripple activity in human tissue. Interestingly slices prepared from human specimens often have a relatively long survival time. This might permit development of slice cultures from human tissue.

It is notoriously difficult to induce seizure like events in human tissue. As discussed above, this is probably due to upregulation of anti-ictogenic mechanisms, in addition to the effects of partial network disassembly. Endogenous protective mechanisms are of interest because studies of these mechanisms could lead to identification of novel anticonvulsant and antiepileptogenic therapies. However it is still possible to induce seizure like events in the hippocampus or temporal neocortex of TLE patients, and in the cortex of patients with developmental disorders. In the hippocampus the most effective method to induce seizures is elevation of potassium concentration. In temporal neocortex seizure-like events can be induced in a subset of preparations by 4-AP, or 4-AP combined with elevated potassium concentration. In our hands the best method for induction of seizure like events in temporal neocortex slices is the use of potassium elevation combined with application of bicuculline (unpublished observation). In studies of epileptiform activity induced by elevated potassium in the hippocampus it was noted that the slices do not respond to CBZ if they come from patients with pharmacoresistant epilepsy but do respond if they come from patients which are not resistant to AEDs such as tumor patients [41]. It is noteworthy that in some instances one slice from a pharmacoresistant

patient may not respond to AEDs while the other does. This heterogeneity offers itself for studies on mechanisms underlying pharmacoresistance. Obviously if one is able to induce seizure like events in slices from pharmacoresistant patients this opens the possibility to test for agents which might alleviate the epilepsy in drug resistant patients. One argument against this strategy is that the obtained material is too heterogenous and that in many centers the incidence of epilepsy surgery is too low to permit for rapid information. However monkey studies are often indeed based on many repeated measures in the same subject. The amount of human tissue available is often large and would permit to study effects of a multitude of agents on the same patient material if logistics can be surmounted. For example in analogy to multi center clinical studies, it might be possible to set up multicenter studies on resected material, although this might require new funding mechanisms.

#### **2.4.4 Slice Cultures as a Model of Traumatic Epilepsy**

##### **2.4.4.1 Ictogenesis**

There are a number of different techniques for preparing organotypic slice cultures from cortex or hippocampus. Their properties depend on the way they are fixed to the substrate material, on the age at preparation and on the media used for maintenance in culture [6, 85]. Most studies related to epileptogenesis are done on organotypic hippocampal slice cultures. Cortical organotypic cultures and hippocampal cultures maintained with B27 artificial media often display spontaneous seizures [2, 10] which can be recorded also while the cultures are in the incubator by different techniques such as MEAs or implanted electrodes. In this preparation, epileptogenesis proceeds at a rapid but predictable time course [25]. Interictal activity precedes the onset of ictal activity by several days. Status epilepticus commencing shortly after the appearance of spontaneous seizures is observable for hours to days [2]. Seizure-induced neuronal death is readily apparent, peaks during status epilepticus [54],

and can be prevented by standard anticonvulsants such as phenytoin [9]. The incidence of epilepsy is nearly 100 % in slice cultures from rats and mice, and in fact a current challenge is developing a culture system with a lower incidence of epilepsy that might make a better predictor of biomarkers and therapeutic agents for human epileptogenesis.

Some investigators prefer to induce seizure like events by lowering Mg thereby activating NMDA receptors, or application of bicuculline thereby reducing inhibition. Application of 4-AP leads to strongly enhanced transmitter release due to the strong expression of 4-AP sensitive Kv1.4 and 1.5 as well as some Kv3 channels on presynaptic terminals. Seizure like events are usually characterized by some initial clonic like discharges, followed by a tonic like and thereafter clonic like period followed by a postictal depression and the recurrence of interictal discharges. During seizure like events, ionic changes occur which mimic those observed during seizures in intact animals. If seizures recur with a high incidence they can convert into late recurrent discharges which are characterized by shorter events with synchronous intracellular depolarizations. Thus slice cultures offer themselves for studies on ictogenesis and factors which facilitate ictogenesis such as reorganization of the neuronal networks under study. Epileptogenesis can also be studied. For example typical epileptic circuitry with recurrent axon collaterals, back projection from CA1 to CA3 or DG can be observed [34, 56]. Slice cultures can be maintained for up to 8 weeks and therefore offer themselves also for long term observations. A drawback is that it is rather difficult to make slice cultures from hippocampal tissue beyond postnatal day 16. There are reports that slice cultures can be made after this date but the chances that these can be maintained for more than 2 weeks are rather slim and therefore very labor intensive [58].

##### **2.4.4.2 Pharmacosensitive vs. Pharmacoresistance**

Depending on duration of culturing and maintenance conditions, evoked seizures can be sensitive to

AEDs or insensitive. In the same slice culture both conditions can coexist: thus while 4-AP and low Mg induced seizure like events in some conditions are pharmacoresistant the seizure like events induced by repetitive stimulation are not [3, 4]. Therefore slice cultures can be used as a model of pharmacoresistant seizures and drugs can be tested which might be useful for the treatment of epilepsies in patients whose seizures cannot be satisfactorily controlled by present medication.

Spontaneously epileptic slice cultures that are not exposed to convulsants respond to anticonvulsants with suppression of ictal but not interictal activities, as is the case clinically [9]. Interestingly, dependent on culture conditions these cultures become resistant to anticonvulsants after 1–3 weeks of exposure, with recrudescence of seizure activity at anticonvulsant concentrations that completely suppress seizure activity in naïve slices of the same age [3, 9]. Thus the organotypic slice culture is a promising tool for the investigation of the phenomenology and pathophysiology of pharmacoresistance.

#### **2.4.4.3 Mechanisms of Ictogenic Cell Death**

Slice cultures offer themselves also for studies on ictogenic cell death. A number of methods are available to monitor cell death. These include the measurements of LDH in the supernatant and also of propidium iodide staining and ethidium bromide staining [9, 54]. Of course it is also possible to test for programmed cell death. One approach is to perform experiments with reduced oxygen supply in slice cultures that are generating stimulation-induced seizure like events. These events develop into spreading depolarization which when oxygen tensions falls to near zero cause cumulative cell death, a situation which is similarly observed also in slices from animals which experienced a stroke [68]. On the other hand with normoxic or hyperoxic perfusion it can be shown that seizure like events are associated with increased free radical production and eventually damage of mitochondria leading to disturbances in the coupling of neuronal and metabolic activity causing cell death because of

lack of sufficient ATP supply. Buffering ROS by different means can be shown to be highly neuroprotective.

Another approach to studying cell death in spontaneously epileptic slice cultures is to assay release of lactate dehydrogenase (LDH) into the culture media, which is changed twice weekly [9, 32]. This is a simple procedure that while not linearly related to cell death, provides a rapid and reliable means to assay cell death in higher-throughput experiments in which toxicity of screened agents and prevention of ictal cell death are important endpoints. More detailed studies of ictal cell death employ either exogenous markers such as propidium iodide, or endogenously expressed fluorescent markers of caspase activation. These studies provide a means to follow cell death over time, and to ask important questions as to the activities and features that precede or predict death in identified neurons.

#### **2.4.4.4 Slice Cultures: A Model of Post-traumatic Epilepsy**

When slice cultures are prepared a large number of connectivities are severed leading to some extent to retrograde degeneration but also to transformation of a three dimensional organization into a two dimensional organization. Thus the slice culture can be considered to comprise a model of (pediatric) traumatic brain injury. When spontaneous seizures emerge in these cultures they can be used for long term monitoring of drug effects thus facilitating detection of changes in efficacy of a given drug. This includes also detection of toxic side effects with nervous tissue [8, 9].

#### **2.4.4.5 Long Term Monitoring of Anti Epileptogenic Effects**

Slice cultures can also be used to study antiepileptogenic strategies. One example is neovascularisation. The density of blood vessels in human and chronic epileptic rodent tissue is often remarkably increased [72]. This makes it possible to address the question as to whether neof ormation of blood vessels can be altered [64]. Surprisingly slice cultures present with many blood vessels which are usually equipped with a



tight blood brain barrier [53]. Most of these vessels remain intact unless there is infection in the tissue. Therefore slice cultures can be used to determine effects of microglial activation on vessel density, and also whether factors that prevent revascularization have neurotoxic effects that might interfere with epileptogenesis.

Slice cultures permit study of signaling cascades and of factors that may serve as antiepileptogenic factors [8]. These can be neuroprotective agents, for example blockers of signaling cascades such as the TGF $\beta$  activated pathways or the mTOR pathways, and agents that interfere with neuronal survival or growth factors. At present most of these strategies are not yet ready for transfer into clinical trials, but this area is a promising area for further *in vitro* and *in vivo* study. Slice cultures are most useful to study drugs whose effects require time to produce anti-seizure or anti-epileptogenic effects. Most drug testing assays used *in vivo* or *in vitro* test for very acute effects although many treatments in psychiatry and epileptogenesis take time to take full efficacy.

#### 2.4.4.6 Use of Transgenic Models of Epileptic Encephalopathies

Many transgenic mice display seizures. Murine models have been developed for several human mutations that cause severe childhood epileptic encephalopathies. In many instances the transgenic models do not live long enough for research on the precise pathogenic cascade. However slice cultures can be prepared from ages ranging from fetal tissue to P16–18. Preparing slice cultures from such animals offers the possibility to look into the precise pathophysiological cascade and to define intervention points by which the epileptogenesis can be prevented. For these studies, the development of chronic slice cultures that do not become epileptic except in the presence of the targeted gene defect would be very useful. However, in the absence of such a slice preparation, the organotypic slice can still be of exceptional utility. For example, slice cultures prepared from transgenic animals expressing cell-type-specific fluorophores that are activated by particular ions, neurotransmitters, or second

messengers, or by cell-type-specific expression of light-sensitive rhodopsins can be studied with targeted path scanning multiphoton microscopy and activity-dependent fluorophores. This provides the means to precisely interrogate critical network elements that are active during ictogenesis and epileptogenesis.

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## 2.5 Some Conclusions

The above discussion is not intended to be a thorough review of epileptogenesis. We tried to illustrate some of the successes and challenges of *in vitro* preparations for translational research. *In vitro* preparations offer a large number of research possibilities to address clinical questions and therapeutic options, including long term observations in slice cultures, detailed cellular analysis, imaging and optogenetic studies, and expression of orphan receptors that permit activation and silencing of select populations. Expression and suppression of specific RNA and proteins can be achieved semi-acutely or chronically. All these technologies can now be employed for studies on ictogenesis, epileptogenesis and aspects of disease such as signaling cascades, development of pharmacoresistance, and neuroprotection. Exploiting the multiple technical possibilities for translational research will be substantially enhanced by improved contact between clinicians and scientists. This would culminate in clinical research executed through coordinated multicenter trials where promising, robust preclinical observations could be readily transformed into clinical proof of principle studies.

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# What Is the Importance of Abnormal “Background” Activity in Seizure Generation?

# 3

Richard J. Staba and Gregory A. Worrell

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## Abstract

Investigations of interictal epileptiform spikes and seizures have played a central role in the study of epilepsy. The background EEG activity, however, has received less attention. In this chapter we discuss the characteristic features of the background activity of the brain when individuals are at rest and awake (resting wake) and during sleep. The characteristic rhythms of the background EEG are presented, and the presence of  $1/f^\beta$  behavior of the EEG power spectral density is discussed and its possible origin and functional significance. The interictal EEG findings of focal epilepsy and the impact of interictal epileptiform spikes on cognition are also discussed.

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## Keywords

Electroencephalogram • Epilepsy • Local Field Potential Oscillations • High Frequency Oscillations • Sleep

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## 3.1 Introduction

The electrical activity of mammalian brain, defined by the electroencephalogram (EEG), has long been a focus of scientific and clinical brain research [16]. The mechanisms underlying various EEG changes associated with brain maturation, behavioral states,

cognition, motor function, and neurological disease represent fundamental discoveries of neuroscience. Epilepsy in particular has benefited from EEG investigations [15]. A disorder characterized by unprovoked recurrent seizures, epilepsy has many underlying pathological causes but is unified by the common clinical expression of seizures and the

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associated pathological brain electrical activity. Not long after the discovery of the human EEG [7], Berger also reported that epileptic seizures had an abnormal EEG signature, and that between the seizures (interictal) there were also transient epileptiform abnormalities not seen in controls (translated in [8]). Thereafter, the significance of interictal epileptiform spikes (IIS) and abnormal transient oscillatory network activity in the development of epilepsy (epileptogenesis), seizure generation (ictogenesis), and associated functional impairments (e.g., cognition, memory, and reaction times) have been active areas of research.

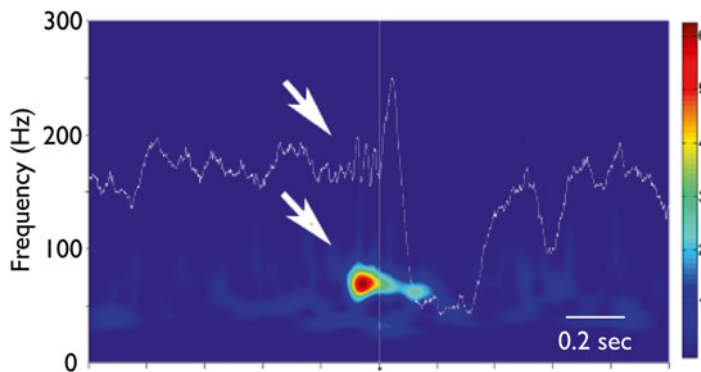
### 3.2 Physiological Electrical Activity in the Normal Mammalian Brain

Since the first observation of the occipital alpha rhythm [7] (translated in [8]) the interest in brain oscillations and their physiological and pathological correlates has occupied a central position in human neuroscience. Historically clinical and basic research focused on specific oscillations that are prominent in the EEG intermittently, for example the occipital alpha rhythm ( $\alpha$ ; 8–12 Hz) recorded at rest with eyes closed, beta ( $\beta$ ; 12–30 Hz) and gamma frequency activity ( $\gamma$ ; 30–50 Hz) during mental and motor tasks, theta frequency

oscillations ( $\theta$ ; 4–8 Hz) during memory tasks or sleep, and delta frequency activity ( $\delta$ ; 0.5–4 Hz) that characterizes slow wave sleep. Similarly, the EEG activity in traditional frequency bands ( $\delta$ ,  $\theta$ ,  $\alpha$ ,  $\beta$ ,  $\gamma$ ) became the focus of EEG research and intensively studied in brain maturation [50], normal function, and disease states. However, it is widely recognized that the brain generates activities well outside these classic EEG bands. In fact, the high amplitude EEG activity below  $\delta$  (<0.5 Hz) including direct current (DC) changes were some of the earliest electrical activities recorded [2].

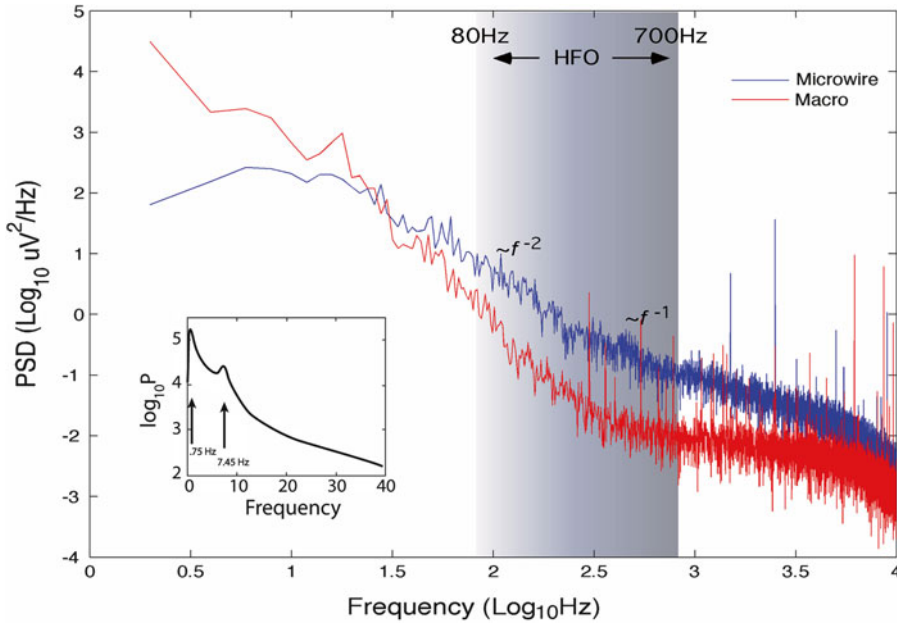
While EEG research has largely focused on narrow band EEG oscillations ( $\delta$ ,  $\theta$ ,  $\alpha$ ,  $\beta$ ,  $\gamma$ ) it is well recognized that there is a broad spectrum of on-going, arrhythmic, background activity that does not contain a dominant characteristic oscillation, but rather is composed of intermixed spectral frequencies [17, 34]. It is out of this arrhythmic background that the traditional EEG oscillations discussed above, e.g. the posterior dominant alpha rhythm, may be evoked or spontaneously emerge (Fig. 3.1). More recently, this broad spectrum of on-going background activity has been a focus of attention [17, 34, 43].

This composition of background electrical activity in mammalian brain generally follows power-law behavior, i.e. the spectral power scales with frequency as  $1/f^\beta$  where  $\beta$  is called a scaling exponent (Fig. 3.2). This  $1/f^\beta$  (“one-over- $f$ ”)



**Fig. 3.1** Interictal Background Activity from human hippocampus recorded with intracranial depth electrode. There is an ongoing background activity followed by a paroxysmal gamma frequency oscillation (*bold arrows*), and an interictal epileptiform spike (IIS, *arrow*). The wavelet transform (1–600 Hz, Morlet basis) spectrogram

of raw intracranial EEG shows the background theta activity, and the emergent low amplitude gamma oscillation preceding the IIS. The scale bar is normalized units standard deviations from background. Time base is 100 msec per division (Courtesy of Liankun Ren, M.D. unpublished)



**Fig. 3.2** Power spectral density (PSD) from 5 min of human hippocampus during sleep recorded with intracranial EEG (0.05–10,000 Hz, sampled at 32 kHz) using micro- and clinical macro-electrodes. The wide bandwidth recording exhibits  $1/f^\beta$  behavior with different scaling regions characterized by different scaling exponents  $\beta=1$

and 2. The inset shows an expansion of the PSD in the 0.05–40 Hz range and spectral peaks from a low delta frequency oscillation ( $\sim 0.75$  Hz) and a theta-alpha frequency oscillation (7.45 Hz). The characteristic 0.75 Hz oscillation is persistent throughout the 5 min and modulates the 7.45 Hz intermittent oscillation (Unpublished Data)

spectrum with lower frequency activities exhibiting higher amplitudes than faster frequencies is characterized by the scaling exponent  $\beta$  that can be obtained by plotting log power vs. log frequency ( $\log(\text{Power})$  v.s.  $-\beta \log(f)$ ) and ranges over  $0 < \beta < 4$  [31, 34, 52, 80]. The spectral peaks embedded in the  $1/f^\beta$  represent ongoing persistent oscillations or organized emergent oscillations that arise out of the ongoing EEG background, such as the traditional EEG rhythms of the human EEG (Fig. 3.2).

The arrhythmic background activity has more recently received attention within the context of the advancing understanding of complex systems. It is recognized that  $1/f^\beta$  behavior in complex systems can be a signature of a self-organized system with scale-free dynamics [31, 34, 55]. It turns out that  $1/f^\beta$  patterns are ubiquitous in nature, from the statistics of earthquakes to stock market dynamics [4]. The origin of  $1/f^\beta$  behavior in EEG and local field potential (LFP) recordings remains unclear [9, 10, 34], but perhaps one of the most intriguing ideas is that it results from hierarchal

nesting of brain activity [19, 34, 73] whereby lower frequency activity modulates higher frequency activity [34]. The modulation of gamma oscillations by theta oscillations is a classic example [6, 14, 18]. At the cellular level, multi-unit activity is correlated with EEG gamma power and phase-locked to the negative-going phase of the delta frequency activity [79]. Synchronization between neuronal assemblies also occurs within arrhythmic brain activity [25, 45, 70].

*Maturation of EEG:* The continuous maturation of EEG activity through young adulthood reflects brain development, e.g. myelination, and organization [50]. In premature infants (24–27 weeks), the EEG is discontinuous and may alternate between periods containing bursts of high amplitude slow (0.1–1 Hz) activity and intermixed faster rhythms (8–14 Hz). From these earliest electrical rhythms in the infant brain there are long periods of continuous development through late childhood ( $\sim 12$  y.o) when the posterior dominant alpha rhythm reaches  $\sim 10$  Hz [50].



### 3.3 Electrical Activity of the Sleeping Brain

There exists substantial evidence for the physiological importance of sleep and in particular the requirement of sleep for normal memory [24]. To better understand how memory benefits from sleep, it would be helpful to first describe briefly the EEG during the two main types of sleep – rapid eye movement (REM) and non-REM sleep – and then how the neurophysiology of sleep might support aspects of memory formation. Since patients with epilepsy often report deficits in sleep and impairment in memory, subsequent sections describe electrophysiological disturbances in the epileptic brain and their likely functional implications for cognition.

*EEG of REM sleep:* During REM or desynchronized sleep, arising from a background of low-voltage, mixed frequency EEG, are spontaneous synchronous bursts of neuronal activity generated by the pontine tegmentum that spread to the lateral geniculate nucleus and visual cortices in the occipital lobe that are termed “PGO waves”. Conspicuous in the EEG of rats and cats and less in humans, PGO waves coincide with rapid eye movements and can become phase-locked with theta oscillations. In rodents, theta oscillations occur with largest amplitude in the hippocampal CA1 area driven by inputs from septum, entorhinal cortex, and CA3. In addition to REM sleep, hippocampal theta can also be observed during awake behaviors in rodents. Theta also occurs in humans during wakefulness, but is more apparent in neocortical areas and less coherent in hippocampal areas.

*EEG of non-REM sleep:* Non-REM sleep is characterized by high-voltage slow wave activity that includes slow oscillations <1 Hz and delta activity. The slow oscillation persists in isolated neocortical tissue and is abolished if thalamocortical cells are deafferented from cortical inputs, suggesting slow oscillations are generated largely within neocortex [60, 71]. In scalp EEG, the alternating sequence of surface positive (depth negative) and negative (depth positive) waves correspond with periods of neuronal membrane

depolarization and hyperpolarization respectively. Periods of membrane depolarization occur within excitatory and inhibitory cells that produces sustained neuronal firing commonly referred to as “UP-states”, whereas periods of membrane hyperpolarization are accompanied by neuronal silence denoted as “DOWN-states”. The mechanisms generating slow oscillations are not yet clear, although evidence to date suggests UP-states could arise from widespread summation of calcium- and persistent sodium inward current-mediated excitatory postsynaptic potentials in cortical cells, while neuronal disfacilitation associated with DOWN-states could be due to calcium- and sodium-dependent potassium currents, inactivation of persistent sodium currents, and possibly GABA-mediated inhibition.

Slow oscillations strongly modulate two other transient oscillations that occur during non-REM sleep – spindles and sharp wave-ripple complexes – and is another classic example of frequency nesting. Spindle waves are beta frequency oscillations that wax and wane between 10 and 16 Hz and last 0.5–2 s that characterize stage 2 of NREM sleep. Spindles arise from interactions between GABA-containing neurons in the thalamic reticular nucleus as well as thalamocortical cells that facilitate the synchrony and spread of spindles throughout neocortex. Human studies have identified two types of spindles designated slow and fast; however, whether these two types of spindles arise from different neuronal mechanisms or reflect the modulation of a common spindle generator is not known. Slow (10–12 Hz) spindles occur primarily over frontal cortical areas and more frequently during slow wave sleep than stage 2 sleep, and fast (13–15 Hz) spindles appear broadly over central and parietal cortices and are often coincident with increased hippocampal activity.

In hippocampus during non-REM sleep, spontaneous extra-hippocampal impulses drive neuronal firing in CA3 that projects forward via Schaffer collaterals onto dendritic processes of CA1 pyramidal cells and some types of interneurons. This briefly irregular (30–120 milliseconds in duration), increase in neuronal firing registers in the depth EEG as a large amplitude sharp wave with maximum negativity in stratum lucidum,

stratum radiatum, and inner third of stratum moleculare corresponding to input layers of CA3, CA1 and dentate gyrus respectively. In CA1, a similarly brief high-frequency oscillation (HFO; 80–200 Hz) termed “ripple” arises from synchronous firing between pyramidal cells and basket cells that is largest in amplitude in stratum pyramidale and superimposed on the sharp wave. During widespread neuronal depolarization associated with the slow oscillation UP-state, hippocampal ripples can co-occur with neocortical fast spindles to form spindle-ripple events with ripples that temporally coincide with the troughs of spindle waves [64].

*Concept of memory function and putative neuronal mechanisms:* Memory function generally involves processes of encoding, consolidation, and retrieval. In the awake brain, encoding occurs when perception of the stimuli produces a new, yet unstable, memory trace. During subsequent sleep, the labile memory trace becomes more stable and eventually integrated into brain networks supporting long-term storage of knowledge in a process termed “consolidation”. During retrieval, the stored memory is accessed and recalled. A number of theories have been proposed on how sleep supports memory consolidation with some more than others supported by compelling data from animal and human studies (for extensive review, see [56]). Central to current theories (e.g., “active system consolidation”) is the concept of reactivation that involves a sleep-related replay of neuronal firing patterns corresponding to the neuronal firing patterns that occurred while encoding, i.e., during prior wakefulness, as well as specific roles for different types of sleep in memory consolidation.

Considerable research has focused on identifying the neuronal mechanisms that provide sleep-related benefits on memory formation. Current models emphasize precisely coordinated neuronal activity between neocortex and hippocampus for hippocampal-dependent memories [26]. During non-REM sleep, neocortical slow oscillation UP-states provide a temporal window for increased ripple activity and associated increase in neuronal firing that could reflect reactivation of hippocampal memories. The coincidence of hippocampal ripples with neocortical

fast spindles (spindle-ripple events) is thought to promote the transfer and ultimately storage of the hippocampal memory to neocortex [64], which is reflected presumably by long-term functional and structural changes that strengthen synaptic transmission (e.g., long-term potentiation). In addition, evidence suggests REM sleep PGO- and theta-related neuronal activity could also be involved with synaptic modifications with theta possibly playing a role in synaptic downscaling, which extends the “synaptic homeostasis” hypothesis that links the regulation of sleep with mechanisms of synaptic plasticity [72].

*Human single neuron correlates of sleep and memory:* Microelectrode unit recordings during natural sleep in humans are few, but available data indicate hippocampal neuronal firing increases during non-REM sleep and declines during REM sleep [57, 65–67]. Furthermore, the propensity for burst discharge is highest during non-REM sleep compared to awake and REM sleep episodes. These results are similar to the rates and pattern of hippocampal pyramidal cell firing during non-REM and REM sleep in rodents [64], and are generally consistent with levels of hippocampal activity that could be involved with reactivation described in the preceding paragraphs. Work using the same microelectrode recordings from single neurons in humans has primarily focused on memory and navigation. These studies have led to the discovery of place cells in the human mesial temporal lobe underlying spatial navigation [26], which resembles the location-specific firing patterns of some pyramidal cells in non-primate hippocampus described in the sections that follow. In addition, studies in humans have found evidence for neurons that encode category specific images [40, 41].

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### 3.4 Abnormal Electrical Activity in the Epileptic Brain

In addition to sleep and wake behavioral states of normal brain, epileptic brain is characterized by interictal state (between seizures), ictal state (seizures), and post-ictal states (after the seizure). It should be noted that while seizures are generally limited to a minute or so, the post-ictal state

as determined by subtle EEG or cognitive and physical changes can be prolonged [30]. In addition to the interictal and post-ictal state, there is emerging evidence for a pre-ictal state that is associated with increase in probability of seizure occurrence [22, 28, 47].

*Interictal Epileptiform Discharges:* Interictal EEG spikes (IIS) are brief, sharply contoured voltage fluctuations of less than 200 msec that are a signature of epileptic brain. The intracellular correlate of IIS is the paroxysmal depolarizing shift [3] seen in the neuronal membrane potential and is associated paroxysmal burst of neuronal population firing, but also involves a more complex interaction of inhibitory and excitatory neurons [3]. Depth electrode recordings during overnight polysomnographic sleep studies show that in patients with temporal lobe epilepsy (TLE), the highest rates of IIS regularly occur during non-REM stage 3, and in some cases stages 1 and 2, sleep compared to waking and REM sleep [42, 59]. In addition, the spatial distribution of IIS is often broader during non-REM sleep than waking or REM sleep, i.e., IIS appear at electrode recording sites within and remote from where seizures begin [59].

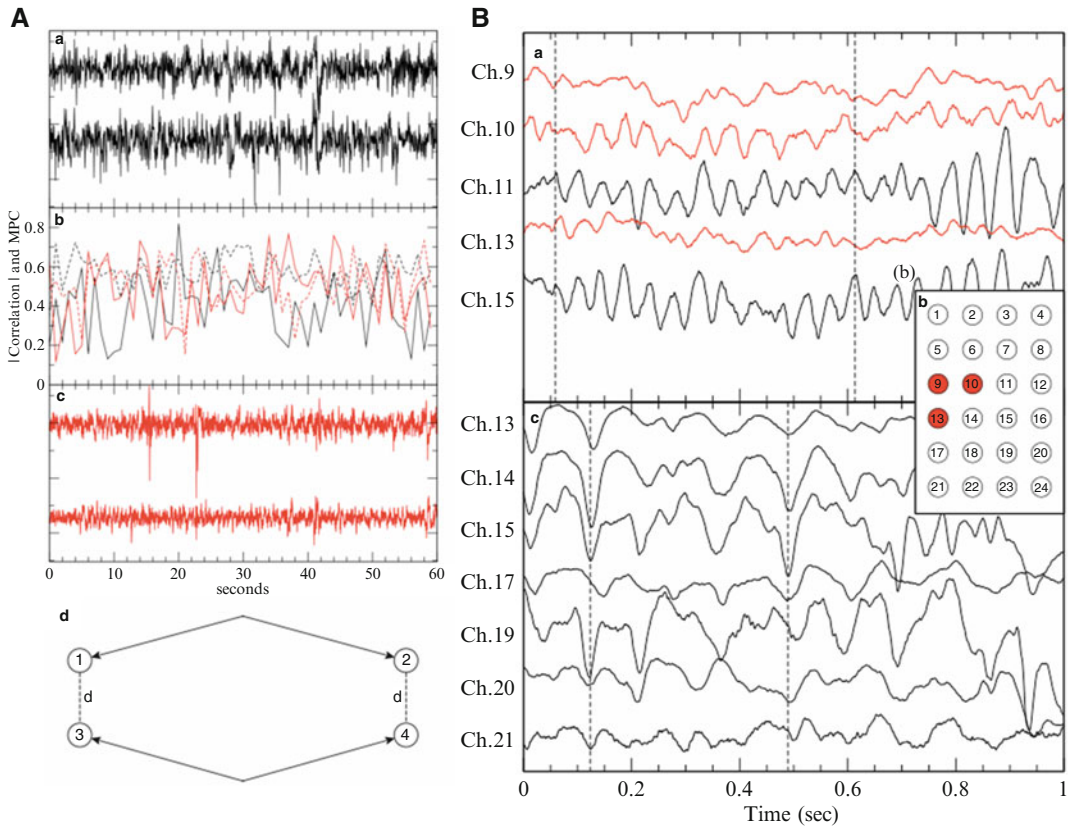
At the level of single neurons, patient studies have not consistently found differences in interictal firing rates and bursting inside versus outside the seizure onset zone (SOZ) during awake episodes [20, 21]. However, during non-REM and REM sleep compared to wakefulness, interictal firing rates, bursting, and synchrony of discharges are significantly higher in mesial temporal lobe (MTL) ipsilateral to the SOZ than contralateral MTL [65–67]. These results provide evidence for sleep-related facilitation of interictal neuronal hyperexcitability within the SOZ of patients with temporal lobe epilepsy.

*Pathological HFO:* In the epileptic brain, transient abnormally synchronous discharges of principal cells can summate in the extracellular space that give rise to a burst of population spikes commonly termed pathological HFO or pHFO [12, 13]. Chronic animal models of epilepsy and studies in patients with epilepsy indicate hippocampal and neocortical pHFOs are strongly

associated with brain areas capable of generating spontaneous seizures [29]. With respect to the wake-sleep cycle, recordings in epileptic rats and patients show the highest rates of hippocampal pHFOs occur during non-REM sleep compared to awake and REM sleep, while equivalent rates can be found during the latter two desynchronized EEG states [65–67]. By contrast, ripples are highest during non-REM sleep, while rates are lower during wakefulness and lowest in REM sleep, which is consistent with their occurrence in the normal non-primate hippocampus [65–67].

*Pathological Synchrony:* Synchronization of neuronal assemblies is thought to underlie normal brain functions such as perception, learning, and cognition. Alterations in neuronal synchrony are thought to underlie the clinical manifestations of many neurological diseases. Hypersynchrony of pathological neuronal assemblies as the generator of epileptiform activity has been a central theme of epileptic brain electrophysiology [53]. Jasper and Penfield speculated that the local high amplitude interictal epileptiform activity recorded directly from human cortex during surgery was generated by a burst of hypersynchronous neuronal activity [53]. Interestingly, however, many seizures appear to begin with an apparent “asynchronous state” – low amplitude LFP activity that evolves into a hypersynchronous state with high amplitude rhythmic activity [53]. Multiple studies have reported increased local synchrony, i.e. hypersynchrony, within epileptic brain using a range of quantitative measures of synchrony, including spectral coherence [74], magnitude squared coherence [81], and mean phase coherence [61]. In addition, investigations of LFP synchrony during spontaneous human seizures have consistently demonstrated a decrease in local LFP synchrony at seizure onset compared to baseline [48, 62, 78].

Analysis of long records of interictal iEEG from patients with focal epilepsy and control subjects with intractable facial pain found that the spatial distribution of LFP synchronization fell rapidly with the distance between electrodes [77]. Consistent with the hypothesis that the generators of normal and pathological HFOs are more spatially localized than lower frequency oscillations,



**Fig. 3.3** Data and from 2 patients, one with intractable facial pain (*black*) and other with focal epilepsy (*gray*). Data from the patient with intractable facial pain and no history of seizures serves as a control recording for quantitative comparison. **(A)** **(a)** Sample signals from two electrodes of the control brain recording. **(b)** The correlation magnitudes (*solid lines*) and mean phase coherence (*dashed lines*) of the signal pairs in **(a)** and **(c)**. Both the correlation and mean phase coherence (MPC) show significant temporal variability over the course of 60 s, with values primarily ranging from (0.2–0.7) **(c)** Sample signals from two electrodes in epileptic cortex outside the seizure onset zone. **(d)** A sample layout of the bipolar reference pair measurement. The dis-

tance  $d$  between one corresponding pair of electrodes 1 and 3 is equal to the distance between the other pair, electrodes 2 and 4, and this is the distance referenced in our bipolar measurements. **(B)** **(a)** Sample interictal iEEG signals from Patient 1 with epilepsy from both inside the seizure onset zone (SOZ), shown in *gray*, and near signals outside the SOZ (*black*). *Dashed line* marks significant phase lag between inside and outside the SOZ. **(b)** Spatial layout of the intracranial electrodes for Patients 1 and A with the SOZ electrodes (9, 10 and 13) of Patient 1 shown in red. **(c)** Sample signals from the control Patient A. The spatial numbering is as shown in **(b)**. For clarification, signals are offset vertically (Reproduced from Ref. [77])

the synchrony fall off is frequency dependent [44]. Synchrony in the epileptic brain, however, was shown to be markedly reduced in electrodes bridging connections between the SOZ and surrounding brain (Fig. 3.3). In effect, the SOZ is functionally disconnected and isolated from surrounding brain regions [77].

**Focal EEG Slowing:** In addition to the IIS and pHFO that have been widely investigated, focal slowing on the EEG is common in the region of

epileptic brain. Focal delta frequency slowing was initially described in patients with focal structural abnormalities, such as tumors and strokes [33, 76], but is also common in TLE [11]. When the slowing occurs as intermittent oscillations of monomorphic delta activity in the temporal lobe region it is termed, temporal intermittent rhythmic delta activity and is a signature of focal epilepsy [58, 68]. Focal delta frequency slowing has also recently been shown to

be more common than IIS following febrile status epilepticus [51].

*Hypsarrhythmia:* Hypsarrhythmia is the an EEG pattern characterized by disorganized high-amplitude spikes and spike- and/or sharp-slow wave discharges and commonly observed in infants with West syndrome (Infantile spasms, Hypsarrhythmia, and Developmental regression) (Fig. 3.4). Hypsarrhythmia is modulated by the sleep-wake cycle. During non-REM sleep, there is a tendency for runs of these high-amplitude discharges to become grouped or clustered with a period consistent with non-REM slow wave activity that typically is then followed by episodes of EEG attenuation [32]. By contrast, during REM sleep, there is a significant reduction and in some cases disappearance of hypsarrhythmia that reappears toward the end of the REM sleep episode, and if awakening then hypsarrhythmia continues often with clinical manifestations.

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### 3.5 IIS Impact on Cognition

While epilepsy in general could have detrimental effects on sleep, sleep-related epileptiform activity in particular could contribute to cognitive impairment [27, 39]. Epileptic encephalopathies refer to conditions in some patients with epilepsy who have neurological deterioration associated with frequency or severity of seizures and/or interictal epileptiform discharges and not due to the original cause or etiology [49]. Hypsarrhythmia in West syndrome and Electrical Status Epilepticus During Sleep associated with continuous spike-wave of sleep and Landau-Kleffner syndrome reflect interictal EEG patterns that predominate during non-REM sleep that could support abnormal activity-dependent synaptic plasticity which in turn produce cognitive impairments. Improved control of seizures and interictal EEG in these patients is often associated with improved cognitive performance that suggests ictal and interictal discharges contribute to these deficits [35]. Indeed, studies in adults and children show IIS can be associated with brief episodes of impaired cognitive functioning

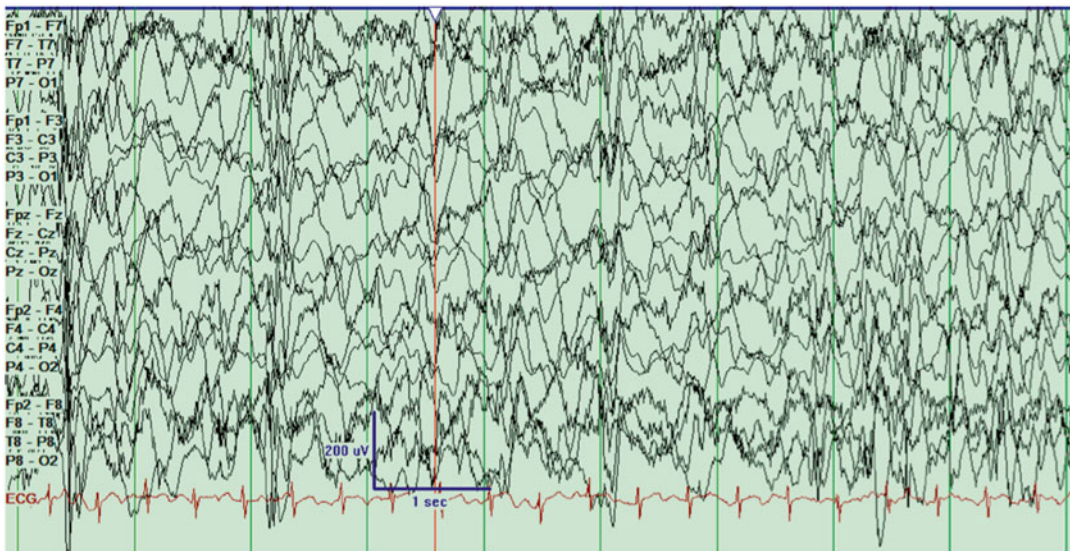
referred to as “transient cognitive impairment” [1]. The functional disruption coincides with the location where the IIS occur, e.g., verbal memory task impairment with left-side IIS, spatial task impairment with right-sided interictal epileptiform discharges (IEDs). Other studies observed IIS in occipital cortices disrupted visual stimuli presented in the contralateral visual field [63]. Work using pilocarpine-treated epileptic rats showed that in a hippocampal-dependent memory task, hippocampal IIS occurring during memory retrieval, but not encoding or maintenance, reduced performance [37, 38]. Similar results were observed in presurgical patients during a short-term memory task [38]. In this study, when depth electrode-recorded hippocampal IIS contralateral to SOZ or bilateral occurred during memory retrieval or maintenance, performance was lower. Recent evidence implicates brain network activity underlying memory processing in suppression of the IEDs, however, which raises an important confound of the interplay between cognitive processing on IIS [46].

How IIS disrupt memory processes is not known, but studies in epileptic rats show hippocampal IIS are associated with a decrease in CA1 cell firing [82]. Moreover, a significant reduction in firing was found after compared to before the IIS in interneurons, but not CA1 pyramidal cells that preferentially discharge when the rat is a specific location in the environment (“place cell”). A separate study of CA3 cell firing found significantly lower firing rates in interneurons and pyramidal cells before and after IIS compare to rates during random episodes [82]. These data indicate that neuronal firing surrounding and particularly after IIS could reflect episodes associated with reduced activity-dependent synaptic plasticity. By contrast, spontaneous pHFOs reflect brief episodes of increased neuronal discharges that are two-fold greater or more than the level of activity that occurs during spontaneous hippocampal ripples. Unlike ripples that involves interneuron-mediated regulation of pyramidal cell firing, it appears that the effects of interneurons are diminished during pHFOs. The abnormally high spatial and temporal coincidence of discharges could contribute to

pathological synaptic plasticity that is functionally disruptive to memory consolidation. One study of resective sclerotic hippocampal tissue, which is often associated with pHFOs and hyperexcitability in patients with drug-resistant temporal lobe epilepsy, found significantly lower levels of long-term potentiation and its synaptic counterpart long-term depression in sclerotic compared to non-sclerotic tissue [5]. These data suggest morphological alterations associated with hippocampal sclerosis and persistent abnormal interictal activity contribute to diminished capacity for physiological activity-dependent synaptic plasticity.

Disruptions in sleep and sleep-related oscillatory activity could interfere with aspects of memory formation. Patients with epilepsy are two times more likely to complain of sleep disturbances, chiefly excessive daytime sleepiness and insomnia that can have a negative influence on quality of life measures [23, 54]. Studies indicate seizures, comorbidity (e.g., sleep apnea),

and anti-seizure drugs can cause disruptions in the amount and architecture of sleep. During nights with nocturnal seizures, patients often have a greater number of nighttime stage-shifts or awakenings, increased amounts of non-REM stage 1 sleep, and reduced amounts of non-REM stage 3 sleep and REM sleep [72]. Daytime seizures, which themselves could prevent or interfere with learning, also reduce the amount of REM sleep during the subsequent night [72]. Furthermore, older types of anti-seizure drugs, such as barbiturates and benzodiazepines, can reduce the amount of REM sleep and in some cases stage 3 non-REM sleep. Newer drugs have no effect or can even increase the amount of REM sleep, although some reduce amounts of non-REM sleep (e.g., Gabapentin, Lamotrigine). Loss of slow wave-rich non-REM sleep or REM sleep in terms of absolute amounts and relative to when daytime learning occurs could negate the time-dependent benefits of sleep on memory formation.



**Fig. 3.4** The pattern of hypsarrhythmia is a specific EEG pattern associated with West Syndrome. First described in detail by Gibbs (66) the pattern consists of high voltage, disorganized EEG with multifocal and generalized epileptiform spikes and sharp waves. The characteristic pattern, often described as disorganized or chaotic, is unique in that the normal pattern of spatial synchrony over multiple brain regions is absent. The EEG tracing from each chan-

nel (e.g. Fp1 and F7) appear independent of each other despite the anatomical proximity. This is distinct from normal brain activity where there is widespread synchronous activity over multiple brain regions. In the hypsarrhythmia pattern the periods of generalized synchrony are due to generalized epileptiform discharges. The epileptiform spikes fluctuate in time and space, and are various focal, multifocal, and generalized

### 3.6 Conclusions

*Is the “background” activity important?* The answer to this question is clearly yes. There is good evidence that the background EEG activity contains important information about brain function and dysfunction in human epilepsy. While the diagnostic importance of IIS and seizures is clear, there is also evidence that the background EEG during sleep and wake is important prognostic tool. In addition, investigations of LFP synchrony and neuronal assemblies are providing mechanistic insights about brain function, cognition, and epilepsy related comorbidities.

*Is interictal background really “normal”?* In some epilepsy cases there are clear EEG background abnormalities. In West syndrome, Electrical Status Epilepticus in Slow-wave Sleep and other epileptic encephalopathies the EEG background is markedly abnormal. Whether there are more subtle abnormalities in the background in primary generalized epilepsy is an area of active study, but often the EEG on visual review appears normal. In focal epilepsy there may be focal slowing in the region of epileptic brain. In drug resistant epilepsy there are abnormalities in LFP synchrony that are present even in the absence of IIS.

Our cognition and behavior rely upon precisely-timed interactions among neurons forming brain networks by coordinated activity of anatomically distributed neuronal networks mediated via rhythmic brain oscillations [75]. Research into the common cognitive [27] and behavioral [36, 69] comorbidities of epilepsy is only beginning to emerge. As our understanding of the cellular mechanisms of cognition and behavior advance the impact on understanding the impact on epilepsy comorbidities should be significant.

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### Abstract

The seizure focus is the site in the brain from which the seizure originated and is most likely equivalent to the epileptogenic zone, defined as the area of cerebral cortex indispensable for the generation of clinical seizures. The boundaries of this region cannot be defined at present by any diagnostic test. Imaging and EEG recording can define regions of functional deficit during the interictal period, regions that generate interictal spikes, regions responsible for the ictal symptoms, regions from which the seizure is triggered, and regions of structural damage. However, these regions define the epileptogenic zone only when they are spatially concordant. The frequent discrepancies suggest the essential involvement of synaptically connected regions, that is a distributive focus, in the origination of most seizures. Here we review supporting evidence from animal studies and studies of persons undergoing surgical resection for medically-intractable epilepsy. We conclude that very few of the common seizures are truly local, but rather depend on nodal interactions that permit spontaneous network excitability and behavioral expression. Recognition of the distributive focus underlying most seizures has motivated many surgical programs to upgrade their intracranial studies to capture activity in as much of the network as possible.

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### Keywords

Epilepsy • Seizure focus • Epileptogenic zone • Neural network • Simple (elementary) partial seizure • Complex partial seizure • Primary generalized seizure • Epilepsy surgery

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#### 4.1 Seizure Focus: Relation to Focal Neocortical Abnormalities

The seizure focus is usually defined as the site in the brain from which the seizure originated or, in the case of focal seizure discharge, the totality of the tissue involved. Localizing the site of seizure onset is critical to the understanding of seizure mechanisms and to the probability of “curing” the epilepsy through surgical resection. The goal of resective surgery is to remove or disconnect the epileptogenic zone, defined as the area of cortex indispensable for the generation of clinical seizures [33]. Should the epileptogenic zone then be considered equivalent to the seizure focus? Most likely yes, but unfortunately the boundaries of this region cannot be defined at present by any diagnostic test. Instead, imaging and EEG recording methods define cortical regions related to, but not necessarily contiguous with, the epileptogenic zone. These regions include the symptomatogenic region (the region that when activated by epileptiform discharge produces the ictal symptoms), the irritative region (the region(s) that generate(s) interictal spikes), the ictal onset region (the region from which the seizure is triggered), the region of functional deficit (the region that is functionally abnormal during the interictal period), and the epileptogenic lesion (structural damage that may be causally related to the seizures). When these regions are spatially concordant, they define the location of the epileptogenic zone (or focus). Frequently, however, there are discrepancies. This finding suggests the essential involvement of synaptically interconnected regions in the origination of most seizures. In these instances, the epileptogenic zone may be composed of the pacemaker or ictal onset region and one or more relay areas required to produce the ictal symptoms [13, 37, 44]. These are qualities more often attributed to generalized seizures. If not only generalized seizures but even the generation of many focal seizures requires activity within a spatially distributed network, then our concept of a seizure focus requires some reassessment.

The focus is related to, but is not synonymous with, the ictal onset region. Normal brain function

requires the correct balance between excitatory and inhibitory processes at every moment in time, and any disruption in this balance that favors excitation over inhibition promotes synchronous discharge of principal neurons. Thus seizures can be evoked in normal brain under conditions that promote such disruption. In focal epilepsy, the excitatory/inhibitory balance is disrupted chronically in some region or regions of the cerebral cortex, such that synchronous discharges arise under appropriate conditions. If these discharges are sufficient to provoke clinical symptoms, the region of chronic imbalance may be regarded as the ictal onset region. The ictal onset region is normally silent, but infrequently generates synchronous action potentials that can evoke afterdischarges. Its location can be approximated by EEG recording or SPECT imaging, but cannot be precisely defined. A great deal of research on animal models of epilepsy in the last 40 years has identified numerous abnormalities that under certain conditions can support the generation of episodic afterdischarges. Pathologic mechanisms found to promote hyperexcitability and seizure generation in animal models include loss or dysfunction of inhibitory neurons [1, 5, 21], creation or expansion of recurrent excitatory circuits [26], dysfunctional Na<sup>+</sup> and/or K<sup>+</sup> channels [12, 14, 31], enhanced intrinsic bursting [34, 35, 43], abnormal expression of HCN channels [29, 32], and altered glial regulation of extracellular fluid composition [9, 15, 39]. Typically, the process of epileptogenesis causes multiple functional and usually also anatomical changes in some region of brain that then becomes a locus for ictal onset. The relative importance of the various changes reported is unclear and is currently an area of active investigation. It is also possible that a single abnormality might trigger episodic seizure discharge when the ictal onset region is stimulated strongly enough or stimulated at an appropriate frequency. Synchronous firing of principal neurons in the ictal onset region provokes afterdischarge in the epileptogenic zone. The epileptogenic zone may be larger or smaller than the ictal onset region. It may include more than one potential ictal onset region differentiated by threshold. The most readily activated region will normally trigger all the seizures, but regions of higher threshold may

become evident if the low threshold region is resected or inhibited selectively. Conversely, resection of the entire epileptogenic zone would eliminate any clinical seizures even if a residual ictal onset region remains intact.

The epileptogenic zone or focus may also be distinguished from any structural lesion detected by MRI or histopathology, as well as from the symptomatogenic zone. Not all cortical lesions detected in a patient or animal having epileptic seizures are themselves epileptogenic. Additional testing is necessary to determine which, if any, is essential for the generation of seizures, and this is not often done. Furthermore, the epileptogenic zone may be larger or smaller than the anatomic lesion. When the epileptogenic zone includes only part of an anatomic lesion, the remaining lesion may not be capable of generating a seizure or may be capable only when driven by the portion of the lesion having a lower afterdischarge threshold. The epileptogenic zone may also extend beyond the anatomic lesions. For example, seizures may arise not because of the lesion itself, but rather from changes in surrounding cortical tissue induced by the lesion. In particular, tumors and vascular malformations induce foci of this type. Alternatively, MRI and even histopathology may not be sensitive enough to detect epileptogenic microlesions that extend some distance from the visible lesion.

Clinical symptoms will arise from the epileptogenic zone only if that zone includes a region of “eloquent cortex,” that is, cortex related to a specific function. Most of the human cortex is symptomatically silent, implying that seizures arise from activation of an epileptogenic zone primarily when epileptiform discharge propagates to a region of eloquent cortex with sufficient strength to elicit clinical symptoms. Thus cortical lesions and ictal symptomatology may, but usually do not, define the seizure focus.

To this point, the discussion of seizure focus has been limited to simple (or elementary) partial seizures, defined as seizures that originate from a limited region of the cerebral cortex and do not impair consciousness. The seizure may be provoked by hyperactivity that occurs spontaneously within a highly localized region or may require activation of a distributed

network. The concept of a distributive onset for many simple partial seizures links circuitry and mechanisms of these seizures with other seizure types suggested to arise from network activity.

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## 4.2 Focus of Complex Partial Seizures

Complex partial seizures, like simple or elementary partial seizures, originate from a limited region of the cerebral cortex (which includes the limbic system), but cause an impairment of consciousness. These seizures typically involve the temporal lobe, but some persons with epilepsy experience extratemporal complex partial seizures. Although attempts to associate epilepsy with a pathological substrate date from antiquity, Bouchet and Cazauviel [10] were the first to describe a “palpable induration of the temporal lobes” in a group of persons with epilepsy. Hughlings Jackson [18] later stimulated decades of searching for a single pathologic source when he replaced the term “psychomotor epilepsy” with temporal lobe epilepsy in the case of Dr. Z who died after having experienced seizures for many years. At autopsy, Hughlings Jackson suspected a lesion in the “taste regions of Ferrier,” and softening of the left uncinate gyrus was indeed discovered. Thus a mechanistic approach to ascribing epileptogenic causality to “the organ of the mind” was born.

Even earlier, Sommer [36] had concluded that hippocampal pathology (neuronal loss and gliosis; hippocampal sclerosis) is an important etiological factor in the subsequent development of seizures. The relationship between hippocampal damage and complex partial seizures has been debated ever since. To what extent does hippocampal pathology lead to epileptic attacks and to what extent do repeated seizures occurring over a period of years result in this pathology [25]? Finally, with the development of EEG, paroxysmal changes were found in the temporal cortex of patients with psychomotor seizures and Jasper et al. [20] used the term “temporal lobe epilepsy” to define these regional electrographic abnormalities.

Interictal spikes in the temporal region prompted some groups to resect portions of the temporal lobe in patients with medically-intractable complex partial seizures. Epileptogenic causality of hippocampal sclerosis remained controversial, however, until Margerison and Corsellis [22] combined EEG, clinical, and autopsy findings in patients institutionalized for epilepsy, and found hippocampal sclerosis in 30 of 34 patients in whom pre-mortem EEG had revealed anterior temporal spikes. The uniformity of neuronal cell loss and gliosis, particularly in hippocampal area CA1, in about 70 % of patients who were subjected to temporal lobectomy because depth electrode recordings had indicated medial temporal ictal onset and the 75 % control of seizures in those patients appeared to corroborate the growing sense that hippocampal sclerosis was the cause of temporal lobe epilepsy. In the mid-1980s to the early 1990s, MRI revolutionized the diagnosis of epilepsy, and unilateral hippocampal atrophy associated with interictal or ictal onset from one temporal lobe replaced most intracranial studies in diagnosing medial temporal lobe epilepsy. As MRI began to reveal an assortment of pathologies associated with suspected focal epileptogenesis, most, if not all, cases of temporal lobe epilepsy were ascribed to well-defined substrate pathologies of mesial temporal sclerosis, neoplasms, vascular lesions, developmental abnormalities, or gliosis from trauma, inflammation, etc. These findings coupled with the intrinsic excitability of the sclerotic hippocampus seen in depth electrode studies and human slice electrophysiology led many in the field to anticipate increasingly better surgical outcomes over time. However, surgical outcomes have not changed dramatically. In fact, several clinical and research observations have emphasized that epileptogenesis and ictal behavior is very likely a network phenomenon of aberrant nodes (review: [42]).

Resection of the anterior temporal lobe on the side of seizure onset, particularly removal of the hippocampus and amygdala, usually leads to cessation or at least reduction in frequency of temporal complex partial seizures. It is therefore often assumed that the epileptogenic zone must

be confined to these regions. Indeed hippocampal area CA3, the entorhinal cortex, and especially the basolateral amygdala exhibit a low threshold for seizure initiation when challenged with electrical stimulation or certain chemoconvulsants [6, 23, 45]. However, the anterior temporal lobe is a rather large block of tissue that includes distinct, but interconnected, brain regions. The epileptogenic zone has not been localized precisely and indeed a single point of onset may not exist. Seizure onset may arise from multiple sites within the temporal lobe [37, 38]. Wherever the onset, the remaining limbic regions are recruited rapidly, resulting in the same behavior regardless. Surgical outcomes also argue for a network as epileptogenic zone, rather than a localized focus. Bitemporal lobe epilepsy, proved by intracranial study, can be cured 50–60 % of the time by removal of only the more dysfunctional temporal lobe, indicating that the spontaneous ictal events in the contralateral lobe depended on network activation. In unilateral temporal lobe epilepsy, seizures are well-controlled after anterior temporal lobectomy in 75 % of patients, but control drops to 50 % when patients are followed for 10 years and falls even lower if antiseizure drugs are not administered. These results suggest that the entire epileptogenic zone is not being resected at least half the time. When one hippocampus is clearly responsible for ictal onset, a restricted mesial temporal resection (hippocampus + amygdala) yields poorer control (50–60 %) than a standard anterior temporal lobectomy. Also, in the most clearly lateralized cases, resection may stop the typical complex partial seizure, but the aura (sensation related to the ictal onset region) persists 15–20 % of the time. Finally, with the exception of some tumors and vascular lesions, patients almost always demonstrate distributed cognitive deficits on neuropsychological testing, again indicative of diseased network not a single diseased region.

Seizure onset may arise from multiple sites within the temporal lobe in animal models of epilepsy as well [7, 11]. In the kindling model, electrical stimulation of many sites within the limbic system evokes the same seizure type,

suggesting that activation of the network *per se* is more important than the precise point at which the network is activated [16]. In the kainic acid model, seizures can originate from either the amygdala or hippocampus at different times in the same rat [30]. These and other observations led to the proposal of a distributive focus for temporal complex partial seizures, which includes the hippocampus, amygdala, entorhinal cortex, anterior and midline thalamic nuclei, and pyriform cortex [8]. This hypothesis is further supported by findings that, in animal models of epilepsy, neurons in all these regions exhibit functional changes expected to promote excitability, creating potential ictal onset regions. In addition, these same brain regions are sites of tissue damage. Although hippocampal sclerosis is the characteristic form of histopathologic damage demonstrable in most patients who have undergone anterior temporal lobectomy, neuronal loss, atrophy, and gliosis have also been reported in the amygdala, entorhinal cortex, and thalamus [3, 28, 46]. Similarly widespread damage to the limbic system is found in commonly used animal models [27, 40]. Pyriform cortex, which is not regularly examined in human tissue specimens, is typically also damaged in animals. Pro-epileptogenic changes documented in animal models include degeneration of inhibitory neurons with subsequent axon sprouting by surviving inhibitory neurons, formation of recurrent excitatory connections by principal neurons, enhanced synaptically-evoked and intrinsic bursting, and altered expression and function of diverse ion channels and neurotransmitter receptors. In animals induced to become epileptic by provoking status epilepticus these changes clearly arise as a consequence of acute seizures. One or more of these changes may also precede the development of epilepsy in genetic models and in models of post-traumatic epilepsy. Regardless of how they were brought about, however, they all probably contribute to the spontaneous seizures, changes in circuit function perhaps being required for the origination of the seizures and histopathologic changes perhaps enhancing their frequency and intensity [17, 47]. The existence of a distributed pathological sub-

strate implies that seizures can arise at multiple points in the limbic circuit and that a certain minimum percentage of the circuit must be activated for the electrical activity to alter perception or behavior.

If temporal complex partial seizures can originate from any of several limbic regions, some of which lie outside the tissue normally resected for medically-intractable seizures, the distributive focus may explain, in part, the somewhat consistent percentage of temporal lobe surgeries (~30 %) that fail to achieve adequate seizure control [41]. Conversely, the success of many such surgeries may be attributable not so much to eliminating the seizure focus, but rather to removing enough of the limbic circuitry that synchronous firing in the remaining seizure onset regions fails to activate an epileptogenic zone.

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### 4.3 Focus of Primary Generalized Seizures

Primary generalized seizures appear to begin simultaneously in both cerebral hemispheres when recorded by scalp EEG, but are probably driven, at least in part, by hyperactivity of subcortical structures. Simultaneous activation of both cerebral hemispheres causes behavioral and perceptual signs and symptoms to be manifested bilaterally and there is always some impairment of consciousness. Although primary generalized seizures are not generally thought of as arising from a distinct focus or epileptogenic zone, the concept of a distributive focus appears applicable. This is perhaps best illustrated by the mechanisms underlying absence seizures. The spike-wave discharges of absence seizures require circuit interactions between the thalamus and neocortex [2, 19, 24]. Interruption of this circuit, such as by cutting the reciprocal pathways that connect the two regions or by inactivating either region alone, abolishes the seizures. Neocortex supplies the excitatory drive that is organized into ictal discharge by bursting thalamic relay neurons. Thus thalamus and neocortex together can be said to constitute an epileptogenic zone with

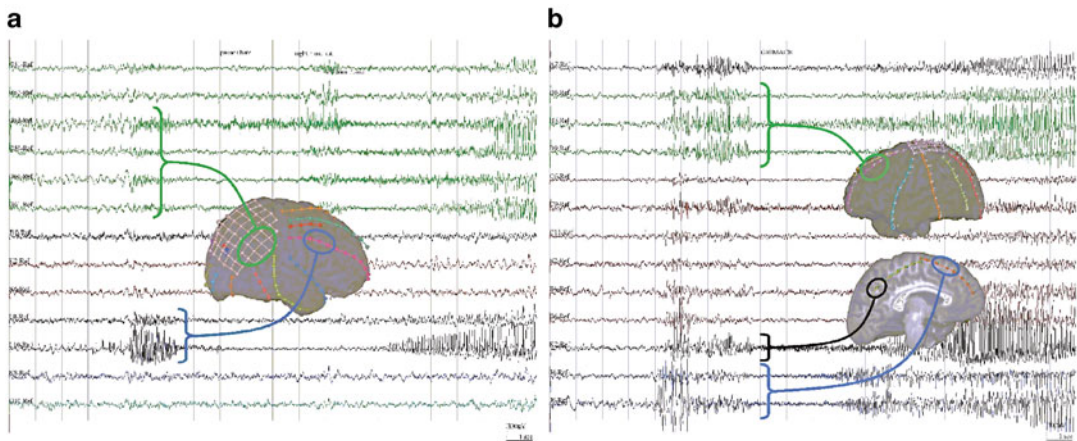
excitatory thalamic nuclei serving as the ictal onset region. Increasing evidence suggests the involvement of subcortical ictal onset regions in other forms of generalized epilepsy as well [4].

#### 4.4 Conclusion

Evolution in the thinking about concordance or non-concordance of the seizure-related cortical regions, has revealed discrepancies between the pathology and physiology of seizures. Definition of the epileptogenic zone remains complex and elusive, and the outcomes of surgical approaches have plateaued. Increasing evidence suggests that epileptogenesis is distributed among multiple foci, usurping known anatomical and functional networks. Very few of the most common seizures may be truly local. Rather, they appear to depend on nodal interactions that permit spontaneous network excitability and behavioral expression. The epilepsy community has not yet succeeded in creating a new paradigm that combines the critical derangements of electrophysi-

ology, pathology, metabolism, genetics, and network communication. New approaches must include better correlation of human data with animal models, wherein the hyperexcitable networks can be more intensively studied and manipulated.

Given the difficulty of defining the epileptogenic zone, many surgical programs have upgraded their intracranial studies to improve analysis of the distributed network. The Yale program, for example, has utilized advanced imaging and navigation systems to increase the number of electrode contacts per patient from <100 in 1991 to 200–250 in 2006 and to even greater numbers since then. Utilization of these electrode arrays has demonstrated many examples of distributive foci, such as those shown in Fig. 4.1. In these instances, the foci are located in functional networks revealed by fMRI – the cognitive control network between lateral parietal and frontal lobes (Fig. 4.1a) and a portion of what has been labeled the “default network,” observed reproducibly when subjects are at rest and not engaged in a task (Fig. 4.1b). Determinations of regional



**Fig. 4.1** Microelectrode array recordings from two surgical patients. **(a)** Reconstructed MRI/CT scan performed after intracranial electrode implantation. Electrode locations are indicated, along with the portion of the EEG recording associated with each electrode. The 28 year old right-handed female, whose MRI scan was normal, was found to have simultaneous ictal onset in the inferior parietal and inferior lateral frontal cortices without involvement of the intervening brain. These regions are the precise cortical areas activated in fMRI cognitive tasks and designated the cognitive network. Seizure control

was effected by resection of both ictal onset regions. **(b)** Similar superposition of MRI/CT scan, electrode locations, and EEG recording in a second patient. The 30 year old right-handed male, whose MRI scan was also normal, was found to have independent ictal onset in the posterior, medial frontal, and medial parietal lobes, with the medial frontal cortex initiating the same behavioral seizure more frequently. Initial treatment with a neurostimulator little affected the behavioral seizures. Subsequent resection of the medial frontal node alone was sufficient to control the seizures

extracellular glutamate concentration and metabolic/energetic studies that utilize 7 T MRS have further supported the concept of a distributive focus. At present, unless patients have a clear tumor or cavernoma on MRI or concordance of all data (electrophysiology, mesial temporal sclerosis on MRI, and neuropsychological studies) indicating unilateral temporal lobe epilepsy, intracranial studies are always performed and directed by anatomic MRI, dynamic imaging (FMR, ictal SPECT, MRS), AVEEG, and seizure semiology for distributed electrode placement. It is only by this persistent adaptation of newly developed technology that we can hope to one day understand and properly treat human epilepsy.

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# What Is a Seizure Network? Long-Range Network Consequences of Focal Seizures

# 5

Hal Blumenfeld

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## Abstract

What defines the spatial and temporal boundaries of seizure activity in brain networks? To fully answer this question a precise and quantitative definition of seizures is needed, which unfortunately remains elusive. Nevertheless, it is possible to ask under conditions where clearly divergent patterns of activity occur in large-scale brain networks whether certain activity patterns are part of the seizure while others are not. Here we examine brain network activity during focal limbic seizures, including diverse regions such as the hippocampus, subcortical arousal systems and fronto-parietal association cortex. Based on work from patients and from animal models we describe a characteristic pattern of intense increases in neuronal firing, cerebral blood flow, cerebral blood volume, blood oxygen level dependent functional magnetic resonance imaging (BOLD fMRI) signals and cerebral metabolic rate of oxygen consumption in the hippocampus during focal limbic seizures. Similar increases are seen in certain closely linked subcortical structures such as the lateral septal nuclei and anterior hypothalamus, which contain inhibitory neurons. In marked contrast, decreases in all of these parameters are seen in the subcortical arousal systems of the upper brainstem and intralaminar thalamus, as well as in the fronto-parietal association cortex. We propose that the seizure proper can be defined as regions showing intense increases, while those areas showing opposite changes are inhibited by the seizure network and constitute long-range network consequences beyond the seizure itself. Importantly, the fronto-parietal cortex shows sleep-like slow wave activity and depressed metabolism under these conditions, associated with

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impaired consciousness. Understanding which brain networks are directly involved in seizures versus which sustain secondary consequences can provide new insights into the mechanisms of brain dysfunction in epilepsy, hopefully leading to innovative treatment approaches.

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### Keywords

Epilepsy • Consciousness • Slow waves • Cortex • Thalamus • Sleep • Hippocampus • Pedunculo-pontine tegmental nucleus • Acetylcholine • Brainstem • Arousal

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## 5.1 Introduction

Seizures are usually defined as an abnormal pattern of neuronal activity which includes excessive synchrony and high frequency firing of neurons. As in most definitions, the obvious cases are easy to recognize. However, in reality there are no distinct boundaries for precisely when neuronal activity become sufficiently synchronous or intense to be considered a seizure. The situation is complicated further by the fact that seizures occur in neuronal networks, which have both local and long-range effects. Network interactions give rise to abnormal activity in local circuits, but in some cases can also influence remote brain regions. Are these remote network changes part of the seizure proper, or are they “side effects” caused by the seizure but not directly involved in the seizure network? To answer this question it is necessary to identify characteristic features that are seen in seizure activity, and to then determine if these same features are present in the remote network regions. If similar characteristic features are present, then the remote regions are likely to be involved in propagation of the seizure itself. If the activity in the remote regions differs drastically from seizure activity, and instead resembles other well-known patterns of non-seizure brain activity (such as coma or sleep), then the activity in the remote region could be considered outside the seizure network, although influenced by it.

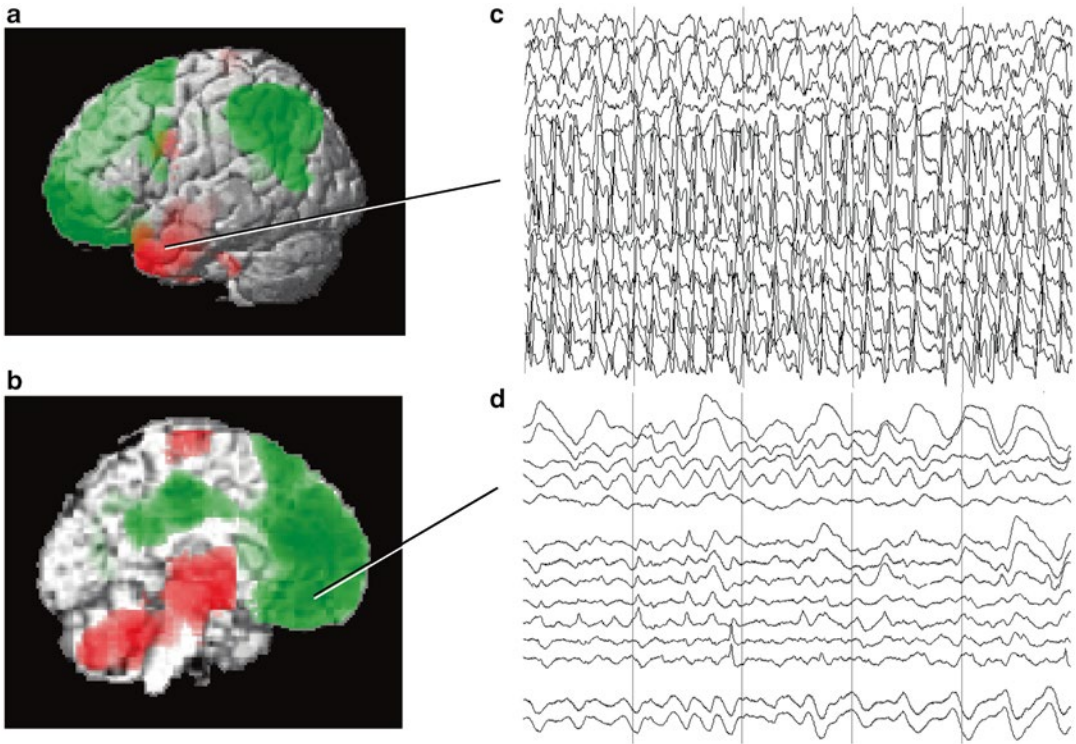
Temporal lobe seizures provide a concrete example of these local and long-range network phenomena. Locally, temporal lobe seizures

produce high frequency rhythmic discharges. At the same time remote regions of the fronto-parietal association cortex exhibit 1–3 Hz slow wave activity resembling coma, sleep or encephalopathy [1–3]. Is this slow wave activity part of the seizure, or is it a distinct state of brain activity caused by the seizure? Here we will examine the detailed characteristics of these remote changes in neocortical networks during focal limbic seizures in both patients and in animal models, and also potential mechanisms for these phenomena. We conclude that these remote effects on neocortical networks are best considered outside the seizure network but strongly influenced by it. Analogous to post-ictal depression, which is closely related to and caused by the seizure itself but occurs at a different time, neocortical slow wave activity is closely related to and caused by focal limbic seizures but occurs in a different space.

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## 5.2 Clinical Data

Intracranial recordings from patients with temporal lobe epilepsy show characteristic low voltage fast activity evolving into rhythmic polyspike-and-wave discharges in the medial temporal lobe limbic circuits, often extending into the lateral temporal cortex (Fig. 5.1c). Simultaneously, remote regions of the frontal and parietal association cortex often show 1–3 Hz slow wave activity (Fig. 5.1d). This ictal neocortical slow wave activity has been interpreted as a propagation pattern in temporal lobe epilepsy [1]. However, several features of the fronto-parietal slow wave activity make it likely that this is a distinct,



**Fig. 5.1** Local and long-range network effects in temporal lobe complex partial seizures. (a, b) Group analysis of SPECT ictal-interictal difference imaging during temporal lobe seizures. CBF increases (*red*) are present in the temporal lobe (a) and in the medial thalamus (b). Decreases (*green*) are seen in the lateral frontoparietal association cortex (a) and in the interhemispheric frontoparietal regions (b). (c, d) Intracranial EEG recordings from a patient during a temporal lobe seizure. High frequency polyspike-and-wave seizure activity is

seen in the temporal lobe (c). The orbital and medial frontal cortex (and other regions, EEG not shown) do not show polyspike activity, but instead large-amplitude, irregular slow rhythms resembling coma or sleep (d). Vertical lines in (c) and (d) denote 1-s intervals. Note that the EEG and SPECT data were from similar patients, but were not simultaneous, and are shown together here for illustrative purposes only ((a, b) Modified from Blumenfeld et al. [2] with permission. (c, d) Modified from Englot et al. [3] with permission)

remote network effect rather than simply seizure propagation, as we discuss below.

Recent work with multiunit recordings in human intracranial EEG has raised new questions about the definition of seizure activity vs. associated changes in surrounding regions. Schevon and colleagues showed that high frequency firing of neurons is highly localized in human seizures [4]. Accompanying local field potential changes measured by conventional intracranial EEG extend over a greater region, but may represent mainly synaptic activity without major changes in local firing of neurons [4]. Whether recording neuronal firing or local field potentials, at least these changes in the vicinity of seizure onset show

high frequency poly-spike activity characteristic of seizure physiology. In contrast, the slow wave activity occurring in distant fronto-parietal regions during temporal lobe seizures occur at a very different frequency (1–3 Hz) from ictal temporal lobe polyspike discharges (broad band >8 Hz) (Fig. 5.1c, d) [2, 3]. Seizure activity on intracranial EEG can be defined as high frequency discharges. Although scalp EEG often exhibits rhythmic theta or delta-frequency slow waves during local seizures [5] direct recording of seizure activity with intracranial electrodes inevitably shows high frequency discharges in these same regions. Therefore, when only slow wave activity is seen in a region *without* high frequency discharges on

intracranial EEG, this likely does not represent seizures. As we discuss in the next section, detailed physiological studies from animal models further support this claim. Such slow wave activity seen in the fronto-parietal cortex during temporal lobe seizures is similar to cortical slow waves in deep sleep, coma or encephalopathy [6, 7]. In these states, cortical function and information processing is depressed, leading to impaired level of consciousness [8].

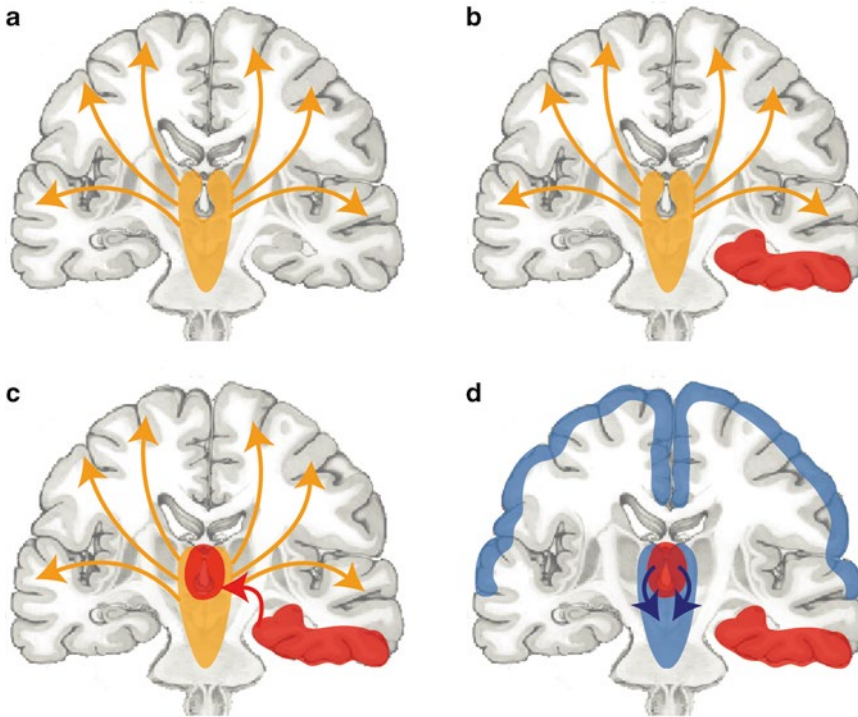
How does focal seizure activity in the temporal lobe lead to remote slow wave activity in the fronto-parietal association cortex? The anatomy and physiology of these changes differs from local “surround inhibition” described for focal cortical seizures [9, 10]. To affect distant lobes, long-range network interactions are required. Some initial clues for the mechanisms of these network changes have come from human cerebral blood flow (CBF) imaging with single photon computed tomography (SPECT) which, unlike fMRI, can be done successfully despite patient movement during seizures. As expected, ictal SPECT in temporal lobe seizures is associated with CBF increases in the temporal lobe (Fig. 5.1a). In addition, *decreases* are seen in frontal and parietal association cortex in the same regions which exhibit slow wave activity (Fig. 5.1a, b) [11–13]. Subcortical networks are also involved in temporal lobe seizures and SPECT imaging shows increases in the medial thalamus and midbrain (Fig. 5.1b) [13–16]. We found that the SPECT increases in the medial thalamus are correlated with the decrease in bilateral fronto-parietal cortex [13], suggesting a mechanistic link between subcortical changes and depressed cortical function in temporal lobe seizures. These long-range network changes in cortical and subcortical function are seen specifically in temporal lobe seizures with impaired consciousness [3, 13, 14, 17, 18]. In contrast, temporal lobe seizures without impaired consciousness are associated with localized seizure activity in the temporal lobe, without these long-range network changes [3, 13].

Based on these findings from patients, we proposed the *network inhibition hypothesis* to explain cortical dysfunction and impaired con-

sciousness in temporal lobe seizures (Fig. 5.2) [19, 20]. Normal cortical function and consciousness is maintained by interactions between the cortex and subcortical arousal systems including the thalamus, brainstem and basal forebrain (Fig. 5.1a). Focal temporal lobe seizure activity in simple partial seizures does not have long-range network impact effects, so cortical function and consciousness are spared (Fig. 5.1b). In temporal lobe complex partial seizures, propagation to subcortical structures (Fig. 5.1c)—such as the anterior hypothalamus, lateral septum and other regions—inhibits subcortical arousal systems (Fig. 5.1d). This in turn removes cortical arousal leading to fronto-parietal slow wave activity and impaired level of consciousness. Note that according to this hypothesis, the cortical slow wave activity is not part of the seizure itself, but instead is a long-range network consequence of depressed subcortical arousal.

Further support for the network inhibition hypothesis has come from recent behavioral observations in patients [21–23]. The network inhibition hypothesis predicts that when focal seizures propagate to subcortical structures, this will cause severe and widespread cortical dysfunction. Therefore focal seizures are expected to usually be associated with either marked impairment of many cognitive functions due to depressed level of consciousness, or alternatively to spare most cognitive functions. In support of this hypothesis, we recently found that behavioral deficits in a wide range of verbal and non-verbal test items during partial seizures are bimodally distributed, such that most seizures either globally impair or spare cognition [21–24].

While human studies have provided clinically relevant correlations between physiology and behavioral changes, and suggest that ictal neocortical slow wave activity is distinct from direct seizure involvement, fundamental mechanistic studies are best performed in animal models. An experimental animal model could enable direct physiological measurements to determine if ictal neocortical slow wave activity is indeed distinct from seizure activity, and would allow further investigation of the mechanisms for this phenomenon.



**Fig. 5.2 Network inhibition hypothesis.** (a) Under normal conditions, the upper-brainstem and diencephalic activating systems interact with the cerebral cortex to maintain normal consciousness. (b) A focal seizure involving the mesial temporal lobe. If the seizure remains localized, a simple partial seizure will occur without impairment of consciousness. (c) Seizure activity often

spreads from the temporal lobe to midline subcortical structures and propagation often extends to the contralateral mesial temporal lobe (not shown). (d) Inhibition of subcortical arousal systems leads to depressed activity in bilateral frontoparietal association cortex and to loss of consciousness (Modified from Englot et al. [3] with permission)

### 5.3 Insights from an Experimental Animal Model

Rodent models of limbic seizures replicate many of the behavioral and physiological characteristics of human temporal lobe epilepsy [25–29]. We found that spontaneous focal limbic seizures in awake chronically epileptic rats following pilocarpine status epilepticus exhibited frontal neocortical 1–2 Hz slow wave activity and behavioral arrest similar to human complex partial temporal lobe seizures [30]. Ictal neocortical slow wave activity in this model resembled slow wave activity during natural slow wave sleep in the same animals. In contrast when limbic seizures secondarily generalized, recordings from the frontal cortex showed 9–12 Hz polyspike

discharges characteristic of ictal activity, instead of slow waves.

Additional physiological and neuroimaging experiments were performed in an acute lightly anesthetized rat model in which seizures could be induced under controlled conditions [30]. Seizures were induced by brief 2 s stimulus trains at 60 Hz to the hippocampus under ketamine/xylazine anesthesia reduced to a stage where the cortex showed physiology near to the waking state. Under these conditions, induced partial limbic seizures produced frontal cortical slow wave activity similar to that seen in awake chronically epileptic rats. This acute model enabled detailed physiological measurements to distinguish ictal neocortical slow waves from seizure activity. Measurements from the hippocampus

during partial limbic seizures revealed dramatic increases in neuronal firing (multiunit activity), cerebral blood flow, blood oxygen dependent (BOLD) functional magnetic resonance imaging (fMRI) signals, cerebral blood volume, and cerebral metabolic rate of oxygen consumption [30]. In marked contrast, during the same seizures the frontal cortex showed *decreases* in all of these measurements along with slow wave activity. These findings provide strong evidence that ictal neocortical slow wave activity is a distinct physiological state, more closely resembling deep sleep or encephalopathy than seizure activity. Indeed, in the same animals slow wave activity under deep anesthesia induced similar changes in neuronal activity in the frontal cortex to those observed during partial limbic seizures.

Further evidence supporting a physiological distinction between ictal neocortical slow waves and seizure activity was provided by secondarily generalized seizures [30]. As in the awake model, when seizures propagated to the frontal cortex, instead of slow waves the frontal cortex showed high frequency polyspike discharges. Unlike the physiological decreases seen during slow wave activity, during secondary generalized seizures the frontal cortex showed marked *increases* in neuronal firing, cerebral blood flow, BOLD fMRI signals, cerebral blood volume, and cerebral metabolic rate of oxygen consumption.

In summary, direct measurements and neuroimaging during focal limbic seizures revealed very distinct physiology for hippocampal or cortical seizure activity which generally showed marked increases in all neurometabolic functions, contrasting markedly with ictal neocortical slow activity which showed opposite changes, with decreases in all markers of neurometabolic function. These findings support the hypothesis that ictal neocortical slow wave activity is not part of the seizure itself, but instead is a consequence arising from long-range network effects producing altered physiology in regions remote from the seizure focus.

The next step has been to identify the network mechanisms by which seizure activity in the hippocampus may produce slow wave activity in the neocortex. As we have already discussed, data from patients suggest that focal hippocampal seizures

may depress subcortical arousal systems, which could lead to cortical slow wave activity resembling deep sleep or coma (Fig. 5.2). Experiments from the rat model have provided further mechanistic details to support this hypothesis [31]. fMRI mapping during focal limbic seizures demonstrated that seizure activity propagates from the hippocampus to subcortical structures including the lateral septal nuclei, anterior hypothalamus, and medial thalamus. Subsequent direct neuronal recordings confirmed increased activity in these subcortical regions during seizures. The lateral septal nuclei and anterior hypothalamus contain gamma amino butyric acid (GABA)-ergic neurons with projections to subcortical arousal structures and are thus well positioned to inhibit cortical arousal during seizures. In support of this model, electrical stimulation of these regions in the absence of seizure activity was able to reproduce cortical slow wave activity and behavioral arrest [31, 32]. Cutting the fornix, the main route of seizure propagation from hippocampus to these subcortical structures, prevented cortical slow wave activity and behavioral arrest during seizures.

Additional studies have confirmed decreased subcortical arousal during focal limbic seizures, specifically in the cholinergic arousal systems [32]. fMRI mapping during focal limbic seizures have shown decreased signals in the midbrain reticular formation, thalamic intralaminar nuclei and possibly the basal forebrain. Juxtacellular recordings from the pedunculopontine tegmental nucleus in the brainstem demonstrated decreased firing of identified cholinergic neurons during frontal cortical slow wave activity in focal limbic seizures [32]. In addition, amperometric measurements of choline signals as a surrogate marker of cholinergic neurotransmission showed decreases in both frontal cortex and intralaminar thalamus during focal limbic seizures, but not during secondarily generalized seizures. While it is likely that in addition to cholinergic arousal other subcortical arousal systems are also involved, these findings provide strong evidence that a well characterized subcortical arousal system is depressed during focal limbic seizures, resembling the decreased function seen in slow wave sleep.

## 5.4 Conclusions and Future Directions

Here we have examined the activity patterns in focal limbic seizures to ask the question: What is a seizure network? In this case, more specifically—which changes in activity during limbic seizures represent the seizure itself and which can be considered long-range network effects arising from, but physiologically distinct from seizure activity? Based on multi-modal measurements including direct recordings of neuronal activity, cerebral blood flow, and neuroimaging-based evaluation of neuroenergetics, we conclude that limbic seizure networks involve intense increases in activity in structures such as the hippocampus and subcortical regions including the lateral septum and anterior hypothalamus. As a long-range network consequence of this abnormal increased activity, there is also abnormal decreased activity in subcortical arousal systems including the brainstem, intralaminar thalamus and basal forebrain which causes the cortex to enter a state resembling deep sleep. These subcortical and cortical decreases in activity are not part of the seizure *per se* since they differ drastically from the increases typically associated with seizures. However, they are an important effect of the seizure network on other parts of the brain, and have a major clinical impact including impaired consciousness.

Important unanswered questions remain about these seizure networks. For example, although the presence of GABAergic neurons in structures involved in seizures (such as the lateral septum or anterior hypothalamus) suggests these may inhibit subcortical arousal systems, direct demonstration of subcortical inhibition has not yet been confirmed. Additional experiments including local infusion of GABAergic agonists and antagonists will be crucial. In addition, while cholinergic arousal was found to be depressed during limbic seizures, the possible involvement of other neurotransmitter systems should be investigated further. Another important direction for future investigation is the development of treatments to prevent long-range network impairment. Although ideally the sei-

zures themselves should be stopped, in some patients this is not possible. In these medically and surgically refractory cases, treatments aimed at preventing the impaired consciousness which accompanies depressed cortical function would be highly beneficial. Possible treatments based on the findings above would include deep brain stimulation targeted at arousal regions such as the thalamic intralaminar region [33, 34] or pharmacological treatments such as modafinil [35] aimed at increasing alertness in the ictal and post-ictal periods. Hopefully, further investigation of the interactions between local seizures and long-range network interactions will make such treatments possible, improving the lives of people with epilepsy.

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# What Is a Seizure Network? Very Fast Oscillations at the Interface Between Normal and Epileptic Brain

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## Abstract

Although there is a great multiplicity of normal brain electrical activities, one can observe defined, relatively abrupt, transitions between apparently normal rhythms and clearly abnormal, higher amplitude, “epileptic” signals; transitions occur over tens of ms to many seconds. Transitional activity typically consists of low-amplitude very fast oscillations (VFO). Examination of this VFO provides insight into system parameters that differentiate the “normal” from the “epileptic.” Remarkably, VFO *in vitro* is generated by principal neuron gap junctions, and occurs readily when chemical synapses are suppressed, tissue pH is elevated, and  $[Ca^{2+}]_o$  is low. Because VFO originates in principal cell axons that fire at high frequencies, excitatory synapses may experience short-term plasticity. If the latter takes the form of potentiation of recurrent synapses on principal cells, and depression of these on inhibitory interneurons, then the stage is set for synchronized bursting – if  $[Ca^{2+}]_o$  recovers sufficiently. Our hypothesis can be tested (in part) in patients, once it is possible to measure brain tissue parameters (pH,  $[Ca^{2+}]_o$ ) simultaneously with ECoG.

## Keywords

pH • Extracellular calcium • Gap junctions • Potentiation • Oscillation • Electrocorticogram

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## Abbreviations

ACSF	Artificial cerebrospinal fluid
DHPG	(S)-3,5-dihydroxyphenylglycine
ECoG	Electrocorticography
TMA	Trimethylamine
VFO	Very fast oscillations (>70 Hz)

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## 6.1 Introduction

The task of defining – or identifying – a seizure network is conceptually very complex and can be approached in a number of different ways. One could, for example, determine which brain regions (and which cell types) are the first to discharge in a “non-normal” fashion that leads to aberrant EEG patterns. This approach has been the conventional one, and has led to the concept of the epileptic “focus” or epileptic “zone.” These concepts have been problematic since it is now clear that – at least in the chronic human epileptic brain – cells in rather widespread brain regions are often linked in their aberrant discharge patterns as seizure are initiated (e.g., Worrell’s work). A related but more recent approach has been to identify those brain regions that generate high frequency oscillations at the onset of seizure activity. The use of such oscillations as a biomarker for “epileptic brain” has received much attention, and seems to provide a useful guideline for surgical intervention (i.e., removal results in “cure”). With this latter approach, it would appear that the “seizure network” is defined as that group of cells that generate these abnormally high frequency EEG patterns. And thus an understanding of these generators would provide a useful handle on defining a seizure network – and for asking such questions as whether such networks are dynamic, are reflective on tissue pathology, are exclusive to networks in epileptic brain (i.e., do not come into play in normal brain when seizures are exogenously generated), etc. We have therefore approached the question of “epileptic networks” via our interest in very fast oscillations (VFO).

The data discussed below suggest that the transition from normal brain rhythms to seizure

is brought about (at least in an immediate sense) by alterations in brain tissue, *in the extracellular environment* rather than by neuronal activities per se – an idea that has been central to the epilepsy scientific endeavor for many years. As shown by many other authors (and also ourselves), very fast oscillations (VFO) – a striking and (we believe) fundamental sort of neuronal activity – are frequently observed prior to and during seizures. What we bring to the table that is new is this: VFO occurs in just those ionic and pH conditions expected to occur after brain activation, and which might in themselves promote seizures. Furthermore, VFO itself could induce synaptic habituation (specifically of pyramidal/interneuron synapses) that would also favor seizures. This emphasis on the extracellular environment, and on the mechanisms for transition from “normal” electrical activity to “seizure” activity, provides perhaps a new view of what might profitably be explored as a defining feature of epileptic networks.

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## 6.2 Very Fast Oscillations in Normal and Epileptic Brain

During epileptic burst complexes (both interictal and ictal), there coexist large field transients (often with simultaneous intracellular depolarizations and multiple action potentials), together with high-frequency field oscillations (“VFO”), the latter sometimes at several hundred Hz [other terms include “ripples”, “fast oscillations”, and HFO or high-frequency oscillations]. This coexistence was observed in penicillin-induced epileptogenesis in cat hippocampus *in vivo*, in 1969 [7]; and not too long afterwards in the *in vitro* hippocampal slice by Philip Schwartzkroin and David Prince [32, 33]. Since then, coexisting large field transients, with superimposed VFO, have been observed in patient EEGs (for example, [42], and see also below), as well as in many experimental contexts ([24]; reviewed in [45]).

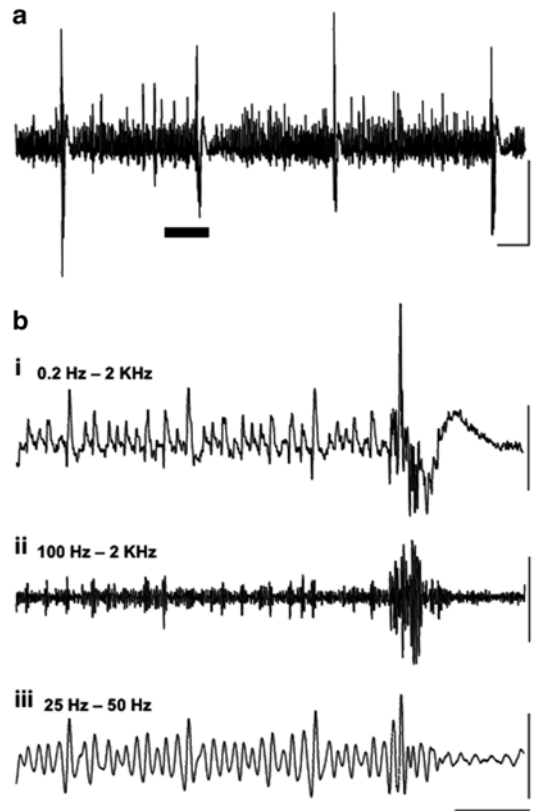
How can one account for the coexistence of these two field patterns, and their relation to normal brain activities, such as gamma (30–70 Hz) rhythms and physiological sharp waves? In this chapter, we shall note that putatively normal-appearing gamma

can alternate with large synchronized bursts, providing a model for the transition between normal and abnormal neuronal population behaviors. Interestingly, in this model – and in many other situations, including in patients – there is a segment of VFO *prior* to the synchronized burst. We shall examine the somewhat surprising conditions in which, experimentally, VFO can occur alone; and we shall review the cellular mechanisms of one experimental type of gamma oscillations (which turns out to be related to VFO). Finally, we shall conclude with an hypothesis as to how the transition from relatively normal activities, to epileptic ones, might take place *in situ*. Imbalance between synaptic excitation and inhibition – the text-book explanation – provides partial, but not complete, understanding. Our hypothesis is testable, at least in part; and, if valid, the hypothesis may have clinical application.

*An example of the alternation between “normal” gamma rhythm and epileptiform bursts.* It was discovered in 1998 (Fisahn et al.) that stable (i.e. lasting hours) gamma oscillations could be induced in properly prepared hippocampal slices, simply by addition of a compound such as carbachol to the bath. Similar oscillations can be produced by other compounds, including kainate, in hippocampus, neocortex, entorhinal cortex, and cerebellum slices (reviewed in [45]). [We shall describe some of the cellular mechanisms below.] Interestingly, a high concentration of the metabotropic glutamate receptor agonist DHPG induces oscillations that alternate with epileptiform bursts, over periods of several seconds (Fig. 6.1). The amplitude of EPSPs in interneurons and in pyramidal cells evolves over the interburst periods, decreasing in interneurons, and increasing in pyramidal cells; and this explains, at least in part, the switch in behaviors [46]. Note, however, that field VFO actually precedes the epileptiform bursts (Fig. 6.1bii, and see also [24]).

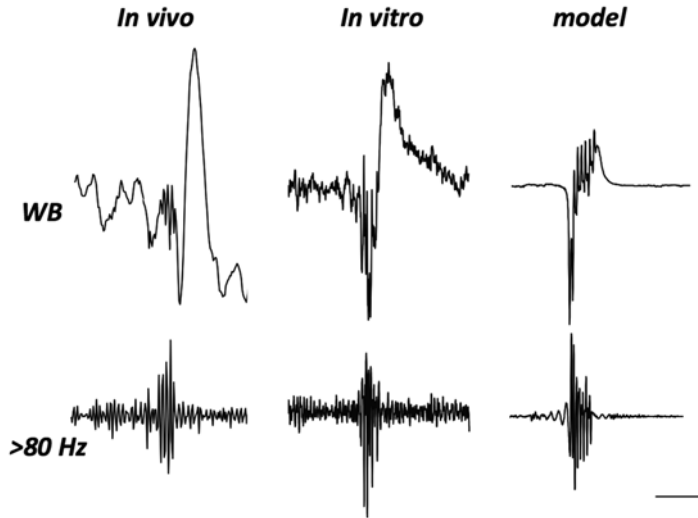
*Further examples of VFO associated with, and prior to, epileptic transients and “full-blown” electrographic seizures.*

The slight advance of VFO, relative to epileptiform bursts, may be a quite general phenomenon (Fig. 6.2), occurring also in human tissue *in situ*, as well as in resected human tissue. Such observa-



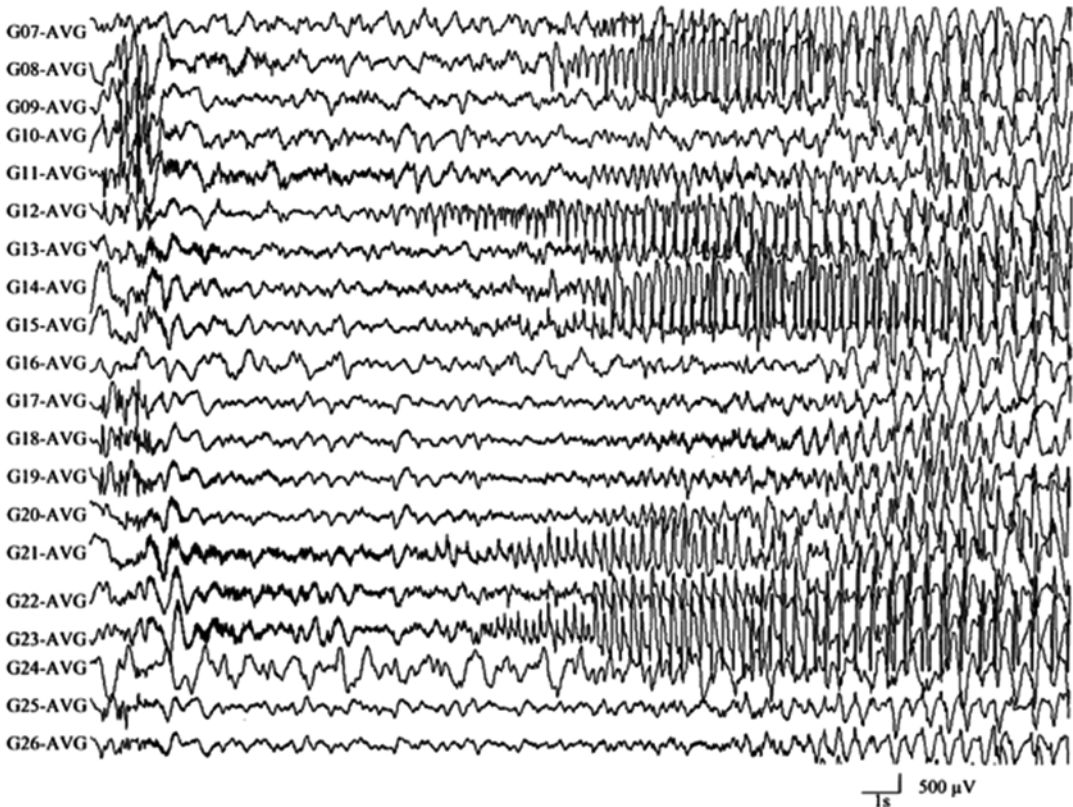
**Fig. 6.1 Alternating gamma oscillation and synchronized epileptiform bursts.** Rat hippocampal slice, CA3 region, bathed in 100  $\mu$ M DHPG (a metabotropic glutamate receptor agonist), s. pyramidal field recordings. (a) long-duration trace showing 4 epileptiform bursts with interspersed gamma oscillations (~30 Hz). Scale bars 0.5 mV, 1 s. (b) the segment corresponding to the bar in (a) is expanded, and filtered to show broad-band (i), VFO (ii), and gamma (iii) signals. Note the brief VFO just prior to the epileptiform burst. Scale bars 0.5, 0.1, 0.2 mV; 200 ms (From Traub et al. [44], reproduced with permission)

tions suggest that perhaps VFO is really the “fundamental” event in epileptic bursts. Indeed, in resected human tissue, it has been shown that blockade of chemical synapses can eliminate the large field transients, while leaving VFO; whereas block of VFO with carbenoxolone also causes loss of the large transients [27]. At least in the experimental conditions there used, it was not possible to observe large transients without VFO, while the reverse could be observed. One wonders, therefore, if VFO at least contributes to the causation of the epileptiform bursts (Fig. 6.2).



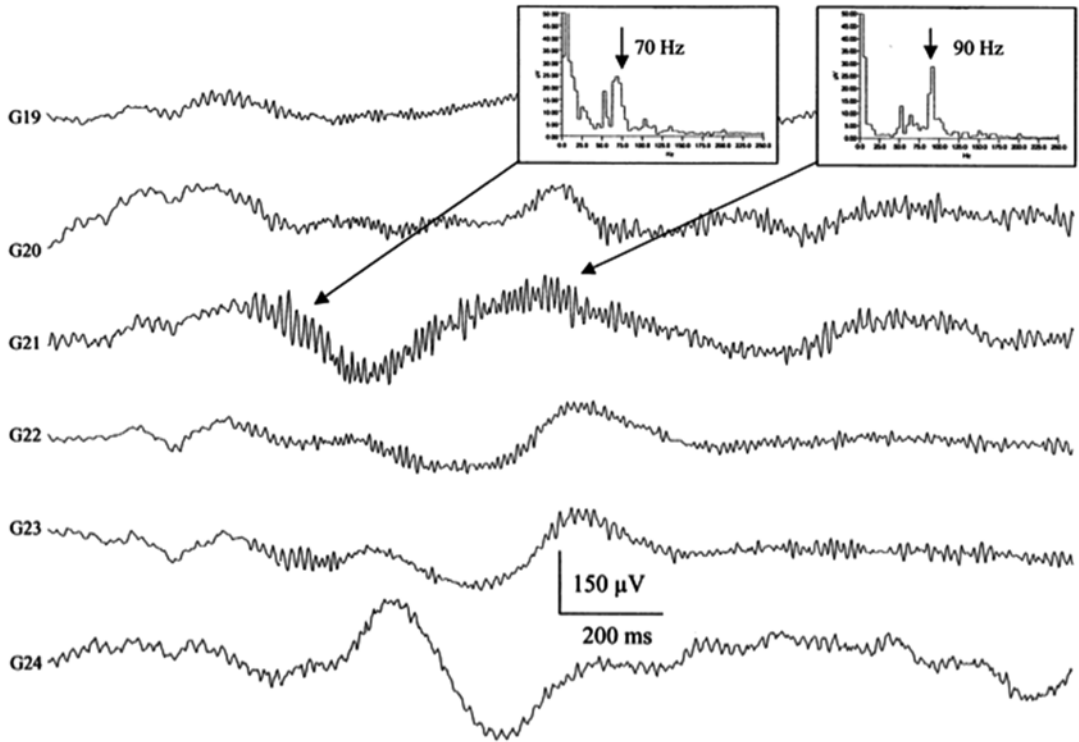
**Fig. 6.2** VFO preceding epileptiform bursts: 3 examples. (“WB”=wide-band.) *Left*, *In vivo*, foramen ovale recording of right temporal interictal activity in a patient with mesial temporal sclerosis. *Middle*, *In vitro*, spontaneous field potential burst in resected temporal neocortex from the same patient. *Right*, model, simulation of net-

work burst in multilayer neocortical circuit model, with multicompartiment neurons interconnected by chemical synapses and by gap junctions. Scale bars 200  $\mu\text{V}$  *in vivo*, 100  $\mu\text{V}$  *in vitro*, arbitrary for model; 100 ms (From Roopun et al. [27], reproduced with permission)



**Fig. 6.3** Subdural grid ECoG recording of an electrographic seizure, preceded by a ~2 s generalized discharge, and then localized, low-amplitude VFO (e.g. G21-G23).

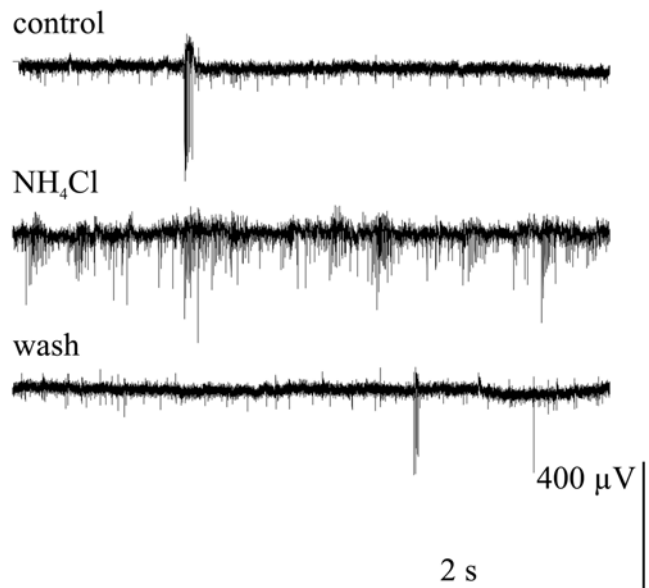
Recordings from a child with a *right frontal* cortical dysplasia and intractable seizures. She responded well to surgery (From Traub et al. [42], reproduced with permission)



**Fig. 6.4** Another run of pre-seizure VFO in ECoG. This example was recorded from the same patient whose ECoG was shown in Fig. 6.3, with the same subdural grid

but different recording technique (From Traub et al. [42], reproduced with permission)

**Fig. 6.5** *In vitro* ~200 Hz ripples are strongly potentiated by tissue alkalinization. Stratum pyramidale recordings of spontaneous VFO in the CA3 region of rat hippocampal slice. VFO occurs transiently in control conditions (*top*), but becomes nearly continuous after tissue alkalinization with 10 mM NH<sub>4</sub>Cl. The effect is reversible (From Draguhn et al. [8], reproduced with permission)



At times, VFO can be sustained for seconds prior to the onset of an electrographic seizure (but not, so far as we are aware, of an interictal

burst). Figure 6.4 shows an example of this phenomenon, in an electrocorticographic (ECoG) recording.

What the above data suggest is that VFO mechanisms may provide a clue as to what is distinctive about normal brain rhythms, as opposed to epileptiform events. In order to explore this idea further, we must make a digression into some of the relevant basic cellular mechanisms.

### 6.3 Cellular Mechanisms of Epileptiform Bursts, VFO, and Gamma Oscillations

Synchronized epileptiform bursts are considered, traditionally, to arise from an imbalance in synaptic excitation and inhibition – an idea perhaps rooted in the experimental observation that blockade of GABA<sub>A</sub> receptors was an effective experimental means of inducing such bursts [7]. The imbalance idea does not explain, however, why one does not simply observe sustained increases in firing rates; instead, epileptiform activity is organized into transient events, lasting tens to hundreds of ms. Furthermore, at least *in vitro*, transient events can be elicited by stimulation of a small number of neurons, sometimes even one neuron [19], although there can be a latency of >100 ms from the stimulus to the population event. This occurs, even though the density of excitatory synaptic connections, *in vitro* in CA3, is of the order of a few per cent. Traub and Wong [39] were able to account for the above observations, if it were postulated that recurrent excitatory connections were sufficiently strong – specifically, that a burst of action potentials could induce a burst in a synaptically connected cell, in the relative absence of synaptic inhibition. This prediction was then verified with paired recordings [20]. Notably, however, the model under consideration did not account for the VFO superimposed on epileptiform bursts. Why is this important? Couldn't it be that the VFO is simply an irrelevant epiphenomenon?

We shall argue that the VFO is important, for a number of reasons, but in the present context, consider the following argument. “Strong” coupling between neurons appears to be important for a synchronized burst to develop. Suppose that gap

junctions were to exist between principal neurons, with coupling powerful to allow a single action potential in one cell to evoke an action potential in another cell. This type of strong electrical coupling does actually exist [18, 48], and it could cooperate with recurrent excitatory chemical synapses. Additionally, as we shall note below, electrical coupling accounts for VFO itself.

VFO: high-frequency oscillations (“ripples”) had been observed in the hippocampus *in vivo*, during physiological sharp waves [4], but distinctive clues to cellular mechanisms came from the discovery that ripples could occur *in vitro*, without sharp waves [8] – the ripples could then be studied in isolation. Remarkably, ripples can occur *in vitro* without chemical synapses, both in hippocampus and in the neocortex [8, 22, 46]. Ripples are coherent (*in vitro*) over hundreds of microns, and so are a true population phenomenon. Extracellular fields (tens of  $\mu\text{V}$ ) are too small to explain them, and a variety of pharmacological manipulations are consistent with gap junctions being fundamental. *In vitro* ripples are also associated with spikelets [8, 46] which, in the hippocampus, are likely of axonal origin [28]. Dye-coupling exists between axons of nearby CA1 pyramidal cells [28], consistent with the occurrence of gap junctions between axons, although not providing definitive proof (by itself) for this concept.

We have shown that *in vitro* VFO, at frequencies up to about 250–300 Hz, can be explained by electrical coupling between axons under certain conditions: first, the coupling is strong enough for a spike in one axon to evoke a spike in a coupled axon (indirectly supported by data of Dhillon and Jones [6], Mercer et al. [18] and Wang et al. [48]); second, each axon couples, on average, to more than one other; finally, that spontaneous axonal action potentials occur at least sometimes. This model accounts for the admixture of spikes and spikelets during VFO, for continuous frequency transitions from gamma to almost 200 Hz, and for spatial patterns of VFO in the neocortex [5, 35, 40, 46]; and, most importantly, it accounts for the propensity of VFO to occur when chemical synapses are

blocked. The model predicts that somatic action potentials during VFO are antidromic [2, 47].

**Persistent gamma oscillations** are traditionally viewed as arising simply from recurrent synaptic excitation to interneurons, and synaptic inhibition to pyramidal neurons. A number of pieces of experimental evidence indicate that the mechanisms are somewhat more complicated. First, while it is true that blockade of AMPA/kainate, or of GABA<sub>A</sub> receptors, will suppress persistent gamma, it is also true that persistent gamma is sensitive to gap junction blockade [11, 41, 42]. Second, the power spectrum of gamma oscillation fields reveals a peak at 70 or more Hz. This activity can be seen in Fig. 6.1bii. This faster peak is not simply a harmonic of the gamma activity, because the high frequency peak persists when gamma is abolished by synaptic receptor blockade [42], or when stratum oriens is separated from stratum pyramidale – in which case VFO persists in s. oriens [43]. Finally, pyramidal cell somata fire rarely during persistent gamma [11].

The above disparate and counter-intuitive observations are readily explained with a model that basically simulates persistent gamma as continuous VFO that is “chopped up” by recurrent synaptic inhibition – something that is possible if axonal gap junctions are not too far from perisomatic sources of inhibition [41]. The model thereby accounts for the pharmacology, the field potential profiles, and the rare somatic firing (the latter because the action potentials that drive the oscillation are generated in axons, and only some of these successfully propagate back to the soma as full spikes). The model predicts that, during persistent gamma, axons fire at higher rates than somata; and that somatic action potentials are antidromic: these predictions have been experimentally verified [10].

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## 6.4 VFO and Origin of Seizures

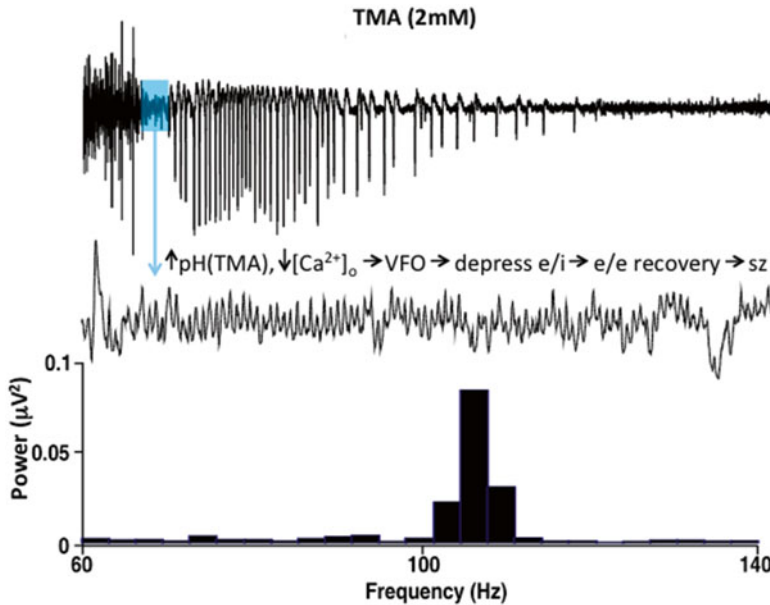
*Experimental VFO is potentiated by alkaline conditions.* A relation between systemic (and presumably brain) pH has long been suspected,

with alkaline pH being epileptogenic: in absence and other seizure types associated with spike-wave [12, 23], and in febrile seizures and their experimental models [29, 30]. In addition, some drugs with anticonvulsant properties, are blockers of carbonic anhydrase (acetazolamide, topiramate, zonisamide) [21, 25]. Remarkably, alkaline pH strongly potentiates *in vitro* VFO (Fig. 6.5, [8, 46]). The effects on VFO are unlikely to result from actions of pH on synaptic transmission, as the effects can occur when synaptic transmission is effectively blocked [46]. A likely cause is the opening of gap junction channels by alkaline pH [36], although it has not been possible yet to prove this directly.

*VFO and calcium.* A class of experimental *in vitro* epilepsy models includes so-called field bursts and related phenomena, in which ionic manipulations are used to suppress synaptic transmission (lowering [Ca<sup>2+</sup>]<sub>i</sub>, use of Mn<sup>2+</sup>), to increase neuronal excitability (for example, elevating extracellular [K<sup>+</sup>]<sub>o</sub>), and probably to open gap junctions with increased pH [13, 34, 38, 51]. Such field bursts likely (in our opinion) depend on gap junctions [26]. The occurrence of field bursts fits in with long-held hypotheses concerning a primary role for glia in epilepsy [9, 14, 37, 50]; and also with long-standing observations that afferent stimulation, as well as seizures themselves, can have significant effects on extracellular ion concentrations, including the lowering of [Ca<sup>2+</sup>]<sub>o</sub> [15].

*Experimental demonstration of gamma/VFO/seizure evolution in alkaline conditions.* Figure 6.6 illustrates a transition from VFO (~110 Hz) to electrographic seizure, suggestive of the human patient data of Figs. 6.3 and 6.4, although the data in Fig. 6.6 are from an *in vitro* hippocampal slice. The slice was bathed in an alkalizing solution, and then a tetanic stimulus was delivered that evokes an epoch of so-called tetanic gamma, during which [K<sup>+</sup>]<sub>o</sub> is expected to rise, and [Ca<sup>2+</sup>]<sub>o</sub> to fall [50]. The gamma is followed by VFO (middle trace in Fig. 6.6), that turns into a brief electrographic seizure. We propose that synaptic excitation of interneurons is depressed during the VFO period, analogous to





**Fig. 6.6** VFO-electrographic seizure transition *in vitro*, in alkaline conditions. Rat hippocampal slices were alkalinized with 2 mM trimethylamine (TMA), and a tetanic stimulus given to s. radiatum of CA1. In s. pyramidal field potential recordings, this stimulation resulted (upper trace, 17 s of activity) in post-tetanic gamma at first, then VFO (1), then ~2 mV epileptiform field transients (2), then further VFO (3). The middle

trace shows, on an expanded time scale (850 ms of activity), the ~0.1 mV VFO potential fluctuations, at ~110 Hz (power spectrum below). Our hypothesized sequence of events is shown above the VFO trace (see text for further details). “e/i”, pyramidal cell-to-interneuron; “e/e”, pyramidal cell-to-pyramidal cell. “sz”, seizure (From Traub et al. [42], reproduced with permission)

what has been shown in the preparation of Fig. 6.1 [44]; such synaptic habituation during VFO remains, however, to be shown directly.

## 6.5 Conclusion and Hypothesis

To summarize some of these data then, our view is that high-frequency firing in the pyramidal cell axon plexus is what drives both VFO and persistent gamma oscillations. **VFO occur under specific extracellular conditions – which we hypothesize to be the initiating factor for seizure activity.** What is now required, we believe, is direct measurement of extracellular tissue parameters [16], in epileptic patients, perhaps now using MRI [1, 17], and preferably in conjunction with EEG or ECoG recordings. If such measurements do indeed indicate, for example, tissue alkalinization just prior to seizure onset, it

will suggest alternative approaches to seizure prevention, and perhaps also better understanding of how present treatments – such as the ketogenic diet – are effective [3, 31].

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# Is There Such a Thing as “Generalized” Epilepsy?

# 7

Gilles van Luijtelaar, Charles Behr,  
and Massimo Avoli

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## Abstract

The distinction between generalized and partial epilepsies is probably one, if not the most, pregnant assertions in modern epileptology. Both absence and generalized tonic-clonic seizures, the prototypic seizures found in generalized epilepsies, are classically seen as the result of a rapid, synchronous recruitment of neuronal networks resulting in impairment of consciousness and/or convulsive semiology. The term generalized also refers to electroencephalographic presentation, with bilateral, synchronous activity, such as the classical 3 Hz spike and wave discharges of typical absence epilepsy. However, findings obtained from electrophysiological and functional imaging studies over the last few years, contradict this view, showing a rather focal onset for most of the so-called generalized seizure types. Therefore, we ask here the question whether “generalized epilepsy” does indeed exist.

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## Keywords

Idiopathic generalized epilepsies • Absence seizures • Generalized tonic clonic seizures • Myoclonic juvenile seizures • Spike and wave discharges • Genetic absence models

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## 7.1 Background

The concept of generalized and partial seizures dates back to conflict during the last century between “universalizers” and “localizers”, the former defending a holistic integrated view of brain function against the “centrencephalic system” of Penfield and Jasper (reviewed in [4]). It was Hughlings Jackson [64] who propose a distinction between generalized and partial seizures. Only much later was the term “generalized” epilepsy itself first employed by Gastaut [20]. Indeed, we will often refer in this chapter to generalized *versus* partial “seizures” rather than “epilepsies” since some experimental results may *stricto sensu* not be applicable to human epilepsy classification and thus they remain seizure-related material. However, since generalized epilepsies are defined as such because of the “generalized” nature of their concomitant seizures, any suspicion with regard to the “generalized” nature of these seizures, will automatically challenge the “generalized” nature of the corresponding epilepsy and *vice versa*.

Generalized seizures are characterized by sudden, often unexpected, manifestations (presumably reflecting the involvement of the entire brain, or at least a large part of the brain) compared to the slower, clinically heterogeneous partial seizures where the patient often remains conscious, at least at the beginning of the seizure. With the development of EEG recordings, this assertion received a formidable confirmation [7]. The electroencephalographic manifestations accompanying generalized absence seizures consist of highly stereotyped pattern of bilateral synchronous, regular and rhythmic spike and wave (SW) discharges at 2.5–4 Hz in children, juveniles and adults, lasting from a few seconds up to 30 s. In contrast, scalp EEG recordings reveal sustained diffused, synchronous, discharges during the *tonic* phase, and interrupted bursts during the *clonic* stage in generalized tonic-clonic seizures (GCTS).

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## 7.2 Evolution of the Classification

Classification in epileptology is a work-in-progress and a simple examination of the past 50 years reveals how the Jacksonian dogma has evolved. In the 1969 classification [19], emphasis was put on the distinction between “seizure that are generalized from the beginning and those that are focal or partial at onset and become generalized secondarily”. In the 1981 classification, it was proposed that generalized seizures have electroclinical patterns that “presumably reflect neuronal discharge which is widespread in both hemispheres”, underlying a conceptual shift from bilateral “onset” to bilateral “spread” [57]. In 1989, a new classification postulated that partial localization-related epilepsies are “epileptic disorders in which seizure semiology or findings at investigation disclose a localized origin of the seizures”, whereas generalized epilepsies are defined by initially bilateral ictal encephalographic patterns [56]. The “generalized” designation was essentially an electroclinical feature, which was discarded in 2010 when terminology was revised.

Indeed, in 2001 and 2006, an ILAE Task Force debated the relevance of this conceptual dichotomy [15, 16]. Recognizing that it was out of date with regard to pathophysiological advances in epileptology, the commission decided to keep its core concept in the 2010 revised classification for convenience. Hence it was proposed that “Generalized and focal are redefined for seizures as occurring in and rapidly engaging bilaterally distributed networks (generalized) and within networks limited to one hemisphere and either discretely localized or more widely distributed (focal)” [6]. Today this convenient scheme, presumably aimed at distinguishing between epilepsies that are recommended for surgery (“surgical”) and others (“non-surgical” epilepsies), represents the first issue to be addressed when diagnosing a person with epilepsy. However, it deserves re-evaluation in the light of recent advances obtained from clinical and basic research studies.

### 7.3 Absence Epilepsy: From the Centrencephalon to the Thalamo-cortical Loop

There is presently compelling evidence based on brain imaging, EEG recording and signal analysis techniques that a key element of generalized epilepsies, the sudden involvement of the whole brain, is highly disputable in typical absence epilepsy [4]. Paradoxically, one of the first electrophysiological studies of the pathophysiology of generalized SW discharges: the Jasper and Droogleever-Fortuyn paper, already suggested a mechanism for focal onset of absence seizures [34]. These authors succeeded in inducing typical 3 Hz SW discharges by local 3 Hz stimulation in the midline and intralaminar nuclei of the thalamus. However, in spite of this evidence, the Montreal school [53] preferred the integrative hypothesis of the centrencephalic system presumably influenced by the recent discovery of the reticular formation [47]. Later, Gloor's team further explored the corticoreticular nature of generalized SW discharges, using the feline generalized penicillin epilepsy model, introduced by Prince and Farrell [55], and established the link between sleep spindles and SW discharges (reviewed in [35]). Those data were still consistent with the thalamo-cortical origin for SW activity and did not challenge or question the "generalized" character of absence epilepsy. Indeed, *in vitro* studies in ferret brain slices and computational models confirmed later that thalamo-cortical oscillations could be driven by an intrathalamic circuit and revealed that some cellular properties of thalamic cells, which are involved in sleep spindles, most likely contribute to SW generation [29].

During the 1980s, in parallel with the growing interest for a genetic etiology for the so-called "idiopathic" epilepsies, animal models with genetic inheritance of absence epilepsy were described. Specifically, both "genetic absence epilepsy in rats from Strasbourg" (GAERS) [73] and "Wistar Albino Glaxo/Rijswijk" (WAG/Rij) rats [68] were identified. For the first time these

models provided the opportunity to directly test hypotheses in animals presenting with spontaneous absence seizures [13, 43, 69]. Both the *in vitro* studies mentioned above [29] and the evidence obtained from genetic models led to the idea of a thalamo-cortico-thalamic network in which the typical SW discharges could elicit spontaneously. This network included the thalamic reticular nucleus in which inhibitory interneurons trigger GABAergic IPSPs on thalamic relay cells. T-type  $\text{Ca}^{2+}$  current are deinactivated, as a consequence of hyperpolarization causing a burst of action potentials that in turn excited both reticular thalamic and cortical cells [5, 14, 31, 32, 52]. Enhanced T-type  $\text{Ca}^{2+}$  currents were recorded in GAERS reticular thalamic cells and thalamic relay cells [67]. In addition, subtle abnormalities in GABAergic transmission were found in the reticular thalamic nucleus of GAERS compared to control rats [8]. However *in vivo* studies using a different model, namely a feline Lennox-Gastaut model, allowed Steriade and coworkers to pinpoint a cortical trigger for the SW discharge, reopening the controversy about whether the cortex or the thalamus were to be responsible for the generation of SW discharges [62].

### 7.4 A Neocortical "Focus" as Trigger of Generalized Absence Seizures

More than 50 years of research were necessary to decrypt the pathophysiology of the thalamo-cortical loop in absence seizures, meaning how this neuronal circuit could "jump" from the physiological sleep spindle to the pathological SW discharge. But the real trigger for an absence seizure was hiding somewhere else, and with its discovery the concept of "generalized" absence epilepsy ended. This discovery began in the 1960s, when it was reported that there was focal neocortical initiation of absence seizures. Marcus and Watson [41] discovered that bilateral application of pro-convulsant drugs to the frontal cortices could produce a pattern of generalized SW discharges

similar to what was observed during an absence seizure. In the 1970 and 1980s, topographical EEG studies performed in patients with absence epilepsy confirmed that SW discharges did not occupy the entire cortex. The *wave* component of the SW discharge was characterized by a maximum localized in frontal areas [58].

These data were soon confirmed in the GAERS model with intracellular recordings of thalamic relay cells where an excitatory drive (EPSCs) was shown, presumably originating from the cortex [54]. The role of the cortex in initiating SW discharges was further established *in vitro* by demonstrating that cortico-thalamic input strength is critical for thalamo-cortical rhythmic activity and for changing a spindle into a SW oscillation [9]. But the evidence that was most compelling ultimately came from experiments performed in the WAG/Rij model where cortico-sub-cortical multiple-site EEG signals were studied using non-linear association analysis [45]. The authors identified a consistent initiation zone in the peri-oral region of the somatosensory cortex (Fig. 7.1), along with a leading role of the cortical projections to the thalamus lasting for the initial 500 ms of the SW discharge. Furthermore, the high degree of bilateral synchronization, characteristic of generalized SW discharges in absence epilepsy, appeared to rely mainly on cortico-cortical connectivity, as indicated by non-linear analysis of inter, intra and thalamo-cortical relationships.

Both initiating and leading roles of the neocortex were further suggested by data showing that SW discharges could be recorded in this structure without concomitant SW activity in thalamus, while the opposite situation never occurred [45, 70]. However, once a seizure evolved, both structures oscillated in concert, suggesting a stereotypical scenario where the *primum movens* involves the peri-oral somatosensory cortex, which secondarily “switches on” the thalamo-cortical loop. The primary role of neocortex in the initiation of SW activity was later confirmed in GAERS rats; in these experiments, Manning and co-workers [40] found that local application of the anti-absence drug ethosuximide has maximal efficacy when this drug is applied into the primary somatosensory peri-oral region while its infusion into the thalamus produced only minor and

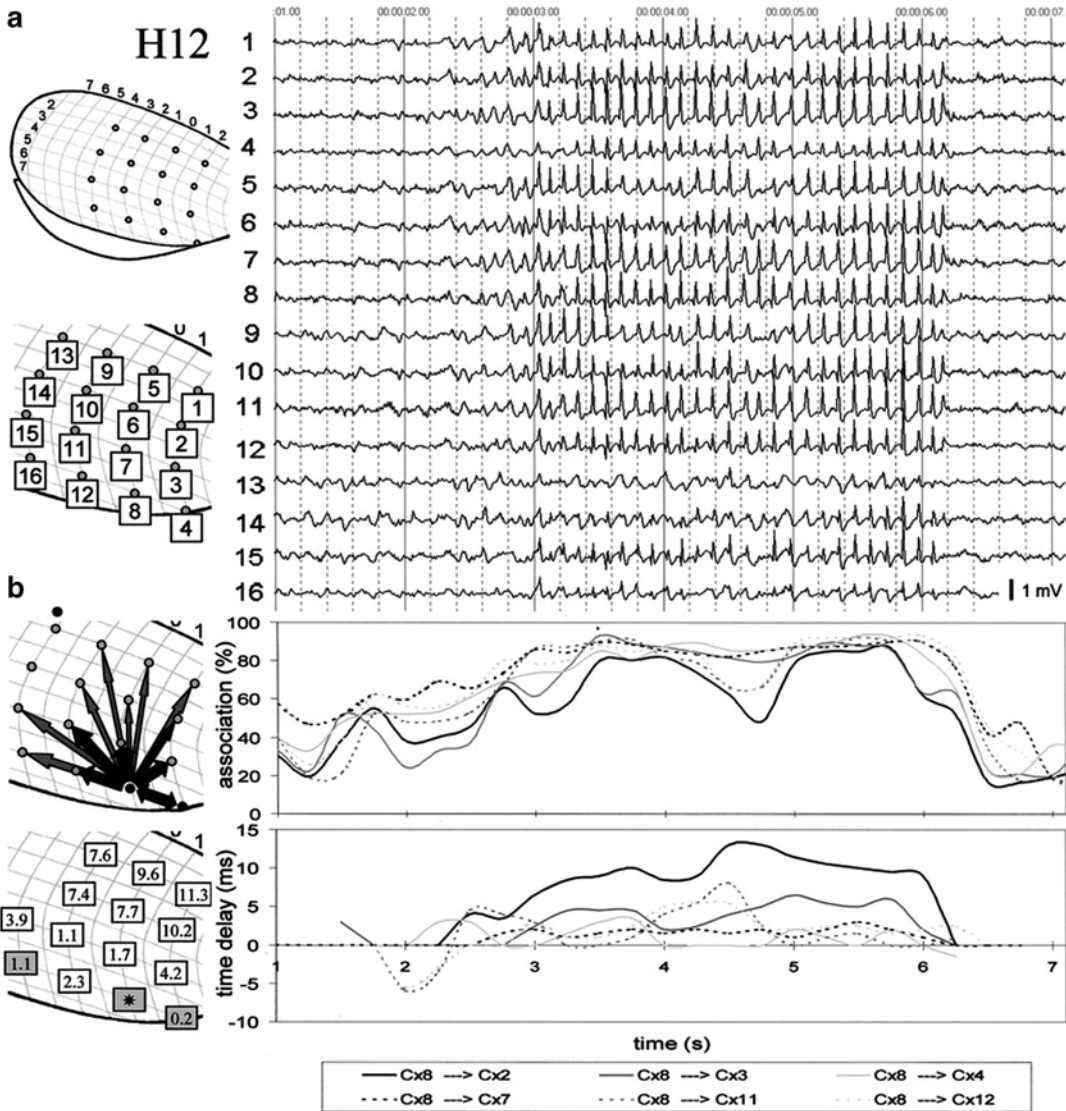
delayed reduction in SW discharges. Thus, SW discharge initiation in the two major genetic models of absence epilepsy occurs in the same restricted area of the somatosensory cortex. Why and how, however, remain to be clarified.

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## 7.5 Focal Cortical Origin of Absence Seizures in Humans

Human studies on ictal generalized discharges in absence epilepsy have shown that patterns of activation and deactivation identified by fMRI are restricted to some cortical (medial frontal cortex, precuneus, lateral parietal, and frontal cortex) and subcortical regions (thalamus, brainstem) [1, 11]. Due to the limited temporal resolution of fMRI, these results did not allow clear confirmation of a cortical initiation site. However, high-resolution EEG and MEG studies in combination with advanced signal analytical techniques have confirmed the existence of a preferential cortical origin of SW discharges. Localized sources were detected either in the frontal cortex, orbito-frontal, medial temporal or parietal lobe [27, 65, 74]. In addition, a rather localized preictal SW rhythm of low frequency (3 Hz) was detected in atypical absence patients [24], whereas a reproducible topography of locally synchronous cortical sources with increased local connectivity was described in a multifocal network, comprising the right prefrontal mesial, left orbito-frontal and left lateral post-central area [2].

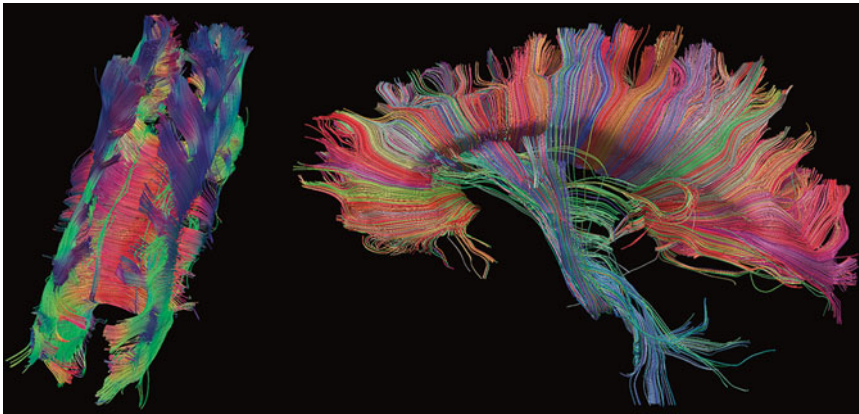
Absence seizures may indeed appear “bilateral and synchronous” (and thus “generalized”) in EEG recordings because of the highly connected inter- and intra-cortical networks that are sustained by a cortico-thalamic-cortical loop thus leading to oscillatory activity. However, the SW discharges appear to originate from specific cortical areas. The velocity of spread between hemispheres is presumably based on extensive monosynaptic inter-hemispheric connections via the corpus callosum (Fig. 7.2). Evidence for the role of callosal interhemispheric connectivity came from callosotomy experiments resulting in the disruption of the bilateral and synchronous SW discharges in several absence seizure and genetic animal models of absence epilepsy [41, 42, 48, 72].



**Fig. 7.1** Evolution of the intra-hemispheric cortico-cortical association strength  $h^2$  (a non-linear correlation coefficient between two signals) was calculated for all electrode combinations as a function of time of shift between the signals) and time delays of the local field potentials signals between electrode pairs. A cortical grid covering a major part of the somatosensory area was used for electrographical seizure recording in 16–22 month old WAG/Rij rats with spontaneous occurring spike-wave discharges. (a) *Left*: Electrode positions (*top*) and electrode labels (*bottom*) on the somatosensory cortex of rat H12. The numbers on the *top graph* refer to coordinates based on the rat’s anatomical brain atlas of Paxinos and Watson. *Right side* refers to frontal. (a) *Right*: A typical 3 s lasting electrographical spike-wave discharge recorded (with negativity up) with the cortical grid that covers a great part of the lateral neocortex

with position of the electrodes and their labels on the *left*. (b) Time courses of the cortico-cortical nonlinear associations (*top panel*) and time delays (*bottom panel*) for several sites (as indicated by the *black arrows* on the *left*) with respect to the focal site (electrode 8). The association and time delays were assessed for successive 50 % overlapping 500 msec epochs. For comparison the pictures on the *left* depict the average overall associations (*top*) and the average overall time delays (*bottom*; in milliseconds). There is a gradual increase in association strength before the start of the seizure and a steep drop in association strength at the end. Before the seizure, time delays are inconsistent, and there is often a zero time lag. During the seizure, time delays are always positive indicating a delay at the different electrode positions compared to position 8, although the magnitude of the delay can vary [45]

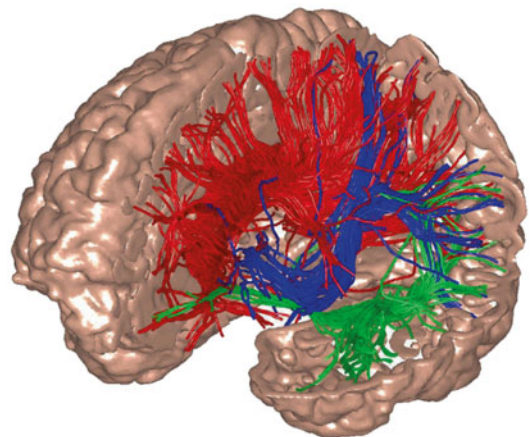




**Fig. 7.2** Diffusion tensor imaging (DTI) of the human corpus callosum colored by end point location. *Left*: viewed from the *top*; Anterior side points to the *bottom*. *Right*: lateral view with anterior part pointing to the *left*.

Notice the massive cross- hemispheric projections through which seizures might get quickly “generalized” (From Tromp [66]. Reprinted with permission from the author)

The cortex is endowed with a variety of excitatory neocortical projection neurons that play a role in the quick information transfer between homotopical regions of the two hemispheres via long myelinated axons through the corpus callosum. Homotopic regions include the somato-sensory cortex [75], providing an anatomical explanation for fast spread and bilateral involvement shown by the electrophysiological results [45]. In addition to their role in integrating homotopic neocortical regions, callosal projection neurons are also responsible for information transfer within each hemisphere [17]. Finally, there is an abundant and widespread thalamo- cortico-thalamic network; the descending projections to the thalamus “are estimated to outnumber thalamo-cortical ones by an order of magnitude” [61]. The extensiveness of the cortico-thalamo-cortical network, visualized with white matter tractography, can be appreciated in Fig. 7.3. The speed of involvement of the thalamus and cortical spread after local cortical initiation is undoubtedly mediated through these massive networks.



**Fig. 7.3** Large-scale model of mammalian thalamo-cortical system based on DTI scans. The massive reciprocal connections between cortex and thalamus are responsible for the quick propagation of SW discharges and other electroencephalographic markers of “generalized” epilepsies from their cortical sites of origin to the thalamus and back. In the illustration, left frontal, parietal, and a part of temporal cortex have been cut to show only a small fraction of white-matter fibers, color-coded according to their destination. *Red*: projections to the frontal cortex, *blue* to parietal cortex, *green* to temporal cortex implying different sources of reentrant axonal fibers connecting one part of the cortex to another [30]

## 7.6 Generalized Convulsive Seizures

Aside from the archetypical absence epilepsy, a rather heterogeneous ensemble of syndromes constitute the group of primary (idiopathic)

generalized epilepsies such as Juvenile Myoclonic Epilepsy, Juvenile Absence epilepsy, and Lennox-Gastaut syndrome, to mention only a few. If growing evidence points to the focal onset for typical absence epilepsy, what

about other types of generalized epilepsies? Our pathophysiological understanding of idiopathic generalized epilepsies mainly relies on the ability of the aforementioned and incriminated networks to generate paroxysmal discharges. This susceptibility would thus result from the combination of a paroxysm-inducing mechanism such as arousal (“dyshormia”) or photosensitivity and a genetically-prone network [50]. Myoclonus, absence and ultimately GTCS would thus represent a crescendo of clinical manifestations related to this genetic predisposition for generalized paroxysms.

### 7.6.1 Generalized Myoclonic Seizures

Myoclonus, on the one hand, can be either focal or generalized, and of either cortical, thalamo-cortical, reticular reflex or negative nature, i.e., characterized by the inhibition of muscular activity [49]. In idiopathic generalized epilepsy, such as juvenile myoclonic epilepsy and absence myoclonic epilepsy, myoclonus is supposedly of thalamo-cortical nature and is associated with generalized EEG discharges. Interestingly mild peri-oral myoclonus has also been described in typical absence epilepsy [26], thus concerning the same cortical regions supposedly driving SW discharges in the genetic absence models [45].

Experimentally, myoclonic seizures can be triggered in rodents, either by electrical or pharmacological stimulation by GABA<sub>A</sub> receptor antagonists such as bicuculline, picrotoxin and pentylentetrazole or flurothyl; any of these procedures induce GTCSs with an initial, variable myoclonic phase, thus being slightly different from human generalized myoclonic epilepsy. Local application of most pro-convulsant drugs onto the cortex also elicits myoclonus of focal origin [71]. In a genetic model such as the photosensitive *Papio papio* baboons, generalized myoclonic discharges appear to start in the fronto-rolandic cortex [18]. In human idiopathic generalized epilepsies, evidence for asymmetry, asynchrony and ultimately focal onset of EEG generalized discharges has been gathered through the years; patients with heterogeneous primary generalized epilepsy have been studied using

repetitive EEG showing the consistent presence of focal features [39, 50]. A restricted cortical network has been described during typical “generalized” 4–6 Hz seizure propagation in juvenile myoclonic patients; this includes regions of frontal and temporal cortex [28]. Another study using Jerk-locked averaging in JME patients pinpointed a frontal cortical generator [51]. The association of JME with some particular personality type, as described by Janz [33], has been related to fronto-cortical disturbances. Neuropsychological studies confirmed verbal and visual memory impairment along with disturbed visuospatial processing and working memory alteration [63].

Neuropathological studies have also revealed microdysgenesis in idiopathic generalized epilepsy [44]. Hence, these results highlight an early cortical involvement in juvenile myoclonic epilepsy. FMRI studies in patients with idiopathic generalized epilepsy, again due to the poor temporal resolution of fMRI, have failed to demonstrate early focal activation. However, a consistent pattern of thalamic activation and cortical default-mode network deactivation were described during idiopathic generalized epilepsy [22, 38, 46], suggesting a common pathophysiology for generalized SW discharges among idiopathic generalized epilepsies. Suspension of default-mode network represent the earliest BOLD signal change to be observed and may thus hide a more discrete cortical onset, whereas later thalamic activation account for the sustained SW discharge. And indeed, BOLD pattern recorded during photoparoxysmal generalized discharges, a rather cortical electroencephalographic trait of photosensible generalized epilepsy, does not concern thalamus [46].

### 7.6.2 Primary Generalized Tonic Clonic Seizures

Primary GTCS, the third major type of seizure present in generalized epilepsy is classically differentiated from secondary GTCS that occur in partial epilepsy. Use of the same term, for both primary and secondary GTCS seems contradictory despite clinical similarities, since pathophysiology

is obviously rather different. MEG studies have shown some discrepancies among the two types of GCTS with regards to levels of close and distant ictal synchronization. Distant synchrony appears higher in primary GTCS whereas increased local synchrony is reported in secondary GCTS [23]. Additionally interhemispheric coherence during secondary GCTS is surprisingly low [21], and variable during the time-course of a seizure [36]. A single-photon positron emission tomography study comparing spontaneous GCTS with electroconvulsive-therapy-induced GCTS showed specific fronto-parieto-temporal along with thalamic activation in bilaterally electroconvulsive-therapy-induced GCTS [10]. Infantile “generalized” spasms can be focal in its etiology [12], and even involve the brainstem, as it is clear from generalized symmetric seizures in hemispherectomized children [37].

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## 7.7 Conclusive Remarks

Focal versus generalized epilepsy is a classical dichotomy inherited from the Jacksonian era, and somewhat confirmed by standard EEG. However, increasing evidence from both structural and functional imaging studies has been gathered though the years to call into question the concept of generalized epilepsy. Recent studies on rodent genetic absence models [40, 45] prove that absence epilepsy, considered as the prototype for generalized epilepsy, may originate from a rather focal, cortico-frontal region. In human, recordings using high density EEG/MEG studies with proper signal analytical techniques [27, 28, 60, 74] also revealed focal features. The traditional view of a widespread recruitment in absence seizures has also been contradicted by fMRI studies demonstrating a rather restricted network of activation and deactivation, mainly corresponding to alterations in the default-mode network [1, 3, 25, 46, 59]. Refuting the concept of generalized epilepsy however, remains almost impossible *per se*, as it consists of hundreds of different epileptic syndromes. Looking for a focal onset in all those syndromes is not realistic and probably unnecessary.

Most studies on “generalized epilepsies” mainly include patients with idiopathic generalized epilepsy as this group represent the majority of so-called generalized epilepsy.

Overtaking the “centrencephalic” theory of the past century bares new ideas about the nature of generalized epilepsies. It is a safe bet that avoiding the use of the term “generalized epilepsy” will benefit for the next generation of epileptologists and patients. Future classifications based on networks properties, along with more specific information about etiology may decrease the emphasis on the classical electro-clinical distinction of partial vs. generalized epilepsy. Nowadays the remaining distinction will thus reflect differences in terms of spreading velocity properties of the underlying network rather than of that network size itself. But as conceptual evolution has a tendency to spread rather slowly in the medical community, one can predict that epilepsy will remain to be characterized as either partial or generalized for some time to come.

**Acknowledgments** We dedicate this manuscript to Phil Schwartzkroin, a friend and colleague, who had a great impact on neurobiological research and on our own studies during the last four decades. He created opportunities for others to promote new concepts and theories. His challenging approach to epilepsy research continues to motivate many of us, and it is indeed mirrored by the question addressed here.

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## Part II

# Synaptic Plasticity

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# Are There Really “Epileptogenic” Mechanisms or Only Corruptions of “Normal” Plasticity?

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Giuliano Avanzini, Patrick A. Forcelli,  
and Karen Gale

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## Abstract

Plasticity in the nervous system, whether for establishing connections and networks during development, repairing networks after injury, or modifying connections based on experience, relies primarily on highly coordinated patterns of neural activity. Rhythmic, synchronized bursting of neuronal ensembles is a fundamental component of the activity-dependent plasticity responsible for the wiring and rewiring of neural circuits in the CNS. It is therefore not surprising that the architecture of the CNS supports the generation of highly synchronized bursts of neuronal activity in non-pathological conditions, even though the activity resembles the ictal and interictal events that are the hallmark symptoms of epilepsy. To prevent such natural epileptiform events from becoming pathological, multiple layers of homeostatic control operate on cellular and network levels. Many data on plastic changes that occur in different brain structures during the processes by which the epileptogenic aggregate is constituted have been accumulated but their role in counteracting or promoting such processes is still controversial. In this chapter we will review experimental and clinical evidence on the role of neural plasticity in the development of epilepsy. We will address questions such as: is epilepsy a progressive disorder? What do we know about mechanism(s) accounting for progression? Have we reliable biomarkers of epilepsy-related plastic processes? Do seizure-associated plastic changes protect against injury and aid in recovery? As a necessary premise we will consider the value of seizure-like activity in the context of normal neural development.

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## Keywords

Seizure • Epilepsy • Epileptogenesis • Neural plasticity • Synaptic plasticity

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The first evidence for experimentally induced plastic changes in nervous system was provided by Minea in 1907. In his thesis, he described metamorphic phenomena in sensory neurons provoked by the compression and transplantation of ganglia into various organs [84, quoted by Marinesco 1909]. In the years that have passed, the study of neural plasticity has become a very important line of research in the neurosciences, with the aim of uncovering the neurobiological bases for the exquisite capability of the nervous system to adapt to environmental changes. Plasticity in the nervous system, whether for establishing connections and networks during development, repairing networks after injury, or modifying connections based on experience, relies primarily on highly coordinated patterns of neural activity. Rhythmic, synchronized bursting of neuronal ensembles is a fundamental component of the activity-dependent plasticity responsible for the wiring and rewiring of neural circuits in the CNS. It is therefore not surprising that the architecture of the CNS supports the generation of highly synchronized bursts of neuronal activity in non-pathological conditions, even though the activity resembles the ictal and interictal events that are the hallmark symptoms of epilepsy. To prevent such natural epileptiform events from becoming pathological, multiple layers of homeostatic control operate on cellular and network levels. While there are extensive data concerning the plastic changes that occur in brain structures during the process of epileptogenesis, the role of these changes in counteracting or promoting epileptogenic processes remains controversial.

In this chapter we will review experimental and clinical evidence on the role of neural plasticity in the development of epilepsy. We will address questions such as: is epilepsy a progressive disorder? What do we know about mechanism(s) accounting for progression? Have we reliable biomarkers of epilepsy-related plastic processes? Do seizure-associated plastic changes protect against injury and aid in recovery? Moreover, as a necessary premise we will consider the value of seizure-like activity in the context of normal neural development.

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## 8.1 Modeling and Remodeling of Network Architecture During Development

The architecture of neuronal connectivity in the CNS is shaped through a process of functional validation. During postnatal development, this process depends primarily on environmental input and sensory stimulation, while during prenatal development, spontaneous patterned activity is largely generated intrinsically. Neuronal activity is essential for guiding synapse formation, remodeling, and elimination, so as to establish optimal connectivity. For example, well in advance of eye opening, embryonic retinal ganglion cells generate rhythmic bursts of action potentials in both rodent [83, 127] and primate [70]; this highly correlated bursting is required for establishing retinotopic maps in the connections across the neuraxis. This activity is highly synchronized within populations of neighboring neurons, and propagates throughout the visual pathway [2], so that waves of stimulation in defined regions of the retina can then coordinate the activity-dependent refinement of corresponding eye-specific layers in the lateral geniculate nucleus [83, 127]. Moreover, the spontaneous bursting is relayed via thalamocortical projections to visual cortex, where it can shape the emerging ocular dominance columns [52]. A similar pattern of correlated bursting activity occurs pre-hearing in the developing auditory system, from the level of the cochlear ganglion cells to the brainstem auditory pathways [57, 58]. These spontaneous, highly synchronized and propagated rhythmic bursting patterns, which share many characteristics of ictal and interictal events, are a classic example of the developmental principle that neurons that fire together, wire together [23, 47].

Another classic example of highly synchronized rhythmic bursting of neuronal populations in utero takes place in spinal cord motoneurons. In fact, the spontaneous waves of hypersynchronous activity in this system have been characterized as epileptiform activity [101]. These discharge patterns propagate through the spinal cord,

triggering transregional synchronization and fast rhythmic repetitive limb movements described as clonus and convulsive-like in nature [14, 94]. While clearly a normal and adaptive feature of prenatal development, a similar pattern of activity in a postnatal organism would be considered pathological.

Before discussing the role of plastic mechanisms in epileptogenesis, we will review the evidence for a progressive course of epilepsies. For the purposes of this chapter we will rely to the following arbitrary definitions:

*Epileptic mechanisms* responsible for seizure generation consist of changes in cellular excitability leading to excessive, disordered discharges underlying ictal manifestation.

*Epileptogenic mechanisms* responsible for epilepsy, i.e. an enduring propensity to generate epileptic seizures [32] consist of some hypothetical neurobiological processes leading to a permanently dysfunction of the neuronal network/system.

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## 8.2 Epilepsy as a Progressive Disorder

Clinical and experimental observations suggest that an acute “initial event” (e.g. traumatic, infectious) can set in motion a series of degenerative, regenerative and inflammatory changes resulting in a permanent epileptic neuronal aggregate. A crucial role is attributed to the epileptic activity both in the initial event (e.g. febrile seizure/status) and in the ensuing process leading to chronic epilepsy. Evidence in some patients for a progressive increase in the risk for seizures with increasing number of seizures is currently quoted in support of Gower’s statement that seizures beget seizures [24]. Indeed, in support of the notion that seizures beget seizures, Hauser and Lee [44] found a significant increase in risk for subsequent seizures with increasing seizure number in a population of patients who are generally considered to have a good prognosis for going into remission: those with unprovoked seizures of unknown cause, normal neurological examination,

and normal EEG. It is worth saying, however, that there are many types of epilepsies that do not progress, in spite of seizure repetition. These non-progressive epileptic syndromes include benign childhood epilepsies with centrotemporal spikes [136], benign occipital epilepsies, childhood and juvenile absence epilepsies [137], juvenile myoclonic epilepsy, benign familial neonatal, infant and neonatal-infant epilepsies [124], and autosomal dominant nocturnal frontal lobe epilepsies. For many of these syndromes seizure activity decreases or disappears with age [40, 136].

A progressive course toward drug refractoriness can be observed only in some types of human epilepsies currently grouped under the definition of epileptic encephalopathies (EEs) [6]. As for mesial temporal lobe epilepsy, a role of repeated seizures for inducing progressive cumulative alterations in neural circuits, resulting in progression of epilepsy severity, has been assumed (based on experimental results with rodent kindling models) but never demonstrated. Patients with mesial temporal lobe epilepsy (MTLE) and EEs substantially contribute to the 30–40 % of patients with epilepsy who show drug resistance [27]; this population represents the main unsolved problem in clinical epileptology. This explains the great investment in research lines aimed at unraveling the neural mechanisms responsible for these types of epilepsies and the need to elaborate strategies capable of preventing their development.

*The latent period.* In several instances the natural history of MTLE indicates an initial precipitating event as an underlying cause of a chronic epilepsy [79, 80]. In some cases (trauma, infection, autoimmune process), the initial event is associated with repeated seizures often presenting as status epilepticus (SE). Moreover, a significant proportion of patients with mesial temporal sclerosis and MTLE had antecedents of complex febrile seizures in early childhood [19]. Between the initial event and the onset of the chronic epilepsy there is an interval of variable duration currently referred to as latent period. During this period, biological changes may occur that are considered to substantiate the

epileptogenic process. For these reasons much interest is focused on animal models that reproduce the typical sequence of initial event-latent period-chronic epilepsies.

The pilocarpine and kainic acid models are both obtained by the acute administration of a chemoconvulsant agent (pilocarpine or kainic acid) to rodents; this treatment induces a state of prolonged SE, followed by spontaneous recurrent seizures beginning after a variable latency (15–20 days) [18, 92, 128]. It must be said that the existence of the latent period in chemoconvulsant rodent models has been disputed [118] and that, in view of the inter-individual variability of its duration, the possibility that it simply reflects the outer fringe of a probabilistic spread seizure latencies must be taken into account. Indeed an impressive bulk of published results suggest that during the latent period several changes occur in hippocampal structures that are associated with the alteration of excitability and synchronization and may hypothetically account for epileptogenesis. For example, axonal sprouting, synaptic reorganization, gene and protein expression, neurogenesis, gliosis and functional glial alterations, inflammation, and angiogenesis have all been suggested to contribute to epileptogenesis in these models (see [125] for a review). Interestingly these changes can also be found in temporal lobe tissue samples from patients who underwent epilepsy surgery for drug refractory MTLE (see [22] for a review). Obviously, the fact that these changes occur during an ongoing epileptogenic process does not prove that they are necessary contributors to disease pathogenesis until the prevention of any of them is unequivocally proved to prevent the development of later epilepsy. For example, the mossy fiber sprouting that characteristically occurs after SE and is thought to be a hallmark feature of the post-SE neuroplasticity, has been demonstrated not to be necessary for the development of spontaneous recurrent seizures [25, 48, 86, 95, 140]. Moreover the histopathological alterations observed in pilocarpine and kainic models are not limited to the mesial temporal lobe structures, raising a question about their validity as MTLE models (e.g., [17, 21, 66, 129]).

From the clinical standpoint, it is not clear that MTLE results from a process that is sustained or facilitated by epileptic activity. The analysis of the natural history of MTLE cannot answer the question of whether unfavorable outcomes are due to the persistence of epileptic activity (which is usually undetectable in the latent period between the initial event and the chronic phase), or if it is instead a product of the underlying neuroplasticity set in motion by the initial event. While previous prospective longitudinal analyses have been inconclusive [113], the results of the ongoing FEBSTAT (Consequences of Prolonged Febrile Seizures) prospective study [72, 90] may clarify this issue. Several studies have shown that only symptomatic SEs correlate with brain damage and late epilepsy [45, 64, 103, 114]. This makes it impossible to point to a necessary role of epileptic activity above and beyond (or apart from) the role of underlying lesion causing SE for the initiation and maintenance of the epileptogenic process [46].

The statement “seizures beget seizures” implies a role of epileptic activity not only in initiating the epileptogenic process but also in maintaining it, suggesting a further progression of epilepsy toward a more severe state. Whereas there is some evidence consistent with possible acute seizure-associated epileptogenic changes (see review in [15]), a subsequent correlation of recurrent seizures and progression to the clinico-pathological picture of drug refractory MTLE with hippocampal sclerosis has not been unequivocally demonstrated. A prospective analysis of 103 patients with newly diagnosed focal epilepsy [107] showed a decrease in hippocampal volume after 1–3 years in 13 % of patients. Here again, the coexistence of uncontrolled seizures and hippocampal atrophy in a limited subset of patients is insufficient to prove a clear causal relationship between recurrent seizures and atrophy. In fact, sporadic reports on acute, possibly inflammatory, damage to the hippocampus (localized limbic encephalitis; [11] and febrile SE [113]) suggest that acute seizures and hippocampal damage can develop within weeks/months, resulting in a MTLE pattern [59, 121] with little, if any, signs of further progression of

structural damage as detected by imaging, in spite of the persistence of seizure activity [71].

Animal models of MTLE based on acute induction of SE by pilocarpine and kainic acid do not clarify this issue, because once the spontaneous recurrent seizures have fully developed in the late chronic period, they do not tend to worsen. In fact, in a similar model of focal tetanus toxin-induced spontaneous seizures in hippocampus, the spontaneous seizures last for about 6 weeks and then tend to subside [55]. More recent data obtained in an animal model of post-traumatic epilepsy, indicates that following brain injury induced in mice or rats by a controlled cortical impact (CCI), once spontaneous seizures appear, they maintain a fairly constant frequency and severity and do not appear to worsen over an extended time period [13].

It is likely that numerous types of plasticity accompany the process by which epilepsy develops. For example, in addition to changes that occur in association with the primary site of epileptogenesis, there is experimental evidence for secondary epileptogenesis occurring in distant sites as a result of the abnormal activation of synaptic projections coming from the primary site (primary focus). Thus, the primary focus induces similar paroxysmal behavior (secondary focus) in the cellular elements of the otherwise normal network [87]. This may explain some of the cases in which epileptic seizures appear to become more severe, as a result of recruiting additional pathways into the network of seizure propagation.

At the same time, some of the plasticity associated with repeated seizure activity may give rise to compensatory processes that serve to limit or prevent the development of chronic state of seizures or epilepsy. Similarly, other aspects of the plasticity may compensate for the original damage that triggered the seizure activity. These types of plasticity and their mechanisms have been investigated in experimental animals in which repeated brief seizures (induced by electrical kindling stimulation, focal tetanus toxin, or electroshock treatments) cause little or no neuronal damage [10, 38, 56, 68, 78, 82, 130]. These repeated seizures have been shown to activate a host of genomic responses in the adult brain,

ranging from immediate early genes, genes for neurotrophic factors and neuropeptides, as well as multiple alterations in the regulation of neurotransmitters and their receptors in various brain regions [5, 12, 33, 41, 53, 65, 67, 73, 85, 88, 89, 97, 102, 104, 106, 110, 117, 126, 139]. In most cases, these responses include the induction or modulation of numerous trophic and neuroprotective factors such as bFGF[(FGF-2) [41]], NGF and BDNF [5], GDNF [3], and heparin-binding EGF-like growth factor [115] in various brain regions. These factors are responsible for triggering neuroplasticity, synaptogenesis and even neurogenesis, at the same time that they confer resistance to injury.

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### 8.3 Neuroprotection, Repair and Recovery After Brain Injury

A consequence of the induction of trophic and neuroprotective factors following repeated non-injurious seizures is a dramatic neuroprotective state in which the seizure-exposed animals become resistant to neuronal damage as evidenced by histopathology and sensitive molecular markers of cell death [63, 66, 78, 98]. Because neuroprotection requires multiple seizure treatments over several days (a single treatment is not protective), there is likely to be a cumulative buildup of resistance to injury. This suggests that seizures in the adult brain may be an endogenous therapeutic mechanism to recruit trophic cascades that promote neuronal survival and recovery in the face of degenerative insults or traumatic injury.

It is noteworthy that the type of seizures that are effective in conferring a neuroprotective effect on forebrain regions are seizures that last only several seconds and engage limbic forebrain networks either through kindling [63, 98] or by minimal electroshock administered via corneal electrodes [66, 78]. These stimuli produce characteristic signs of limbic-motor seizures (facial and forelimb clonus with rearing in the rodent), typically without the tonic-clonic motor manifestations characteristic of seizures that have spread to brainstem seizure-generating sites. In sharp contrast, repeated exposure to seizures that only engage brainstem seizure circuitry

(and not forebrain limbic networks) and evoke only tonic-clonic motor responses (without limbic-motor seizures) [16] fails to confer neuroprotection and may even worsen neural injury [4]. These observations emphasize the network-specificity of the protective responses.

Consistent with this concept of network specificity, the regional induction of mRNA for bFGF is very different following maximal electroshock seizures as compared to low-intensity (minimal) electroshock seizures, even when the two types of seizures are induced by corneal electrodes [33]. The minimal seizures increased bFGF mRNA levels by 350 % in entorhinal cortex by 5 h, whereas at the same timepoint after maximal seizures, the increase was only 200 %. Similarly, the increase in bFGF mRNA in hippocampus was greater after minimal seizures than after maximal seizures. In contrast, maximal seizures, but not minimal seizures, induced increases in bFGF mRNA in striatum and cerebellum [33]. This suggests that the minimal seizures, which are more selective for activating limbic forebrain networks may be more efficacious in triggering neuroplastic changes in those networks as compared with more generalized seizures. The circuit-specificity of seizure-induced neuroprotection also indicates that nonspecific responses such as stress associated with repeated seizures, various endocrine changes and other nonspecific physiological responses to seizures cannot account for the neuroprotection that appears to be selective for seizures involving limbic forebrain activation. Instead, adaptive changes restricted to the network through which the seizures propagate appear to be required for the neuroprotective state.

The fact that exogenous infusion of bFGF directly into hippocampus can protect against excitotoxic neuronal injury [75, 76] indicates that an increase in bFGF protein, as observed after several days of electroshock seizures [41] may account for a component of seizure-induced neuroprotection. It is also likely that multiple neuroprotective adaptive responses are engaged by repeated seizures and that the relative importance of any given factor may vary with cell type and brain region.

Because very brief seizures are remarkably protective even in the complete absence any evidence of injury or cellular stress, it may be that seizures serve to trigger activity-dependent mechanisms of neuroplasticity that recapitulate the injury-resistant and resilient conditions characteristic of development. In the face of insults to the nervous system, transient recurrent seizure activity could serve to attenuate neurodegeneration and promote regrowth and remodeling in the network affected by an insult. An especially robust demonstration of this type of protective action comes from a study in which daily electroshock seizures were administered to adrenalectomized rats [78]. Removal of the adrenal glands in adult rats leads to a highly selective apoptotic degeneration of the dentate granule cells in the hippocampus [74, 119, 120], reflecting the fact that these neurons are directly or indirectly dependent upon adrenal corticosteroids [39]. These seizures completely prevented the dentate granule cell degeneration, while sham treatments or daily exposure to restraint stress did not alter the profile of degeneration seen in the adrenalectomized animals.

Moreover, daily exposure to brief seizures has been shown to accelerate recovery of function following cortical damage, probably by enhancing post-injury plasticity in local and distant networks connected to the site of injury [43, 50]. Thus, clinical, and possibly subclinical, seizures that occur transiently during the post-traumatic period may serve an adaptive function: to reduce injury, promote repair and trigger compensatory plasticity. If this is the case, then the frequent procedure of placing patients on prophylactic anticonvulsant therapy immediately following either head injury or neurosurgical intervention may potentially retard or diminish functional recovery [49, 111]. In this context, seizures may be analogous to fever—a symptom that may have adaptive and protective value in specific pathological settings; and like fever, seizures can become maladaptive and injurious in their own right in the rare cases where they go beyond a self-limiting state and evolve into a chronic epileptic condition.

During nervous system development, a dynamic continuum between neuronal plasticity and neuronal death exists and coordinated synaptic stimulation can ‘protect’ or ‘select’ the population of neurons or synapses that will be maintained. In an analogous fashion, seizure evoked stimulation may allow the sparing of otherwise vulnerable populations of neurons in the injured adult brain, a phenomenon that we have referred to as ‘excitotrophic’ [35]. This could account for the therapeutic benefits of controlled administration of electroshock seizures in various neurodegenerative disorders including Parkinson’s and Huntington’s Disease [1, 8, 9, 30, 34, 60, 62, 91, 99, 108, 122], but it remains to be determined if the seizures slow the disease progression. Currently electroshock seizures are used in the treatment of bipolar affective disorders and their therapeutic impact may derive from neuroprotective actions [77]. Further characterization of the mechanisms contributing to the neuroprotective impact of brief seizure episodes may generate novel strategies of neuroprotection and recovery of function following excitotoxic insults and other forms of injury to the central nervous system.

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#### **8.4 Controlled Patterns of Hyper-synchronous Discharge in Certain Subcortical Networks**

As Stevens had observed, “Rapid neuronal discharge (bursting), although typical of epileptic discharge, is part of the normal brain repertoire and does not necessarily signal pathology” [123]. These events can be distinguished from pathological seizure activity in that they occur in highly confined areas and/or during highly restricted time periods (such as during certain phases of sleep). The networks of the limbic system, brainstem, and diencephalon are organized in such a way as to give rise to highly synchronized bursting in discrete nuclei in association with certain physiological states or functions such as parturition, growth hormone release, milk ejection, ovulation, and

orgasm (see discussion in [123]). In association with the estrus cycle, neuronal spiking and coordinated burst discharges in nuclei of the basal hypothalamus and forebrain limbic preoptic area appear to coordinate the cyclical and pulsatile release of reproductive hormones [61]. Stevens [123] described this type of highly regulated hypersynchronous discharge as “microseizures” that serve to augment signal transmission for critical species-specific survival functions, but do not propagate beyond highly restricted circuitry due to surrounding inhibitory control. In conditions in which the inhibitory control mechanisms are compromised, such microseizures could potentially propagate beyond their physiologically appropriate boundaries. The fact that certain phases of sleep [54, 96, 116] or hormonal cycles [7, 26, 31, 112] are associated with increased vulnerability to seizures may be a reflection of this natural fluctuation in physiological microseizure discharge.

The limbic system network is also organized to generate synchronized, reverberatory discharge characteristic of Hebbian cell assemblies [47] that instantiate memories via temporal lobe circuitry. This core feature of associative learning is reminiscent of the activity-dependent plasticity that drives the shaping of neuronal connectivity during development. But in the case of learning in the mature brain, there is a selective strengthening of specific synaptic connections within a network in a highly defined spatiotemporal pattern. The limbic network comprised of the hippocampus, amygdala, mediodorsal thalamus, and piriform and rhinal cortices is especially suited to the amplification of discharge patterns via reverberatory loops between the nuclei. The ability to amplify repetitive discharge originating at one site, such as occurs during the process of kindling from the amygdala or other sites within the limbic network, is a reflection of the propensity for activity-dependent plasticity in this system. In fact, kindling has been used as a model of learning and memory [36], especially because once an animal is fully kindled, the remodeling of the network that supports the kindled state is relatively permanent [37]. At the same time, it is curious that

the network would be poised to amplify the stimulation to the point that propagated seizures emerge, considering the fact that these seizures have no clear adaptive function. Perhaps by providing near-physiological stimulation in a repetitive manner, the kindling process lures the network into an amplification process until it becomes hijacked by the long-lasting modifications associated with the repeated seizures.

The transfer effect in kindling, in which pre-kindling from one area reduces the number of stimulations necessary to kindle from another area, may likewise reflect plasticity within the limbic seizure network (or the recruitment of new components to the network, as discussed above in the context of secondary seizure foci). However, transfer of kindling within the limbic system appears to require other networks (e.g., brainstem). For example, transfer of amygdala kindling is impaired by transection of or damage to midbrain and brainstem [20, 42, 51, 132]. This suggests that repeated limbic stimulation can alter the functions of other networks, either by actively recruiting them into a transfer process, or perhaps by disrupting endogenous seizure-suppressive functions of these extra-limbic circuits.

The fact that the process of kindling can be retarded or suppressed by the occurrence of generalized convulsive seizures [69, 93, 100] emphasizes the importance of homeostatic mechanisms for the process. The same is true for kindling using chemoconvulsants or corneal electroshock [28, 29, 105, 109, 138]: repeated minimal (threshold) limbic seizures become amplified over time, while repeated maximal seizures induce a seizure-resistance over time.

It is, however, essential to recognize that the vulnerability to kindling is highly species-specific, with the rate of kindling taking days to weeks in rodents [37, 81], weeks to months in cats [37, 42, 135], and months to years (with a relatively low success rate) in primates [37, 131, 133, 134]. These species differences may reflect the extent to which the limbic network is under inhibitory control from an increasingly elaborated frontal cortex. Thus, the kindling phenomenon, which

has been most thoroughly characterized in the rodent, may not readily generalize to humans or to human clinical conditions.

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## 8.5 Conclusions

Abundant evidence supports a neuroplasticity-inducing action of seizures and seizure-like events in the CNS. However, the extent to which the neuroplasticity serves an adaptive function vs. a maladaptive function depends on the context in which the seizure activity occurs. During fetal CNS maturation, neuroplasticity induced by naturally-occurring ictal activity and seizure-like phenomena promotes the formation of neural connections. Similarly, in the aftermath of injury in the mature CNS, limited seizure activity may promote neural repair and compensation and serve a neuroprotective role. However, the circumstances in which seizure activity serves a “normal” function typically involve seizure activity that is highly limited temporally and/or spatially (e.g., in specific circuitry, during specific developmental stages, or within a short period post injury). In the small percentage of cases in which the seizure activity does not remain highly controlled and limited, it becomes pathological, repeatedly interrupting normal function with maladaptive, and even potentially injurious consequences. The epilepsies represent this type of pathological seizure occurrence, and it is likely that some of the associated neuroplasticity impairs normal CNS function. Whether the neuroplasticity is also an essential component of the process of epileptogenesis remains to be determined, but since chronic epilepsy occurs only in a small percentage of individuals, we first need to understand the unique features that render those individuals susceptible, and whether the unique features change the nature of the seizure-induced plasticity in those individuals.

We can therefore conclude that epileptogenic mechanisms may indeed be a corruption of normal, adaptive neuroplasticity. If the normal neuroplasticity associated with limited, controlled seizure activity is largely helpful, turning

pathophysiological in rare circumstances, this raises several challenging questions for future epilepsy research to address:

1. Do adaptive and maladaptive neuroplasticity differ, and if so, how?
2. If there are distinctions between adaptive and maladaptive neuroplasticity, can we selectively prevent the maladaptive with compromising the adaptive?
3. Which control mechanisms that normally prevent seizures from becoming repetitive and self-sustaining become compromised in individuals susceptible to epilepsy? Is it possible that it is the failure of these control mechanisms, rather than neuroplasticity, that is essential for epileptogenesis?

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# When and How Do Seizures Kill Neurons, and Is Cell Death Relevant to Epileptogenesis?

9

Ray Dingledine, Nicholas H. Varvel,  
and F. Edward Dudek

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## Abstract

The effect of seizures on neuronal death and the role of seizure-induced neuronal death in acquired epileptogenesis have been debated for decades. Isolated brief seizures probably do not kill neurons; however, severe and repetitive seizures (i.e., status epilepticus) certainly do. Because status epilepticus both kills neurons and also leads to chronic epilepsy, neuronal death has been proposed to be an integral part of acquired epileptogenesis. Several studies, particularly in the immature brain, have suggested that neuronal death is not necessary for acquired epileptogenesis; however, the lack of neuronal death is difficult if not impossible to prove, and more recent studies have challenged this concept. Novel mechanisms of cell death, beyond the traditional concepts of necrosis and apoptosis, include autophagy, phagoptosis, necroptosis, and pyroptosis. The traditional proposal for why neuronal death may be necessary for epileptogenesis is based on the *recapitulation of development hypothesis*, where a loss of synaptic input from the dying neurons is considered a critical signal to induce axonal sprouting and synaptic-circuit reorganization. We propose a second hypothesis – the *neuronal death pathway hypothesis*, which states that the biochemical pathways causing programmed neurodegeneration, rather than neuronal death *per se*, are responsible for or contribute to epileptogenesis. The reprogramming of neuronal death pathways – if true – is proposed to derive from necroptosis or pyroptosis. The proposed new hypothesis may inform on why neuronal death seems closely linked to epileptogenesis, but may not always be.

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## Keywords

Neurodegeneration • Epilepsy • Necrosis • Apoptosis • Autophagy  
• Phagoptosis • Necroptosis • Pyroptosis

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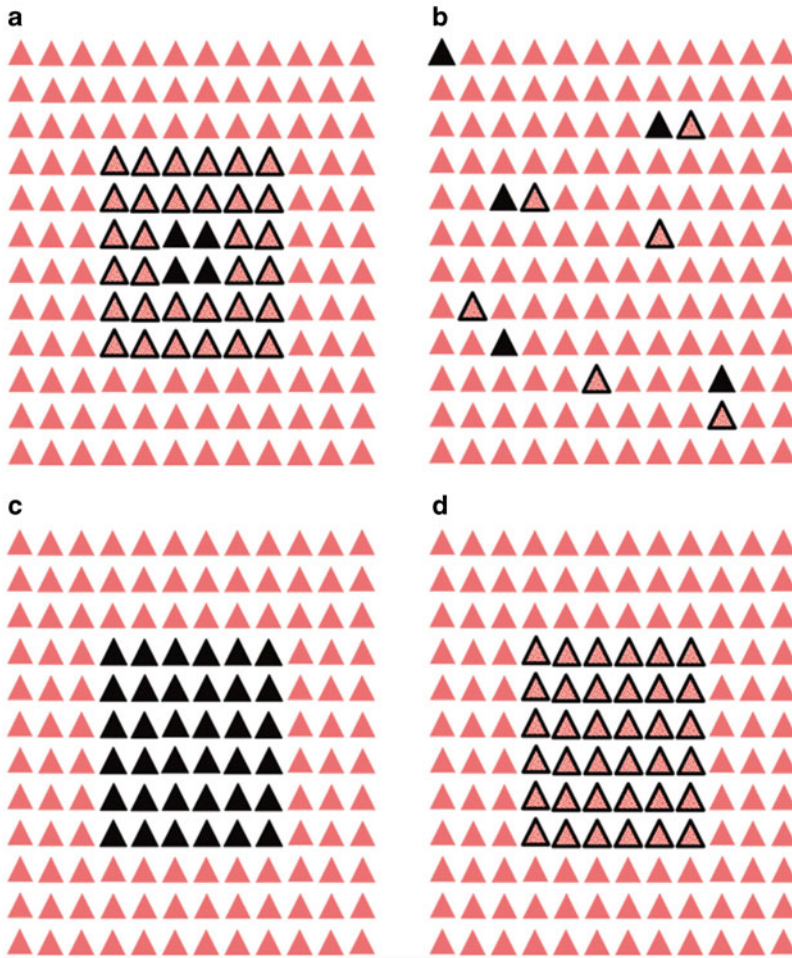
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## 9.1 Seizures and Neuronal Death: When, Where, and What?

Debates and controversies concerning the interplay among seizures, neuronal death and epilepsy continue to occur. Over several decades, many epilepsy researchers have focused on various aspects of the issue of whether seizures cause neuronal death, and conversely, whether neuronal death is necessary and/or sufficient to cause epilepsy. For example, a classic – yet still ongoing – debate is the degree to which GABAergic interneurons are lost in tissue from patients and animal models of temporal lobe epilepsy, and the consequence of such loss. In spite of the longevity and intensity of the previous debates, the relationship between seizures, neuronal death and epilepsy remains one of the most disputed in translational neuroscience, particularly as it relates to possible mechanisms of acquired epileptogenesis and the clinical interactions and consequences of seizures and neuronal death. We will discuss, as the title implies, two important and longstanding hypotheses of contemporary epilepsy research – important because the degree to which seizures cause brain damage and the hypothetical role of neuronal death in the development of epilepsy are inter-related and could underlie the often quoted statement “Seizures beget seizures” [33]. These two issues are not “black and white”; rather, they probably form an interactive continuum and are quite complicated; and furthermore, technical limitations and interpretational difficulties plague any analysis of them. The key questions include *when* do seizures kill neurons, *where* in the brain are neurons most susceptible to seizure activity, and *what* is the identity of the neurons that are preferentially killed? Answers to a fourth issue – “*How* do seizures kill neurons?” – may hold a key to understanding at least one component of epileptogenesis, as described below. We will begin with a brief summary of some of the key questions and controversial topics; then, we will review more recent views of the many possible mechanisms whereby seizures may kill neurons; and finally, we will conclude with a brief discussion of some of the ongoing issues and controversies in this area.

### 9.1.1 When

A large and long-standing body of experimental and clinical data indicates that some types of seizures lead to neuronal death, while other types do not. In either experimental animals or humans, whenever seizures are long enough in duration and occur repetitively for prolonged periods, some neurons – particularly in adults – are killed. In terms of the temporal features of the seizures that are thought to cause neuronal death, relatively brief seizures – such as typical *absence* seizures in children (usually lasting 5–10 s) – do not appear to cause overt brain damage. However, the more prolonged seizures characteristic of temporal lobe epilepsy, such as the traditional complex partial seizures (i.e., dyscognitive focal seizures) that may progress to tonic-clonic convulsive seizures, are much more likely to lead to neuronal loss [84]. Finally, the prolonged and repetitive seizures that define status epilepticus typically cause brain damage, often with extensive neuronal death [10, 15, 29, 40, 57, 62, 67, 85]. Interestingly, however, status epilepticus in the immature brain causes far less neuronal death [16, 38, 59, 68, 74, 75, 80, 81], and appears less likely to cause epileptogenesis [51, 74]. The long-standing observation that experimental status epilepticus in laboratory animals, mostly rodents, leads to a chronic epileptic state raises the following question: Is the occurrence of neuronal death during status epilepticus a critical part of the epileptogenesis? In terms of epilepsy, one could view seizure clusters, where some of the interseizure intervals are much shorter than the typical interseizure intervals [36, 37], as essentially a reduced form of status epilepticus. The difference between status epilepticus and a seizure cluster in a patient with epilepsy is not always so clear. Thus, a fundamental question in clinical epilepsy is: Do the spontaneous recurrent seizures kill neurons – particularly when the seizures occur in clusters? If so, under what conditions does this contribute to a worsening of epilepsy? Are seizure clusters a particular concern in terms of neuronal death and brain injury? These are some of the unanswered questions that are both clinically important and can theoretically be addressed with animal models.



**Fig. 9.1** Schematic diagrams showing hypothetical relationships of neuronal populations after a brain insult that activates cellular mechanisms of neuronal death. In the four panels of the figure, two or three populations of neurons are depicted in a schematic manner. Dead neurons (*filled black triangles*) are shown within a network of live and completely-normal neurons (*filled red triangles*). Among these two populations of cells is another group of neurons, which form the core of this hypothesis; these neurons have undergone only the initial steps of a neuronal-death and/or are under the molecular influence of the neuronal death process (*black triangular outline with red stiples* inside). (a) Focal neuronal loss. A small cluster of dead neurons is shown to be clumped together within a network of normal neurons, as would be expected to occur during an infarct. Between these two completely different neuronal populations is the group of neurons

that are hypothetically epileptogenic, because they have undergone the first part of a neuronal-death process and/or are under the molecular influence of the neuronal death process. (b) Diffuse neuronal loss. Using the same code to define the members of the neuronal population, this diagram illustrates scattered neuronal loss, as would be expected to occur after status epilepticus (vs an infarct in (a)). (c) Occurrence of neuronal death without generation of neurons altered or influenced by death-process mechanisms, which theoretically represents the occurrence of frank brain damage without subsequent epilepsy. (d) Absence of neuronal death after a brain insult, but with the presence of death-pathway neurons. In this case, the death-pathway neurons are hypothesized to become epileptogenic, and they generate spontaneous recurrent seizures without the prior occurrence of overt neuronal death

### 9.1.2 Where?

If we focus on the seizures that characterize temporal lobe epilepsy and other forms of severe acquired epilepsy (e.g., after hypoxic-ischemic

encephalopathy), many specific areas appear to be particularly prone to seizure-induced neuronal death. Depending on the etiology, neuronal death can be relatively circumscribed, as with an infarct (Fig. 9.1a), or it can be diffuse (Fig. 9.1b).

Seizures, particularly repetitive seizures, cause substantial brain damage in highly susceptible areas, such as parts of the hippocampus, entorhinal cortex, amygdala, thalamus and other limbic structures; however, neuronal death after seizures can be more widespread and is generally quite variable (e.g., [24, 77]).

### 9.1.3 What?

A focus in epilepsy research has been – and remains – the unequivocal identification of the type(s) of neurons that are killed: glutamatergic principal neurons, such as cortical pyramidal cells, and subpopulations of GABAergic interneurons, which comprise 5–10 % of the neurons in epilepsy-relevant brain regions and are highly heterogeneous in their anatomy and electrophysiology [4]. Regardless of the type of brain insult, the potential loss of interneurons is obviously a special case, because the loss of interneurons, if uncompensated by inhibitory axonal sprouting, can translate to a reduction in GABAergic tone.

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## 9.2 What Are Some of the Important Technical and Experimental-Design Issues?

The challenges and controversies concerning how to evaluate whether neuronal death has occurred and how to quantify it are substantial. Even when one only considers a fraction of the methodological and protocol-related issues, the additional factors involving “what, where, and when” of neuronal death (“how” is discussed below) add further complexity to the potential analyses and interpretations. Additional disagreement surrounds the question “What is a seizure?” and the problem of what comprises an adequate animal model of acquired epilepsy.

An important issue in regard to considerations of neuronal death in epilepsy – as with most other research – involves the complimentary concerns of false positives (specificity) and

false negatives (sensitivity). For example, two of the main approaches to analyzing neuronal death involve staining (1) those neurons that *remain* after seizures and (2) the neurons that are *destined to die*. Staining the remaining neurons involves a variety of traditional techniques such as cresyl violet staining of Nissl substance, and/or more specific methods including but certainly not limited to immunocytochemical staining of specific cell types, such as GABAergic interneurons. This most basic level of methodology has numerous caveats – some of which are obvious, and others not. For example, what does it mean when one finds no significant (i.e., statistical) difference between an experimental condition or animal model and the control group? On first principles, one has to ask: Does this mean that no neuronal loss (death) has occurred? Or, could it mean the amount of neuronal death was so small that it could not be detected? Issues such as how the tissue was sectioned (section thickness, where in the brain, but also orientation) are relevant, not to mention that extensive cell loss in epilepsy is associated with tissue shrinkage. Thus, cell number can be quite different than cell density. In regard to use of histological stains that mark “dying” neurons, such as FluoroJade B (FJB), one must also consider their advantages and disadvantages. For example, one has to question our confidence that they will actually die – can FJB-labeled neurons remain viable for a prolonged period before death? If we assume that all of the FJB neurons are going to die, or even most of them, then this approach has the important advantage that it can reveal situations in which only a small fraction of the neurons will die, which is simply not feasible with stains that mark the “remaining” neurons. Another issue, however, is that the FJB technique will only stain neurons that are dying at that particular time; so therefore, euthanasia, fixation, and staining must be performed at the appropriate time; neurons could have died at other times, and their death would not be detected with FJB [94]. Thus, although it is quite difficult to quantify neuronal loss, it is even more difficult – if not impossible – to show that neuronal loss has not occurred.



### 9.3 Mechanisms of Seizure-Induced Neuron Injury

In order to explore how seizures could kill neurons, it is first necessary to review cell-death pathways; our understanding of them has expanded well beyond the traditional mechanisms of *apoptosis* and *necrosis*. This seemingly simple endeavor is complicated by the observation that some of the newly identified cell-death pathways share criteria used for identification. For clarity, we classify cell-death processes as non-inflammatory (apoptosis, autophagy, phagoptosis) and inflammatory (necrosis, necroptosis, pyroptosis) (Table 9.1).

#### 9.3.1 Apoptosis

A controlled, programmed process of packaging internal components of the cell for clearing by phagocytes characterizes the apoptotic process. As such, intracellular molecules with the potential to activate immune responses are disposed of rapidly, without initiating an immune response [2]. Apoptosis is also characterized by chromatin and cytoplasmic condensation, plasma membrane blebbing, formation of apoptotic bodies as well as fragmentation of cellular compartments and DNA. Apoptosis occurs naturally during development and serves as a means to facilitate cellular turnover in healthy tissue, and also in response to hormone withdrawal [47]. This programmed series of events is reliant upon the effector functions of activated caspases -3, -6, and -7, which enzymatically cleave intracellular organelles, proteins and DNA. The degraded cellular corpse is then packaged in preparation for phagocytosis by macrophages or

microglia [26]. Processing of intracellular compartments and subsequent removal of cellular debris during apoptosis does not result in a secondary inflammatory response in surrounding tissue as inflammatory mediators are largely sequestered and degraded [2]. Changes in mitochondrial membrane permeability [50] and release of mitochondrial proteins are also observed [87]. Another characteristic of apoptotic cells is the exposure of phosphatidylserine on the extracellular leaflet of their plasma membrane. While phosphatidylserine is normally found exclusively on the cytoplasmic side of the plasma membrane, apoptotic cells present phosphatidylserine on the extracellular surface to serve as an “eat-me” signal for neighboring phagocytes [32, 70]. Cellular shrinkage, likely due to caspase-mediated proteolysis of cytoskeletal proteins, also typifies apoptotic cells [49].

#### 9.3.2 Autophagy

Although autophagy usually serves a protective role, in extreme stress conditions it can contribute to cell death. In similar fashion as apoptosis, autophagic pathways also progress in a series of cellular steps that involve programmed degradation of cellular components. However, intracellular autophagic, largely non-caspase, enzymes are responsible for degradation of organelles or other cytoplasmic proteins within double-membrane vesicles known as autophagosomes [54]. The autophagosome then fuses with intracellular lysosomes to facilitate degradation of the contents within the autophagosome by acid hydrolases. In contrast to apoptosis, caspase activation is not required and chromatin condensation is minor [11].

**Table 9.1** Six mechanisms of cell death

Death process	Programmed?	Inflammatory lysis?	Effector	Shape $\Delta$	TUNEL?
Necrosis	No	Yes	Non-caspase	Swell	No
Necroptosis	Yes	Yes	TNF- $\alpha$ RIPK1	Swell	No
Pyroptosis	Yes	Yes	Caspase-1	Swell	Yes
Autophagy	Yes	No	Lysosomes	?	No
Phagoptosis	Yes	No	Microglia	No	No
Apoptosis	Yes	No	Caspase-3/6/7	Shrink	Yes

In addition to contributing to the death of a cell, autophagic mechanisms also contribute to cellular function and homeostatic maintenance. For example, in the immune system, antigen-presenting cells utilize autophagy to digest intact proteins, creating smaller antigens for subsequent presentation to T lymphocytes [21, 54]. Moreover, mice deficient in proteins involved in autophagy develop spontaneous neurodegeneration [35, 48]. Taken together, these findings indicate that, in addition to cell death, autophagy mediates an important role in the organism's response to pathogens as well as maintenance of cellular homeostasis.

### 9.3.3 Phagoptosis

Many of the identified physiological cell death pathways involve phagocytosis of either whole cells doomed to die or of fractured cellular components. As such, the process of phagocytosis has been viewed as a secondary event, occurring after the death of the cell [70]. However, the process of phagocytosis can also kill living cells. Recent studies have identified a pathway, termed "phagoptosis", wherein phagocytes such as activated microglia actively contribute to the death of viable neurons and other cells [8]. Similar to apoptosis, the otherwise viable cell presents "eat-me" signals, such as phosphatidylserine, on the outer leaflet of its cellular membrane. The "eat-me" signals are then recognized by nearby phagocytes, and cellular uptake ensues followed by digestion of the viable cells. Importantly, cell death can be prevented during phagoptosis by inhibiting phagocytosis [28, 60]. This is because "eat-me" signal exposure is transient and reverses when phagocytosis is prevented. Therefore, neuronal insults not severe enough to initiate apoptotic pathways might be a trigger for phagoptosis due to the temporary exposure of eat-me signals on stressed but viable neurons [8, 28].

### 9.3.4 Necrosis

In contrast to these non-inflammatory modes of neuron death, during necrosis cells lyse, effectively

spilling their internal contents into the interstitial fluid and releasing molecules that can initiate inflammatory cascades. This uncontrolled release of intracellular molecules can potentially damage surrounding tissue and cells [76, 79]. Necrotic cell death is typically initiated by extreme physiological stress or trauma that kills cells quickly. Biochemically, caspase is not involved. Morphologically, condensation or digestion of internal cellular compartments is not observed. Instead, organelles and the entire cell undergo extensive swelling. The cell eventually bursts, spilling its internal contents into the surrounding environment, triggering robust inflammation in the neighboring tissue [45].

### 9.3.5 Necroptosis

While necrosis leads to an uncontrolled cellular death, a variant of necrosis, which has some controllable features, has recently been described. This programmed pathway, termed necroptosis, exhibits characteristics of both programmed cell death and necrosis. The main characteristic distinguishing necrosis from necroptosis is that the latter is initiated by TNF- $\alpha$  and other death receptor activators, which promote the assembly of receptor-interacting protein kinase 1 (RIP1) with RIP3 [86]. Thus, kinase activity controls necroptosis [45]. Interestingly, RIP1 and RIP3 assemble into a functional kinase-containing cell-death complex only in the absence of functional caspase 8 [25, 46]. While the physiological impact of necroptosis is currently under investigation, it is conceivable this pathway may be relevant in the event caspase activity is impeded and thus canonical apoptosis is not possible.

### 9.3.6 Pyroptosis

Perhaps the most extreme example of inflammation-related cell death is pyroptosis (i.e., caspase 1-dependent programmed cell death). While this form of cell death was first described

after infectious stimuli, such as *Salmonella* and *Shigella* infection [6, 18], caspase-1-dependent cell death also occurs in myocytes after myocardial infarction [27] and in the central nervous system [53, 103]. The primary distinguishing feature of pyroptosis is the formation of the inflammasome, an intracellular multimolecular complex that is required for the activation of inflammatory caspases, particularly caspase 1. The activated inflammasome culminates with production of enzymatically active caspase 1, which in turn mediates the maturation and secretion of active IL-1 $\beta$  and IL-18 [2]. Secreted pro-inflammatory cytokines can subsequently influence nearby cells with potentially adverse consequences, such as blood-brain barrier breakdown and possible leukocyte entry into the brain. Although TUNEL-positive breaks in cellular DNA typify both apoptosis and pyroptosis, the latter is entirely reliant upon caspase-1 [7, 17]. This is important for classification purposes because caspase-1 is not involved in apoptosis. Mitochondrial release of cytochrome c, a hallmark of apoptosis, also does not occur during pyroptosis. In contrast to the coordinated packaging of intracellular components observed in apoptosis, cellular lysis and release of inflammatory effector molecules occur during pyroptosis [26].

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## 9.4 How Might Seizure-Induced Neuronal Injury Promote Epileptogenesis?

### 9.4.1 Overview of Two Competing Hypotheses

We envision two conceptually distinct answers to this question. *First*, maladaptive new circuits among neurons could form to replace synapses lost during neuronal death. This mechanism, potentially involving axonal sprouting within excitatory pathways and amplified by loss of inhibitory interneurons, has been described in numerous previous studies and can be termed the “*recapitulation of development*” hypothesis. If replacement of lost synapses is the critical factor underlying this mechanism, then neuronal death

would seem to be an essential component of the process. *Second*, rather than neuron death *per se* being responsible, molecular signals from upstream pathways that mediate some of the more newly recognized forms of cell death might underlie or contribute to epileptogenesis. We call this the “*neuronal death pathway*” hypothesis. We will focus on potential roles for IL-1 $\beta$  and TNF- $\alpha$ . We will also consider whether the inflammasome pathways (caspase-1 activation leading to synthesis of IL-1 $\beta$  and IL-18), normally considered a feature of myeloid cells and innate immunity, might be involved in epilepsy-related neurodegeneration.

In some cases focal inflammation produced by lytic cell death, perhaps involving only a small number of neurons undetectable by normal Nissl stains (e.g., Fig. 9.1b), could promote increased neuronal excitability and perhaps synchronous activity. However, in the absence of any neuronal death (Fig. 9.1d), how might inflammatory cascades be initiated? Understanding how microglia, the innate immune cells of the CNS, respond to injurious or danger signals may provide insights into this undoubtedly complex process. Microglia in the intact, healthy brain continuously palpate the surrounding tissue for subtle disturbances [61], and can rapidly respond to tissue injury or danger signals by altering morphology, proliferating and expressing a wide variety of inflammatory cytokines and chemokines [19, 69]. Microglial activation can be initiated by injured neurons through the release of molecules collectively known as alarmins [4].

One well-characterized alarmin, prostaglandin E<sub>2</sub>, is released by highly active neurons in a COX-2-dependent process. Cyclooxygenase 2 (COX-2) is rapidly upregulated in hippocampal pyramidal cells and dentate granule cells after seizures [55, 73, 98], but the impact of neuronal COX-2 has remained elusive because astrocytes, endothelial cells and probably other cell types in the CNS also express COX-2. To determine the role of neuronal COX-2 after status epilepticus, a neuron specific conditional knockout mouse was utilized wherein principal neurons of the hippocampus, dentate granule cells, amygdala, thalamus and layer-specific neurons in the piriform

and neocortex (layer 5) are devoid of COX-2, while the remaining cell types of the CNS still express functional protein [43, 72]. Interestingly, conditional ablation of COX-2 from neurons resulted in less severe damage to hippocampal neurons after status epilepticus produced by pilocarpine. The intensity of status epilepticus was not diminished in the COX-2 conditional knock-outs, as judged by the temporal evolution of behavioral seizures and by cortical EEG [78], making it unlikely that neuroprotection was caused by a less severe seizure episode. Neuroprotection was accompanied by reduction in multiple markers of neuroinflammation as well as preserved integrity of the blood-brain barrier, suggesting that neuronal COX-2 mediates a broad deleterious role after status epilepticus. These findings provide strong evidence that the neuron itself can contribute to the neuroinflammatory milieu [78]. The beneficial effects of the conditional ablation of COX-2 from principal forebrain neurons were completely recapitulated by systemic administration of a novel antagonist of EP2, a receptor for PGE2 [44].

Injured neurons might indirectly contribute to inflammation after status epilepticus through cell-to-cell signaling with microglia. Multiple lines of evidence indicate that the local microenvironment plays an important role in regulating the microglial phenotype wherein microglia activation is constitutively inhibited by repressive forces [34, 65, 69]. For example, surface proteins on microglia, such as CD200R and CX3CR1 (the fractalkine receptor), normally interact with the neuronal surface protein ligands, CD200 and CX3CL1 (fractalkine), respectively [14, 42]. If interactions between CD200R and CD200 [42, 102] or CX3CR1 and CX3CL1 [3, 13] are disrupted by signals released during neuronal damage or distress, then microglia are unleashed from this constitutive state of inhibition and a more florid microglial response ensues. Enhanced microglial activation is likely attributed to the presence of ITIM motifs (immunoreceptor tyrosine-based inhibitory motif) on both CD200R and CX3CR1 as these motifs function as activators for SHP-1 and SHP-2 phosphatases that can repress further inflammatory signaling [5]. Indeed, CX3CR1-deficient mice exhibit microglia-mediated neurotoxicity,

through enhanced IL-1 $\beta$  secretion, after immune challenge [14]. Interestingly, altered expression of CX3CL1 has been reported in both epileptic patients and animals models after status epilepticus [97].

In addition to the above-mentioned studies, viable neurons might also induce inflammatory cascades. Studies in *Drosophila melanogaster* originated this concept, wherein damaged cells, prevented from dying, release mitotic signals that prompt neighboring cells to divide. Cells in the wing of flies were triggered to die by X-rays, but they were blocked from completing the death process by expression of anti-apoptotic proteins. The authors describe the resulting cells as “undead”. The neighboring cells divide in an apparent attempt to fill the void in the tissue expected to be left by the dying cells [64]. Do similar situations occur in human disease? Interestingly, neuronal populations expected to degenerate in the brains of Alzheimer’s Disease (AD) patients re-express proteins typically encountered in a mitotic cell cycle [12, 58, 92, 93]. Importantly, DNA replication accompanies cell cycle entry [99]. Transgenic mouse models of AD also recapitulate neuronal cell cycle entry [88, 101], suggesting that the same “stressors” that provoke neuronal cell cycle entry in the human AD brain are phenocopied in the mice. However, cycling neurons exhibit little atrophy [100] and robust neuronal loss is absent in AD mice [41, 56], indicating that re-expression of mitotic proteins and DNA synthesis in a post-mitotic neuron is not sufficient to induce death, at least in the lifetime of the mouse. It has been proposed that cycling neurons also might send out mitotic signals, pressuring otherwise healthy neurons to enter the “undead” state [39].

#### 9.4.2 Inflammatory Pathways and Epileptogenesis

How might inflammatory signaling upstream of neurodegeneration increase excitability and subsequent synchronicity? Immune responses in the brain are initiated, maintained and terminated by soluble effector proteins known as cytokines. Although a strong correlation between seizures

and elevated inflammatory cytokines or their mRNA transcripts has been reported [90], emerging experimental evidence indicates that inflammatory cytokines can in turn alter neuronal excitability and synchronicity by modulating receptor function and expression [31, 89]. For example, the pro-inflammatory cytokine TNF- $\alpha$  has also been shown to promote the recruitment of AMPA receptors to postsynaptic membranes. Interestingly, the recruited receptors preferentially lack the GluR2 subunit [52, 63, 82] and consequently the calcium conductance underlying EPSPs is increased. Additionally, TNF- $\alpha$  causes endocytosis of GABA<sub>A</sub> receptors from the cellular surface, decreasing inhibitory synaptic strength [82]. Taken together these findings demonstrate that TNF $\alpha$  can have a profound impact on circuit homeostasis in a manner that can provoke the pathogenesis of seizures.

In addition to TNF- $\alpha$ , multiple lines of evidence directly implicate IL-1 $\beta$  in lowering the seizure threshold, and perhaps in epileptogenesis. First, hippocampal application of IL-1 $\beta$  can increase seizure intensity threefold. This proconvulsant effect is attributed to IL-1 $\beta$ -mediated engagement of Src-family kinases in hippocampal neurons. The activated kinases subsequently phosphorylate the NR2B subunit of the NMDA receptor, leading to seizure exacerbation [1]. Second, IL-1 $\beta$  can inhibit calcium currents through protein kinase C, at least at low concentrations [66]. Finally, IL-1 $\beta$  can also inhibit GABA<sub>A</sub> receptor current, which could underlie neuronal hyperexcitability [95]. These studies, coupled with the findings that pharmacological treatments targeting IL-1 $\beta$  or its activation result in robust anticonvulsant effects [20, 71, 90, 91], indicate that inflammation might play an important role in epileptogenesis and is a viable therapeutic target class.

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## 9.5 Implications of the New Concepts on Neuronal Death for Epileptogenesis

The long-standing *recapitulation-of-development* hypothesis essentially states that neuronal death in acquired epilepsy is linked to a re-activation

of developmental processes, which replace the synapses lost through neuronal death [30]. Initially, most experimental and clinical epileptologists viewed this hypothesis as “mossy fiber sprouting”, which causes the formation of new recurrent excitatory circuits among dentate granule cells. This hypothesis is discussed by Buckmaster [9] and a more general view would be that neuronal death in many areas of the brain, particularly in seizure-sensitive regions, causes multiple networks to form new local excitatory circuits [22, 23, 83]. The data reviewed above suggest a new hypothesis, the *neuronal-death-pathway* hypothesis, whereby the biochemical pathways causing programmed neurodegeneration, rather than neuronal death *per se*, are responsible for or contribute to epileptogenesis. This hypothesis is consistent with the view that frank brain damage (i.e., cases where obvious neuronal death has occurred) leads to epilepsy, and further, that the likelihood of developing intractable epilepsy is linked somehow to the severity of the brain injury. In addition, however, this hypothesis may begin to explain why brain injuries that clearly induce neuronal death do not always appear to lead to epilepsy, since the critical hypothetical mechanism for acquired epileptogenesis would be the linkage between the to-be-defined mechanisms *within the pathways responsible for neuronal death*, as opposed to neuronal death itself (Fig. 9.1c). The identification of these hypothetical processes is an area ripe for future investigation. Finally, this hypothesis could also explain how epilepsy may occur when neuronal death is absent or appears minimal (Fig. 9.1d). It is conceivable that these molecular mechanisms may be aborted or reversed before neuronal death actually occurs, for example, so that specific signaling molecules direct some of the surrounding neurons toward an epileptogenic phenotype, even though the processes of neuronal death may not reach completion. The key point here is the proposal or hypothesis that molecular/genetic signals from neurons that are on a “death pathway” could initiate epileptogenesis independent of the final outcome (i.e., neuronal death).

## 9.6 Concepts and Conclusions

Although much has been learned about when seizures do kill neurons and the conditions when they appear to cause less damage, it is extraordinarily difficult to *rule out* that neuronal death has occurred after seizures. One problem is both a conceptual and technical one, namely, showing that something has not occurred is particularly challenging, if not impossible. We simply do not know if a threshold exists whereby a few, brief seizures – possibly in the seizure-resistant immature brain – cause absolutely no neuronal death. In terms of the question, “When?”, there is no way to show that neuronal death has not occurred during and/or after seizures, except to count the remaining neurons in control and experimental groups; however, the potential error – even in well-powered studies, can be 10 % or more ([10] [see Table 1 and Fig. 2A-B]; [96] [see Fig. 6]) – and yet a loss of just a few percent of the neurons within a brain structure could have a substantive epileptogenic effect. If one considers the problem of “Where?”, it becomes obvious that the answer is “Almost anywhere!”. For the animal models of repetitive seizures and status epilepticus – whether induced by hypoxia, pilocarpine, or some other precipitating insult – numerous seizure-sensitive areas of the brain show neuronal loss, and the structure could be different for individuals within a similarly-treated cohort of animals, further supporting the idea that it is extremely difficult to exclude a role of neuronal death. In terms of, “What types of neurons may be lost?”, excluding loss of part of the critical interneuron pool generally requires specific staining techniques, such as immunohistochemistry with stereology (e.g., [10]). As important, however, is the discovery of new neuronal death pathways that could lead to neuron loss in ways that have previously not been appreciated. This latter set of observations opens up the possibility that a gateway to seizure-induced neuronal loss involves signaling pathways that represent or are influenced by early neuron-death pathways. Thus, we propose that – in addition to the previously proposed *recapitulation-of-development*

mechanisms – another hypothesis could be the *neuronal-death-pathway* hypothesis, whereby the early steps of neuronal death generate signals that promote epileptogenesis even if the neurons ultimately do not die. An attractive feature of this hypothesis is that it could lend itself to classification by molecular markers that reflect these neuronal pathway molecules. This hypothesis might also explain why neuronal death seems so important to acquired epileptogenesis, yet might in some cases be unnecessary.

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# How Is Homeostatic Plasticity Important in Epilepsy?

# 10

John W. Swann and Jong M. Rho

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## Abstract

Maintaining physiological variables within narrow operating limits by homeostatic mechanisms is a fundamental property of most if not all living cells and organisms. In recent years, research from many laboratories has shown that the activity of neurons and neural circuits are also homeostatically regulated. Here, we attempt to apply concepts of homeostasis in general, and more specifically synaptic homeostatic plasticity, to the study of epilepsy. We hypothesize that homeostatic mechanisms are actively engaged in the epileptic brain. These processes attempt to re-establish normal neuronal and network activity, but are opposed by the concurrent mechanisms underlying epileptogenesis. In forms of intractable epilepsy, seizures are so frequent and intense that homeostatic mechanisms are unable to restore normal levels of neuronal activity. In such cases, we contend that homeostatic plasticity mechanisms nevertheless remain active. However, their continuing attempts to reset neuronal activity become maladaptive and results in dyshomeostasis with neurobehavioral consequences. Using the developing hippocampus as a model system, we briefly review experimental results and present a series of arguments to propose that the cognitive neurobehavioral comorbidities of childhood epilepsy result, at least in part, from unchecked homeostatic mechanisms.

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## Keywords

Homeostasis • Synaptic plasticity • Seizures • Synapses • Epileptogenesis • Learning and memory

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Homeostasis or the maintenance of a physiological state – despite external or internal disturbances that would be expected to alter that state – is a biological concept central to both animal and human physiology [27]. This concept was first introduced nearly 150 years ago by Claude Bernard who demonstrated the ability of organisms to maintain a relatively constant internal environment and who stated that the maintenance of “*le milieu interieur*” was essential for life [3]. In 1929, Walter Cannon extended these concepts and coined the term homeostasis or “similar state” [7]. In adopting a systems level approach to physiology, Cannon suggested that coordinated adjustments of interacting systems through feedback systems result in the maintenance of physiological parameters such as body temperature and circulating oxygen levels within a set range.

Today, the concept of homeostasis is fundamental to all studies of physiology from the level of individual cells to that of entire organisms. Indeed, it is such a well-accepted concept that it is taken for granted that the underlying basic mechanisms are essential for survival. Homeostasis includes, but is not limited to, the regulation of blood pH, circulating levels of glucose, body temperature, interstitial level of O<sub>2</sub> and CO<sub>2</sub> as well as critical electrolytes such as Na<sup>+</sup>, Cl<sup>-</sup>, K<sup>+</sup> and Ca<sup>+2</sup>. In keeping with the work of Cannon, feedback systems are thought to be the primary regulatory mechanisms underlying homeostasis. In general, these feedback systems must first detect a change in a parameter that needs to be held within narrow limits – also referred to as a set point. If it deviates from these limits, the system activates mechanisms to return the parameter to the set point. Such systems consist of a sensor that is able to measure changes in the parameter, an integrator that compares the detected information to the desired set point and an effector that generates the compensatory response to return the parameter to homeostasis. Circulating levels of Ca<sup>+2</sup> are a good example of such a feedback system. When blood Ca<sup>+2</sup> falls below its set point, Ca<sup>+2</sup>-sensing receptors in cells of the parathyroid gland are activated. This results in the release of parathyroid hormone which acts to increase circulating Ca<sup>+2</sup> levels. Multiple effector mechanisms

are induced, including increased absorption from the gastrointestinal tract and reabsorption from urine. Another important source of Ca<sup>+2</sup> is bone. Parathyroid hormone increases the activity of bone-degrading osteoblasts which release Ca<sup>+2</sup> from bone and thereby return circulating Ca<sup>+2</sup> to its set point.

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## 10.1 Neuronal Homeostatic Plasticity

Over the past 15 years, a great deal of evidence has accumulated to suggest that the physiological activity of neurons and neuronal networks are homeostatically regulated [33, 34]. Neuronal networks of the central nervous system (CNS) are highly dynamic. This is easily observed in variations in human and animal EEG recordings over a 24-h period. For example, the dramatic alterations in recordings at transitions from NREM sleep to the awake state reflect marked changes in the activity of individual neurons and the operations of their networks. At these times, an organism in interacting with its environment will store information for future use. Hebbian synaptic plasticity mechanisms such as long-term potentiation (LTP) and long-term depression (LTD) are widely thought to underlie the processes for learning and the storage of memories. However, for some time, theoretical neurophysiologists have recognized that Hebbian plasticity should destabilize and consequently interfere with network operations [1]. The idea behind this claim is that once a group of excitatory synapses undergo a use-dependent form of plasticity – like LTP – they will in the future produce larger excitatory post-synaptic potentials (EPSPs) which will more likely induce action potentials in the postsynaptic neuron. This in turn results in even larger EPSPs and more neuronal firing, and a self-perpetuating cascade of ever-increasing synaptic strengthening and ultimately increasing network excitability.

Homeostatic synaptic plasticity has been proposed as a stabilizing mechanism to counter the potential run-away excitation of Hebbian plasticity [33]. As would be expected, this relatively

new field of homeostatic plasticity has borrowed concepts from other forms of physiological homeostasis. For example, synaptic homeostasis has been defined as “a form of plasticity that acts to stabilize the activity of a neuron or neuronal circuit around some *set point* value” [33]. Possibly the best demonstration of homeostatic plasticity comes from studies of dissociated cultures of CNS neurons. In these models, when the networks of cortical neurons are pharmacologically challenged by application of a  $\gamma$ -aminobutyric acid type A (GABA<sub>A</sub>) receptor antagonist such as bicuculline, the firing rates of individual neurons initially increase. However, over a period of many hours to days, firing rates return to their original rate, which is interpreted to be the homeostatic set point of affected neurons and their networks. Similarly, when activity is suppressed, firing rates are initially very low but are restored over time [35].

There have been many studies of the cellular and molecular events underlying this form of neuronal plasticity. It has become clear that there are numerous mechanisms that can act independently to regulate post-synaptic and pre-synaptic strength as well as mechanisms that operate at the level of individual synapses, and others that act in parallel but on a more global scale. The most studied of this type on neuronal plasticity is synaptic homeostasis – particularly at excitatory, glutamatergic synapses. In these studies, following periods of pharmacologically-induced heightened neuronal activity, the amplitude or strength of miniature excitatory post-synaptic currents (mEPSCs) has been shown to decrease at the times when neuronal firing rates had returned to their set point [35]. These alterations in mEPSCs were also found to parallel decreases in the amplitude of evoked synaptic events. Much evidence has emerged suggesting that synaptic homeostasis results from post-synaptic alterations in glutamatergic subunit expression and localization. Under some experimental conditions, a decrease in the number of glutamatergic synapses has also been demonstrated [13]. In other cases and circumstances, pre-synaptic changes in transmitter release have been reported [4, 6].

In addition to synaptic homeostasis of glutamatergic synapses, the function of inhibitory synapses and inhibitory interneurons appear to be homeostatically regulated, as are the intrinsic excitability properties of individual neurons. As might be expected, synaptic inhibition is regulated in the opposite direction of excitation. For instance, when neuronal activity is experimentally depressed as the amplitude of mEPSCs in pyramidal cells is increased, miniature inhibitory post-synaptic currents (mIPSCs) decrease in strength [15, 20]. Both pre-synaptic and post-synaptic changes appear to contribute to homeostatic regulation of synaptic inhibition. The variety of mechanisms underlying this form of regulation may be a reflection of the diversity of inhibitory synapses and inhibitory interneurons in the CNS. For instance, when ascending activity to the visual cortex is experimentally lowered (in an attempt to mimic activity suppression *in vitro*), the amplitude of inhibitory synapses onto layer 4 pyramidal cells from fast spiking interneurons is reduced [22]. However, inhibitory synapses from other interneuronal subtypes appear to be stronger although fewer in number.

In terms of the intrinsic excitability of neurons, it has been shown that when the activity of cultured neurons is experimentally suppressed, the intrinsic excitability of excitatory neurons is enhanced [9]. So these neurons are able to generate more action potentials in response to a given synaptic input than untreated control neurons. Thus, at the same time that excitatory synaptic transmission is increased and synaptic inhibition is decreased, alterations in the expression and function of ion channels (likely both inward and outward voltage-dependent currents) further enhance neuronal excitability in attempting to re-establish normal neuronal and ultimately network activity.

In summary, numerous studies over the past 15 years have not only repeatedly demonstrated the ability of neurons to homeostatically adapt to experimentally-induced alterations in their activity but have also shown that the cells have a wide array of mechanisms at their disposal to stabilize

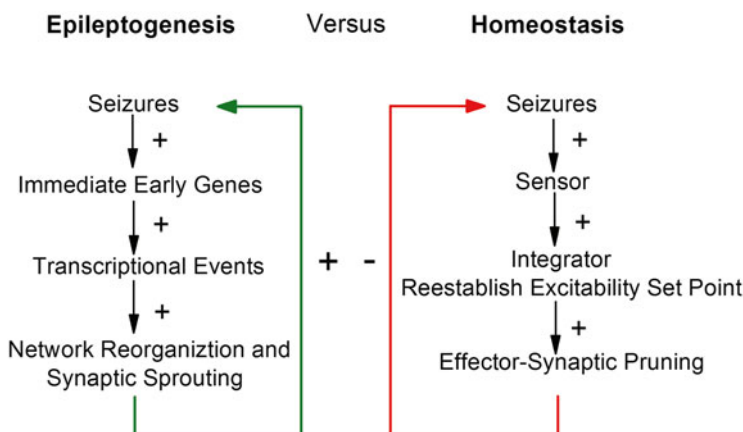
their activity in the face of forces like Hebbian synaptic plasticity that can potentially lead to neuronal and network instability.

## 10.2 Homeostatic Plasticity Versus Epileptogenesis

The neuroscience community has learned a great deal from studies of the basic mechanisms of homeostatic synaptic plasticity. This has been propelled by the use of relatively simple culture systems that are amenable to rigorous experimental manipulations and testing of hypotheses. A number of other studies have been performed *in vivo* in attempts to extend information from *in vitro* studies to the intact CNS. However, how these results impact our understanding of epilepsy is now only beginning to be explored. For at least the past 15 years, a large proportion of experimental epilepsy research efforts have been focused on understanding the mechanisms underlying epileptogenesis, a process that at least superficially appears to be the antithesis of homeostasis. The term epileptogenesis has been defined as a chronic process by which normal brain

is transformed into tissue capable of generating spontaneous recurrent seizures [17]. In the acquired epilepsies (e.g. following traumatic brain injury), the seizure-prone state is thought to arise from a progressive series of molecular, cellular and circuit changes that evolve over time.

Results from long-term continuous video-EEG recordings in several animal models of acquired epilepsy have emphasized the progressive nature of epileptogenesis [11]. Within a week after injury, nonconvulsive seizures are first observed. A week thereafter they become convulsive. Seizure frequency can gradually increase nearly tenfold over the ensuing 3–4 months. Such results are consistent with the idea first proposed by Gowers in 1888 that “seizures beget seizures” [14]. Potential steps in the progression of epileptogenesis are illustrated by the positive feedback system in Fig. 10.1. Here, seizures are envisioned to induce a cascade of molecular and cellular events that lead to sprouting of glutamatergic synapses and other forms of network reorganization that further enhance network excitability and the genesis of more seizures with increasing frequency. Juxtaposed to this is a diagram of the negative feedback loop that is thought to characterize



**Fig. 10.1** Diagram outlining the hypothesized opposing forces of epileptogenesis and homeostasis. A positive feedback loop is envisioned to mediate epileptogenesis. Examples of some of the potential molecular events are named that lead to the network reorganization and synaptic sprouting that is thought to contribute to recurring seizures.

Homeostasis is suggested to oppose epileptogenesis through negative feedback loops that are designed to re-establish normal neuronal and neural circuit excitability. Pruning of glutamatergic synapses is but one example of an effector mechanism that would reduce seizure generation

homeostasis in general and homeostatic plasticity more specifically. In this context, seizures are envisioned to activate sensory processes – just as increased neuronal activity is thought to during synaptic homeostasis. A molecular integrator compares neuronal activity to a set point value and induces changes in an effector, which in this example results in the pruning of excitatory synapses [13] and which in turn would be predicted to reduce neuronal and network excitability and reduce seizure frequency.

If homeostasis is such a fundamental property of animal and human physiology, why does it apparently fail in epilepsy? One possibility is that it does not always fail since seizures do spontaneously remit in some forms of epilepsy without any apparent reason. Many of the so-called benign epilepsies of infancy and childhood carry a favorable prognosis. In these instances, children are simply said to “outgrow their seizures”. The mechanisms accounting for these observations are unknown. One possibility is that as the brain matures the developmental factors that enhance seizure susceptibility are no longer operant. Alternatively, ongoing homeostatic mechanisms may play a significant role in these remissions. In contrast, in more severe and intractable epilepsy, it seems possible that homeostatic mechanisms are actively engaged in epilepsy but in many cases the precipitating injury (or in the case of genetic forms of epilepsy, i.e., the consequences of gene mutations) are so severe that homeostatic mechanisms are simply unable to re-establish neuronal activity to the desired set point and seizure progression continues unabated by the processes underlying epileptogenesis. However, if this were the case, then as seizures recur, homeostatic mechanisms would also be repeatedly induced in an attempt to re-establish normal neuronal excitability and network stability.

There are a number of observations made in animal models of epilepsy that appear somewhat paradoxical in that molecular and anatomical changes observed would be expected to prevent seizures, not promote them. Increases in GABA<sub>A</sub> receptor subunits [16] and potassium channel expression [26, 28] as well as dendritic spine loss

in hippocampal and neocortical pyramidal cells [5, 30] are just a few examples of alterations that researchers have sometimes referred to as “paradoxical” and possibly “compensatory” responses to on-going seizure activity. It is not hard to imagine that there are many other such paradoxical findings that remain unpublished since they could not be explained in the context of epileptogenesis or the seizures that were being studied. However, such observations could be indicators of homeostatic processes taking place.

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### 10.3 Homeostasis and Seizures During Brain Development

The developing brain is well known to be highly susceptible to seizures. However, during the first 2–3 weeks of life in rats and mice, in general neither prolonged seizures nor recurrent seizures usually lead to the genesis of epilepsy later in life (but this remains unclear and controversial in the clinical setting). Nonetheless, these seizures are not without significant consequences since numerous studies have shown that they routinely produce deficits in learning and memory – particularly in spatial learning [21]. Several labs have begun to explore the underlying mechanisms. For example, recent studies of hippocampal CA1 pyramidal cells suggest that place cell function is impacted by recurrent early-life seizures. Place cells are thought to provide an animal with a spatial map of its environment and serve as surrogate markers for spatial memory. Among other observations, investigators have shown that place cells are unable to form stable maps in animals that experienced early-life seizures [19]. Further, rats exposed to early prolonged hyperthermia-induced seizures exhibit a significant increase in hippocampal T2 MRI signal intensity which is associated with spatial memory deficits [10].

In exploring the underlying mechanisms of learning and memory deficits, investigators have understandably focused on alterations in central excitatory and inhibitory synapses. Early-life seizures have been shown to profoundly affect

synaptic signaling through both glutamate and GABA<sub>A</sub> receptors. For example, hypoxia-induced seizures in postnatal day 10 rats results in a decrease in silent N-methyl-D-aspartate (NMDA) synapses and an attenuation of hippocampal LTP that persists into adulthood [36].

Changes in inhibitory neurotransmission can also play an important role in processes critical for learning and memory. Enhancement of GABA<sub>A</sub> receptor signaling is known to impair LTP, and studies have shown increased inhibition after early-life seizures. For example, after both hyperthermia- and kainate-induced seizures, there is enhanced paired-pulse inhibition in the hippocampus [29] and selective increases in specific GABA<sub>A</sub> receptor subunits, notably the  $\alpha 1$  subunit, after status epilepticus induced by either lithium-pilocarpine or kainate at postnatal day 20 [25]. However, it should be noted that GABA<sub>A</sub> receptor subunit changes following seizure activity are age-dependent as are the responses to agonists. Importantly, GABA<sub>A</sub> receptor activation in neonatal neurons results in membrane depolarization, in contrast to the normal hyperpolarizing response seen in mature neurons – a result of differential expression of the cation-co-transporters KCC2 and NKCC1 in early post-natal brain development which establishes the transmembrane chloride electrochemical gradient [2]. Early-life seizures have been reported to promote the developmental switch from depolarizing to hyperpolarizing, one consequence of which may be impaired spatial learning and memory [12].

In addition to molecular receptor changes affecting both excitatory and inhibitory neurotransmission, a number of studies have reported decreases not only in dendritic spine density but also dendrite length and branching complexity in hippocampal pyramidal cells [18, 23]. Similar abnormalities in dendrite morphology have been reported in human epilepsy [30]. However, in experimental studies of seizure induction in early-life, dendritic changes have been observed after a series of seizures that do not lead to epilepsy later in life and presumably do not induce a significant epileptogenic process. For instance,

when 15 brief (~3 min in duration) seizures are induced over a 5-day period (3 seizures per day) in 1 week old mice, dendrite length and branching complexity are reduced by 25 % compared to control mice, and as adults these same mice are learning impaired [23]. Changes in CA1 dendrite arborization are observed within 1 week after the last seizure and have been shown to be the result of dendrite growth suppression. Very similar observations of growth suppression have been made in hippocampal slice cultures [24]. In these instances, slice cultures from 5 day-old mice are grown under conditions that produce recurring seizure-like activity. Within 24 h of initiating epileptiform activity, CA1 pyramidal cell dendrites are shorter in length and have fewer branches than pyramidal cells from sister control cultures. Moreover, over time, while dendrites in control slices continue to grow, dendrites in slices that are undergoing seizure-like activity do not. Similar to the studies of synaptic homeostatic plasticity in dissociated cultures discussed earlier, mEPSC amplitude and frequency (recorded in pyramidal cells) are reduced in slice cultures following a few days of treatment [31]. Remarkably, a very recent report has shown that similar changes in excitatory synaptic transmission and dendrite arborization can be observed after only a few hours of synchronized epileptiform activity [8]. Collectively, these results suggest that seizures may not only suppress on-going dendrite growth but acutely may even induce a retraction of growing dendritic branches.

One interpretation of these results is that the seizures *in vivo* and seizure-like activity *in vitro* are activating homeostatic mechanisms in attempts to limit neuronal excitability, re-establish network excitability *in vitro* and prevent the occurrence of future seizures. However, by limiting the branching complexity of hippocampal pyramidal cell dendrites, the number of excitatory glutamatergic synapse present on dendrites should also be reduced. Indeed, biochemical results have consistently shown reduced expression of markers for glutamatergic synapses, such as PSD95, in the hippocampus taken from mice that have experienced recurring early-life seizures and in slice



culture that have undergone chronic epileptiform activity [31, 32]. With a reduction in the number of glutamatergic synapses, one might predict deficits in hippocampal-based learning and memory. This is because these synapses are well known to undergo Hebbian forms of synaptic plasticity, such as LTP and LTD, which are thought to contribute in important ways to the formation of memories (see earlier discussion). Thus, in attempting to re-establish normal neuronal and network excitability, homeostatic mechanisms may also limit an animal's capacity to learn since some of the anatomical substrates for learning have been eliminated.

In such situations, homeostatic mechanisms could become maladaptive or dyshomeostatic, where these mechanisms are driven to such extremes that they have undesirable consequences. Regulating the circulating levels of  $\text{Ca}^{+2}$  that was discussed earlier provides an example of such a phenomenon. In some clinical situations, blood  $\text{Ca}^{+2}$  levels can fall below its set point for prolonged periods of time. Dietary deficiency is one cause of low blood  $\text{Ca}^{+2}$ . Under these circumstances, calcium sensing cells in the parathyroid gland release parathyroid hormone which activates osteoblasts in bone resulting in the release of  $\text{Ca}^{+2}$  into the blood in attempt to restore circulating  $\text{Ca}^{+2}$ . However, intense activation by parathyroid hormone will eventually lead to bone dissolution, cavitations of the skeleton and increased susceptibility to bone fractures. Similarly, it seems possible that uncontrolled seizures may induce neuronal homeostatic responses that in attempting to limit neuronal hyperexcitability results in impaired synaptic plasticity and learning deficits.

It is thought that synaptic homeostatic plasticity and Hebbian synaptic plasticity are normally complementary processes. While Hebbian plasticity occurs from moment-to-moment, homeostatic mechanisms occur more slowly, over hours and days and function to prevent runaway excitation or inhibition but do not to interfere with rapid information transfer and storage. However, in epilepsy where abnormal – and often extreme neuronal hyperexcitability – exists, homeostatic

mechanisms appear unable to reset neuronal excitability levels to something approaching normal. But by continually attempting to reset normal levels, homeostasis may be driven to such extremes that it limits Hebbian plasticity and interferes with information processing.

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## 10.4 Concluding Remarks

At this time, some may not be convinced that homeostatic mechanisms are active in the epileptic brain and more direct evidence is needed to support the notion that the cognitive neurobehavioral comorbidities of epilepsy are at least in part a consequence of homeostasis and homeostatic imbalance. Currently, the challenge is in developing ways to study such hypothetical seizure-induced homeostatic mechanisms in relative isolation and in greater detail without the confound of the myriad molecular, cellular and genetic processes that are active in epileptogenesis. The developing hippocampus may serve as a useful model system in this regard since at least under some experimental conditions homeostatic mechanisms appear to predominate over mechanisms of epileptogenesis. The future may provide better experimental opportunities and researchers should be prepared to exploit them. Ultimately, a full understanding of the molecular mechanisms underlying seizure-induced homeostasis will be required. It seems that employment of relatively simple *in vitro* systems (e.g. dissociated or slice cultures) would accelerate discovery. However, key findings *in vitro* will always need to be validated *in vivo*. Under some experimental situations (e.g. the prolonged seizures of status epilepticus) neuronal injury and death may occur and should be avoided if possible. Being able to discriminate between injury-induced changes and homeostatic-induced mechanisms will be critical. However, currently neuroscience researchers have a wealth of new and powerful cellular and molecular tools at their disposal that should make such studies possible. Live time-lapse imaging of neurons in which molecular biomarkers of suspected key contributors of seizure-induced homeostasis can be visualized

is but one example of the types of experiments that should be possible. Once homeostatic mechanisms have been well characterized and ways to selectively eliminate them have been discovered, returning to more complex situations where epileptogenesis and homeostatic plasticity co-exist will be important not only to definitively prove that homeostasis is active in epilepsy but also to understand the costs and benefits of suppressing or enhancing these homeostatic processes.

**Acknowledgement** From John Swann: I remember first meeting you at the 1981 SfN Meeting. At that point, I had inadvertently stumbled into epilepsy research but became intrigued by how little was known about the basic mechanisms of the childhood epilepsies. You had recently published your first papers on epileptiform activity in immature hippocampal slices and I was in the midst of somewhat similar experiments as a newly minted independent investigator. During our conversation, you encouraged me to continue my line of investigation even though we were potential competitors. Your generous gesture contributed importantly to my commitment to epilepsy research and I have valued you as a friend and colleague throughout the intervening years. I think this book in many ways reflects the positive influence you have had on the careers of so many scientists – in epilepsy and the neurosciences more generally. This is a legacy to be proud of and emulated by your students and the future generations of their students.

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# Is Plasticity of GABAergic Mechanisms Relevant to Epileptogenesis?

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## Abstract

Numerous changes in GABAergic neurons, receptors, and inhibitory mechanisms have been described in temporal lobe epilepsy (TLE), either in humans or in animal models. Nevertheless, there remains a common assumption that epilepsy can be explained by simply an insufficiency of GABAergic inhibition. Alternatively, investigators have suggested that there is hyperinhibition that masks an underlying hyperexcitability. Here we examine the status epilepticus (SE) models of TLE and focus on the dentate gyrus of the hippocampus, where a great deal of data have been collected. The types of GABAergic neurons and GABA<sub>A</sub> receptors are summarized under normal conditions and after SE. The role of GABA in development and in adult neurogenesis is discussed. We suggest that instead of “too little or too much” GABA there is a complexity of changes after SE that makes the emergence of chronic seizures (epileptogenesis) difficult to understand mechanistically, and difficult to treat. We also suggest that this complexity arises, at least in part, because of the remarkable plasticity of GABAergic neurons and GABA<sub>A</sub> receptors in response to insult or injury.

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## Keywords

GABA • GABA<sub>A</sub> receptor • Interneuron •  $\alpha$ 1 subunit • Chloride channel • Granule cell • Adult neurogenesis • Status epilepticus • Febrile seizures • Aberrant neurogenesis • Ectopic granule cell

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## 11.1 Introduction

In the nineteenth century, the idea that epilepsy was a brain disorder arose as a consequence of the relatively new discipline of neurology. In the latter half of the twentieth century, many studies showed that chemicals such as penicillin, a GABA<sub>A</sub> receptor (GABA<sub>A</sub>R) antagonist, caused experimental seizures or epileptiform activity when applied to the neocortex of animals. Philip Schwartzkroin played a major role in the development and refinement of these ideas by the use of the hippocampal slice preparation [131, 132, 152]. One view that emerged was that epilepsy might be caused by defects in inhibition, which was supported by pharmacological experiments showing that several anticonvulsants, such as the barbiturates and benzodiazepines, exerted their actions by facilitating the actions of GABA at GABA<sub>A</sub>Rs [88, 109].

The idea that epilepsy is caused by insufficient GABAergic inhibition has developed more support as it has become clear that some types of GABAergic neurons are vulnerable in animal models of epilepsy, or lost in tissue resected surgically from patients with intractable epilepsy [78, 126, 127]. In addition, mutations in the subunits of the GABA<sub>A</sub> receptor have been identified as a basis of some genetic epilepsy syndromes, such as Genetic Epilepsy with Febrile Seizures+ (GEFS+) which can be caused by a point mutation in the *GABRG<sub>2</sub>* gene which normally encodes the  $\gamma$  subunit of the GABA<sub>A</sub>R [4, 159]. However, many arguments have also been made that epilepsy cannot be explained solely by a defect in GABAR-mediated inhibition. Some of the opposing views have come from studies of GABAergic agonists, which exacerbate some types of seizures instead of inhibiting them. For example, drugs that enhance GABAergic inhibition increase absence seizures instead of suppressing them. The explanation is related to the actions of GABA at GABA<sub>B</sub> receptors on thalamocortical relay cells. By enhancing the actions of GABA to hyperpolarize relay cells, T-type Ca<sup>2+</sup> current in relay cells are strongly deactivated, leading to more robust bursts of

action potentials in relay cells when the hyperpolarizations end; these rebound bursts drive the thalamocortical oscillation [58, 141].

In the last 20 years, a wealth of new information about GABA and GABA<sub>A</sub>Rs has been published using animal models of epilepsy and clinical research. One of the complexities that has emerged is the plasticity of GABAergic mechanisms. This plasticity is remarkable because it involves many aspects of GABAergic transmission: the numbers of GABAergic neurons and the locations of their axons; the synthesis, release and uptake of GABA; and alterations in GABA receptors. Although the contribution of GABAergic mechanisms, and their plasticity, to epilepsy is still an area of active research, it seems unlikely that there is simply too little GABA in epilepsy – or too much. Instead, GABAergic transmission is very different in epilepsy compared to the normal brain. This concept, that GABAergic inhibition is not simply deficient in epilepsy, is consistent with the relatively normal function of individuals with epilepsy during the interictal state.

We discuss below the basic characteristics of GABAergic transmission in the normal and epileptic condition to clarify this idea. For the epileptic condition, we focus on temporal lobe epilepsy (TLE) where this concept appears to be particularly relevant. We also focus on the dentate gyrus (DG) in animal models where status epilepticus (SE) is used to produce spontaneous recurrent seizures and simulate acquired TLE. The reason for this focus is that the data that are available for this context are extensive. However, these models have been criticized because they do not simulate all aspects of TLE.

Most of the discussion below addresses the ways that GABAergic circuitry are changed by SE and alterations in GABA<sub>A</sub>Rs in DG granule cells (GCs). Presynaptic GABA<sub>A</sub>Rs and effects of GABA<sub>A</sub>Rs on other cell types are also important to consider in the context of the DG and epilepsy, and are reviewed elsewhere [70]. Regulation of GABA<sub>A</sub>Rs by phosphorylation also has implications for the dynamics of GABAergic transmission in epilepsy; effects relevant to the DG are discussed below and

additional issues are described elsewhere [83, 155]. Finally, GABA<sub>B</sub>Rs clearly have a role in epilepsy, but are outside the scope of this discussion and readers are referred to excellent reviews published previously [14, 84].

## 11.2 GABAergic Transmission in the Normal Adult Dentate Gyrus (DG)

### 11.2.1 GABAergic Neurons in the DG of the Adult Rodent

Figure 11.1 illustrates the fundamental circuitry of the DG in the normal adult rodent [2]. The principal cell of the DG is the granule cell (GC), which uses glutamate as its primary neurotransmitter, but also has the capacity to synthesize GABA, especially after seizures (discussed further below). GCs also synthesize numerous peptides that are packaged in dense core vesicles and behave as co-transmitters [55]. The peptides are numerous: dynorphin [25], leu-enkephalin [153], brain-derived neurotrophic factor [125], and others. The major afferent input to the GCs is the perforant path projection from entorhinal cortical neurons in layer II [161]. The GCs form the major output from the DG, the “mossy fiber” pathway, which innervates neurons in the hilus and area CA3 [2]. There is another glutamatergic neuron in the DG, located in the hilus, which is called a mossy cell (for reviews see [53, 126]). The major afferent input to mossy cells comes from the GCs, and mossy cells project to GCs and GABAergic neurons within the DG [126].

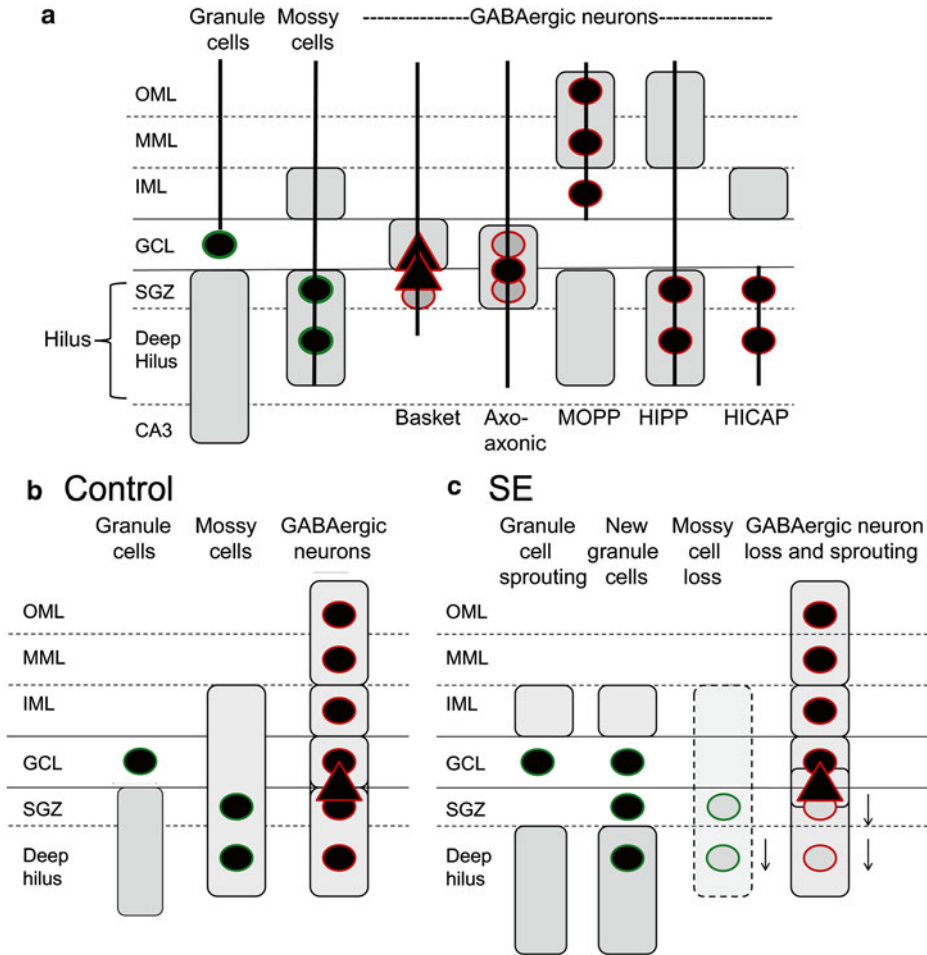
There are many other types of neurons in the DG, and they use GABA as a neurotransmitter. Most of the GABAergic neurons have an axon that projects primarily in the area surrounding the cell body, similar to other cortical circuits where most of the GABAergic neurons are local interneurons. However, there are several subtypes of DG interneurons that also have axons that project to distant areas of the DG, such as the contralateral DG [34, 49]. Like GCs, GABAergic neurons of the DG also use peptides as co-transmitters [55, 138], and after seizures, some of the peptides in

GCs are the same peptides as those in GABAergic neurons (e.g., neuropeptide Y; NPY; [120]).

The primary type of GABAergic neuron in the DG is the basket cell, which makes basket-like endings around GC somata. It initially was described as a pyramidal-shaped neuron with somata at the base of the GC layer (on the border of the GC layer and the hilus) but the location, somatic morphology and other characteristics are actually diverse [115]. Furthermore, some of the basket cells with pyramidal shaped somata have axons that project to the contralateral DG [49]. There also is variation in neuropeptide content in pyramidal-shaped GABAergic neurons, ranging from parvalbumin, cholecystokinin, to substance P [55, 81, 139]. Electrophysiologically, these cells also vary, although they fit the general characteristics of interneurons because they have a very large afterhyperpolarization following single action potentials [115]. They inhibit their postsynaptic targets by opening chloride channels of GABA<sub>A</sub>Rs at the soma. Because the resting potential of GCs is close to the reversal potential for chloride or hyperpolarized to it, chloride entry depolarizes the GC rather than hyperpolarizing it, shunting currents that would otherwise reach threshold for action potential (AP) generation; for this reason, “shunting inhibition” is probably the main inhibitory effect of basket cells, rather than hyperpolarization.

Another very important inhibitory cell type also inhibits AP generation of GCs, but is slightly different because it primarily innervates the axon hillock, rather than the somata of GCs. This cell type, the axo-axonic cell, is similar to chandelier cells in neocortex [142] in that chandelier-type endings envelope the axon hillock of GCs. The cell bodies of axo-axonic cells are variable and many types of neuropeptides are co-localized with GABA. The intrinsic electrophysiology of axo-axonic cells is consistent with fast-spiking interneurons [22].

Another type of DG interneuron is the so-called HIPP cell, named because it has a *H*ilar cell body and projects to the outer 2/3 of the molecular layer, where the *p*erforant *p*ath projection terminates. This neuronal subtype usually expresses somatostatin and NPY [145] and has



**Fig. 11.1 DG circuitry in the normal adult rodent and following status epilepticus (SE).** (a) Circuitry of the normal rodent DG is shown schematically. Cell bodies outlined in *green* are glutamatergic; those cells outlined in *red* are GABAergic. *Black circles* indicate the primary location of the somata; *grey circles* are secondary locations. *Grey rectangles* indicate the location of the axon terminals. Abbreviations of the lamina of the DG are as follows: *OML* outer molecular layer, *MML* middle molecular layer, *IML* inner molecular layer, *GCL* granule cell layer, *SGZ* subgranular zone. MOPP, molecular layer cell body, axon in the terminal field of the perforant

path; HIPP, hilar cell body, axon in the terminal field of the perforant path. HICAP, hilar cell body, axon in the terminal field of the commissural/associational projection (Adapted from Freund and Buzsaki [42]). (b) A summary of a. (c) Changes in the DG circuitry following SE are diagrammed. After SE, changes are as follows: GC axons sprout into the IML; newborn GCs are born and some migrate into the hilus and GCL; many mossy cells are lost (indicated by the *arrow*, *light cell body color* and *dotted line* around the axon plexus); some GABAergic neurons are lost and others sprout into several layers (For references, see text)

axon collaterals primarily in the molecular layer [52], with a less dense projection in the hilus [35]. It has been suggested that it inhibits the EPSPs produced by the perforant path input, presumably by innervating GC dendrites and shunting EPSPs traveling to the GC soma. HIPP cells may

also inhibit glutamate release from perforant path terminals because they make synapses on the terminals [80]. The electrophysiology of HIPP cells is characteristic of interneurons generally [44], but it has been noted that they are relatively slow spiking [2, 115] and have a pronounced

'sag' in response to hyperpolarizing current commands [89]. This cell type has attracted a lot of attention in epilepsy research because these cells are relatively vulnerable to insults or injury [116, 126]. Several mechanisms have been proposed for their vulnerability, such as STAT3 expression [29]. It has also been shown that p75<sup>NTR</sup> receptors are present on the septocholinergic terminals that innervate the HIPP cells, and can cause their death when the septocholinergic pathway is lesioned [37, 38].

Analysis of the numbers of GABAergic neurons using immunocytochemical markers and stereological techniques has led to estimations that the majority of DG interneurons are basket cells or axo-axonic cells, which express parvalbumin or CCK. The other major subtype of DG interneuron is hilar HIPP cells, which co-express GABA and NPY or somatostatin (for reviews see [55, 81]). However, many other types of DG interneurons exist: MOPP cells [28], ivy cells and neurogliaform cells [3] and hilar neurons that innervate the inner molecular layer (HICAP cells; [51, 52]).

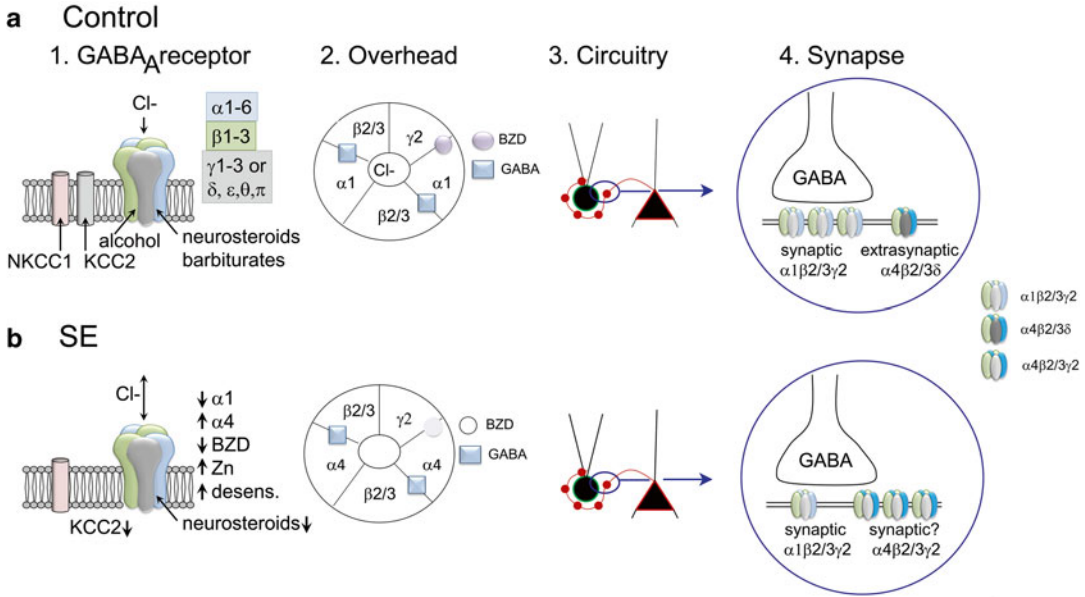
The major afferents to DG interneurons are the perforant path, GCs, and mossy cells. In addition, there is extrinsic input from the ascending serotonergic, cholinergic, and noradrenergic nuclei. The primary effects appear to be inhibitory [41]. In addition, there are additional inputs to the DG from areas outside the hippocampus that are not well understood functionally, such as the supramammillary input [74]. Many neuromodulators, such as endocannabinoids, have been shown to exert striking effects in the DG [40], but how all the neuromodulators act in concert in the awake behaving animal is still unclear.

### 11.2.2 GABA Receptors in the Normal Adult GC

Post-synaptic GABA<sub>A</sub>Rs mediate most fast synaptic inhibition in the forebrain (Fig. 11.2). GABA<sub>A</sub>Rs are heteromeric protein complexes composed of multiple subunits that form ligand-gated, anion-selective channels whose properties

are modulated by barbiturates, benzodiazepines, zinc, ethanol, anesthetics and neurosteroids. There are several different GABA<sub>A</sub>R subunit families and multiple subtypes exist within each of these subtypes ( $\alpha$ 1-6,  $\beta$ 1-4,  $\gamma$ 1-3,  $\delta$ ,  $\epsilon$ ,  $\pi$ ,  $\Phi$ ). The most common GABA<sub>A</sub>R is the  $\alpha$ 1 $\beta$ 2 $\gamma$ 2 subtype, but multiple subtype combinations exist and they vary in different brain regions and cell types, and during different times in development [73, 111, 134]. Subunit composition of GABA<sub>A</sub>Rs plays a major role in determining the intrinsic properties of each channel, including affinity for GABA, kinetics, conductance, allosteric modulation, probability of channel opening, interaction with modulatory proteins, and subcellular distribution [77, 97, 134]. For example, alterations in the  $\alpha$ -subtype results in differences in receptor kinetics, membrane localization and GABA<sub>A</sub>R modulation by benzodiazepines and zinc [87, 97, 140, 154]. In the GC, GABA<sub>A</sub>Rs that contain  $\alpha$ 1 subunits paired with  $\gamma$ 2 subunits are sensitive to benzodiazepines and generally located at the synapse, contributing to phasic inhibition, a term that refers to the effects of GABA released at GABAergic synapses that binds to postsynaptic receptors located at the synaptic cleft. These effects are primarily related to increased conductance when chloride channels open, and hyperpolarization of postsynaptic membrane potential when chloride influx occurs. However, as mentioned above, when the postsynaptic membrane potential is hyperpolarized relative to  $E_{Cl^-}$ , which may occur in GCs, there is a depolarization. GABA<sub>A</sub>Rs that contain  $\alpha$ 4 subunits have unique pharmacological properties, such as insensitivity to benzodiazepines and increased sensitivity to zinc blockade. Receptors containing  $\alpha$ 4 subunits are most often found with the  $\delta$  rather than the  $\gamma$  subunit in combination with  $\alpha\beta$ . These  $\alpha$ 4 $\beta\delta$  GABA<sub>A</sub>Rs are localized to extrasynaptic sites and contribute to tonic inhibition, which refers to the basal inhibitory current produced by low concentrations of extracellular GABA that are present outside of the synapse (resulting from diffusion from synaptic to extrasynaptic space). Under physiological conditions, only a minor population of  $\alpha$ 4 $\beta\gamma$ 2 GABA<sub>A</sub>Rs are found at synapses of GABAergic neurons on





**Fig. 11.2 GABA<sub>A</sub> receptor subunits in dentate gyrus (DG) granule cells (GCs) in the normal adult rodent and following SE.** (a) Control conditions. (1) The subunits of the GABA<sub>A</sub> receptor (GABA<sub>A</sub>R) are diagrammed, with sites of modulation noted. The location of the K<sup>+</sup>Cl<sup>-</sup> cotransporters NKCC1 and KCC2 are depicted schematically. (2) An overhead view of a typical GABA<sub>A</sub>R in a normal adult GC. It has  $\alpha 1$ ,  $\beta 2/3$  and  $\gamma 2$  subunits with two sites for GABA and a benzodiazepine (BZD) site for modulation. (3) The prototypical GABAergic neuron in the DG is the basket cell (triangle) which has an axon that encircles GC somata, making periodic GABAergic synapses. (4) A schematic of the GABAergic synapse in control conditions has synaptic  $\alpha 1\beta 2/3\gamma 2$  receptors and extrasynaptic

receptors that contain different subunits ( $\alpha 4\beta 2/3\delta$ ). (b) After SE, KCC2 expression decreases and the direction of chloride flux may change as a result. The expression of  $\alpha 1$  subunits decrease and  $\alpha 4$  subunits increase. Other changes are altered sensitivity to modulators. (2) One of the changes in the GABA<sub>A</sub>Rs in the DG after SE is loss of benzodiazepine sensitivity. (3) The pyramidal basket cell and its basket plexus appears to be similar after SE, although other GABAergic neurons are altered, and there may be changes in expression of various peptides. (4) The GABAergic synapse after SE has fewer  $\alpha 1$  subunits and increased  $\alpha 4$  subunits, which may become perisynaptic (indicated by a ?) (References are listed in the text. Parts 1–2 of this figure were adapted from Jacob et al. [59])

GCs, where they are proposed to affect both the rise time and decay of synaptic currents [71].

### 11.2.3 Regulation of [Cl]<sub>i</sub> in Early Development and Its Relevance to TLE

One of the characteristics of GABAergic inhibition at GABA<sub>A</sub>Rs that has implications for epilepsy – and has been studied extensively in the hippocampus in TLE – is the regulation of chloride flux through the GABA<sub>A</sub>R. The direction of chloride flux is regulated by many factors, and one source of regulation that has attracted a great deal of attention is the K<sup>+</sup>-Cl<sup>-</sup> cotransporters

KCC2 and NKCC1. KCC2 extrudes chloride normally, and NKCC1 transports chloride into the cell [7]. In early life, KCC2 expression is low and there is a relatively high concentration of intracellular chloride; chloride efflux occurs when GABA binds to the GABA<sub>A</sub>R, leading to a depolarization [8, 27]. After maturation, KCC2 expression increases and this leads to a lower [Cl]<sub>i</sub> and chloride influx when GABA binds to GABA<sub>A</sub>Rs, leading to a hyperpolarization [106]. As mentioned above, an exception is the GC, which has a resting potential (–70 to –80 mV) that is usually negative to E<sub>Cl<sup>-</sup></sub>. Therefore, in early life, a strong depolarization of GCs by GABA is predicted, and a smaller depolarization in adulthood compared to adulthood.

The idea that GABA is depolarizing in early postnatal life has recently been contested because most data that led to the idea were collected in slices where truncation of neuronal processes leads to elevated  $[Cl^-]_i$  [15]. However, *in vivo* studies have been conducted that are consistent with a depolarizing action of GABA in pyramidal neurons in neonatal life [9]. It remains to be determined exactly at what age these depolarizing effects end; in rodents it seems likely to be the first or second postnatal week [9, 15].

In the DG, one might expect that the switch from depolarizing to hyperpolarizing effects of GABA would not be as important because GABA typically has a depolarizing effect on GCs regardless. However, the size of the depolarization will be substantially greater if KCC2 expression is low, and moreover, there are many cells besides GCs in the DG that will be affected; only the GC has a very high resting potential. There are also many types of GABAergic inhibition, not only postsynaptic. If the GABA<sub>A</sub>R is presynaptic, for example, the net effect could very different if the terminal is depolarized or hyperpolarized by GABA.

There is also another process in the DG that is likely to be affected if the effects of GABA “switch” from depolarizing to hyperpolarizing – the maturation of GCs that are born postnatally, i.e., postnatal or “adult” neurogenesis [67]. GABA is a critical regulator of the maturation and migration of immature neurons in early life [24, 160]. GABA also influences maturation and migration of adult-born GCs [36]. In acquired TLE this is potentially important because animal models of TLE have shown that there is a large increase in proliferation of adult-born GCs after seizures [90], and the young GCs often mis-migrate (discussed further below). It has been suggested that these mis-migrated GCs contribute to chronic seizures (discussed further below).

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### 11.3 Alterations in GABAergic Transmission in Animal Models of TLE

There are many types of TLE, and one of the ways to classify the types is based on whether the epilepsy appears to have been “acquired.”

The term ‘acquired’ indicates that an insult or injury occurred prior to seizures and is likely to have caused the epilepsy. Acquired TLE has been simulated in laboratory animals by various insults or injuries that lead to a pattern of brain damage that is typical of TLE, called mesial temporal sclerosis (MTS; [127]). In general, MTS involves loss of a large number of CA1 and CA3 pyramidal cells, with sparing of CA2 and GCs. Many hilar neurons are lost, and these include both mossy cells and HIPP cells [116]. Notably, there are individuals with acquired TLE that do not have this classic description of MTS, and animal models vary in the extent they simulate MTS [127]. However, the pattern has been the focus of the most research in TLE, based on the assumption that this general pattern of neuropathology causes TLE or is very important to TLE.

One method that leads to a MTS-like pattern of neuropathology in adult rodents is induction of SE, either by injection of a chemoconvulsant such as kainic acid or pilocarpine, or electrical stimulation of hippocampus [31, 85, 95]. Here we will focus primarily on the SE models to study TLE in adult rodents, and use the data from SE models to address changes in GABAergic inhibition. We suggest that these changes involve plasticity of GABAergic mechanisms rather than simply an erosion or increase in the effects of GABA.

#### 11.3.1 Alterations in GABAergic Neurons After SE

Early observations that GABAergic neurons were decreased in neocortical epileptic foci produced by alumina gel in monkeys supported ideas that disinhibition may be the cause of epilepsy [100–102], particularly because the reduction in GABAergic neurons preceded epilepsy [56, 103]. Chandelier cells appeared to be one of the subtypes that was affected, and it was suggested that loss of the chandelier subtype of GABAergic neuron would be most likely to cause disinhibition of cortical pyramidal cells because loss of only a few axo-axonic cells would substantially change the number of GABAergic terminals at the axon hillock [33].

However, as more animal models were examined, there was less enthusiasm for the idea that disinhibition was the fundamental cause of seizures. In seizure-sensitive gerbils [93], the audiogenic seizure model [110], and kainic acid model [32], GABAergic neurons were not always decreased [54]. In fact, some GABAergic neurons increased their axon arbors, exhibiting axon sprouting (discussed further below). When GABA<sub>A</sub>R-mediated inhibition was examined, it was often strong rather than weak [11]. Therefore, even if some changes in these animal models involve disinhibition acutely, GABAergic neurons and GABA<sub>A</sub>R-dependent inhibition often show recovery and plasticity.

In the DG, an alternative hypothesis to disinhibition was suggested to address an animal model of TLE in which the perforant path of adult rats was stimulated electrically to simulate the precipitating insult in TLE. In this animal model, a 24 h period of intermittent perforant path stimulation in urethane-anesthetized rats led to a loss of 'paired-pulse' inhibition. Based on the results from these experiments, investigators suggested that the basket cells, (defined by parvalbumin expression) were spared but there was loss of HIPP cells (defined by somatostatin expression) and mossy cells [135]. Because mossy cells appeared to be decreased in numbers, and there were suggestions in the literature that they innervated basket cells, it was hypothesized that the parvalbumin-expressing basket cells lost afferent input from mossy cells and became 'dormant' and this led to disinhibition of GCs [136]. The hypothesis became known as 'the dormant basket cell hypothesis.' It was suggested that the hypothesis explained epileptogenesis in acquired TLE: if an early insult or injury led to loss of vulnerable mossy cells and HIPP cells, but GCs and basket cells were spared, the result would be disinhibition of GCs [6, 75].

However, later studies led to some doubt that this hypothesis could explain acquired TLE [12]. An alternative hypothesis – the 'irritable mossy cell hypothesis' – suggested that mossy cells could cause GC hyperexcitability because the mossy cells, which project directly to GCs, developed increased excitability. This hypothesis was

developed on the basis of recordings from mossy cells in slices after post-traumatic injury [113, 114], another type of precipitating insult that leads to TLE. In addition, mossy cell hyperexcitability was shown subsequently in slices from epileptic rats after SE [128].

A result that argued against these two hypotheses came from studies of animals with chronic epilepsy after kainic acid-induced SE. These experiments showed that there was an increase in paired-pulse inhibition of GCs, not a decrease [139]. In addition, slices from animals after SE did not exhibit spontaneous seizure-like activity, suggesting they had intact inhibition rather than weak inhibition. This was unlikely to be due to the differences in the SE model since 'irritable mossy cells' were observed, at least in one study of SE [128]. In slices, exposure of slices to GABA<sub>A</sub>R antagonists led to seizure-like activity that was more prolonged in slices from animals that had SE than slices from control rats. From these experiments, it was suggested that slices from animals with SE were hyperexcitable but it was normally masked by GABA<sub>A</sub>R-mediated inhibition [129, 147]. In slices from humans with intractable TLE, there was enhanced sensitivity to bicuculline [39]. These observations and others led to the idea that increased inhibition was present to compensate for underlying hyperexcitability [147, 162]. Although in some cases the studies of animals with SE and intractable TLE reflect differences in the models or the subtypes of TLE, here the data from different models and humans was consistent, making the observations compelling.

Although an attractive idea, GABAergic inhibition in the animal models of SE does not necessarily seem to be too strong, masking underlying hyperexcitability. For example, interneurons exhibit axonal sprouting in the DG in animal models of TLE [5, 32, 151]. It is not clear that they simply extend their output, inhibiting more glutamatergic neurons than normal, because they innervate inhibitory neurons as well [137]. Interneurons develop abnormal glutamatergic input from sprouting of the GCs into the inner molecular layer (mossy fiber sprouting; for review see [19]). The evidence for this is based on

staining of the mossy fibers with Timm stain [137]. Electron microscopy of the mossy fiber boutons in the inner molecular layer supported the idea that the sprouted mossy fibers activate GABAergic basket cells [43]. In further support of this idea, it was suggested that normal mossy fibers in the hilus and area CA3 primarily innervate GABAergic neurons and primarily have an inhibitory effect on CA3 [1]. Moreover, GCs express GABA as well as glutamate after SE [50] and GABA release from GCs can be inhibitory [158] although the latest studies suggest this may be limited to GCs at an early stage of development [23]. The vast majority of studies show that GCs in normal hippocampus excite their target cells [60, 122, 156]. In addition, when mossy fiber synapses in the epileptic rat were quantified in the inner molecular layer, the majority were located on GCs, not interneurons [19, 20].

One way to reconcile the different data is to suggest that mossy fibers have a large dynamic range, with filopodia that excite interneurons and massive boutons that excite principal cells. The outcome may depend on recent activity, which can potentially upregulate GABA expression, or alter the peptide content of the massive boutons so that they are more excitatory [123]. Other hypotheses suggest that mossy fibers can be inhibitory to area CA3 pyramidal cells depending on the firing mode of GCs – after bursts of GC action potentials, excitation of pyramidal cells is transiently suppressed [82].

As our experimental techniques improved, our understanding of the underlying changes became clearer. For example, initial assays to assess inhibition measured paired-pulse inhibition which uses extracellular recordings and is not an extremely reliable measurement, because small changes in the stimulating or recording sites can alter the extent of inhibition even in the same preparation [157]. As patch clamp recordings developed, more indices of pre- and postsynaptic GABAergic inhibition became possible, and the results have shown that the GABAergic system in the DG is changed in diverse ways after SE, not always consistent with disinhibition of GCs, and not always consistent with hyperinhibition (Fig. 11.1b, c).

### 11.3.2 Alterations in GABA Receptors in GCs After SE

During SE, inhibitory GABAergic synaptic transmission in the DG becomes compromised, presumably due to the dramatic increase in activation of GABAergic neurons. Miniature inhibitory post-synaptic currents (mIPSCs) are reduced in GCs and the number of active GABA<sub>A</sub>Rs per GC decreases [26, 47, 86] via enhanced clathrin-dependent GABA<sub>A</sub>R internalization [48, 59]. In vitro studies using hippocampal neurons, stimulated with a buffer containing low magnesium to induce spontaneous recurrent epileptiform discharges, showed a large decrease in GABA-gated chloride currents that correlated with reduced cell surface expression and intracellular accumulation of GABA<sub>A</sub>Rs [13, 48]. In vivo studies using chemoconvulsants have shown that SE promotes a rapid reduction in the number of physiologically active GABA<sub>A</sub>Rs in GCs that correlated with a reduction in the level of  $\beta 2/\beta 3$  and  $\gamma 2$  immunoreactivity present in the vicinity of a pre-synaptic marker [86]. In fact, SE appears to trigger subunit specific events to regulate the trafficking of GABA<sub>A</sub>Rs by promoting the dephosphorylation of  $\beta 3$  subunits [47, 150]. Decreased phosphorylation of  $\beta 3$  increases the interaction of GABA<sub>A</sub>Rs with the clathrin-adaptor protein 2 (AP2), facilitating the recruitment of GABA<sub>A</sub>Rs into clathrin-coated pits and promoting their removal from the plasma membrane [47, 150]. In hippocampal slices obtained from mice after SE, increased GABA<sub>A</sub>R phosphorylation or blockade of normal AP2 function resulted in GABA<sub>A</sub>R accumulation at the plasma membrane and increased synaptic inhibition [150].

Alterations in GABA<sub>A</sub>R subunit composition occur subsequent to SE in a number of animal models, and there is evidence that these changes may contribute to epileptogenesis [18, 72, 76, 92, 144, 166]. SE results in changes in the expression and membrane localization (i.e., extrasynaptic vs. synaptic) of several GABA<sub>A</sub>R subunits (e.g.,  $\alpha 1$ ,  $\alpha 4$ ,  $\gamma 2$ , and  $\delta$ ) in GCs. Beginning soon after SE and continuing until and after the animals become epileptic, these alterations are associated with changes in phasic and tonic GABA<sub>A</sub>R-

mediated inhibition, and in GABA<sub>A</sub>R pharmacology [21, 30, 45]. After pilocarpine-induced SE, GABA<sub>A</sub>R  $\alpha 1$  subunit mRNA expression decreases, and GABA<sub>A</sub>R  $\alpha 4$  subunit mRNA expression increases [18]. Changes in GABA<sub>A</sub>R function and subunit expression have also been observed in neurons from surgically resected hippocampus of patients with intractable TLE; [17, 143]. These alterations are associated with an increase in  $\alpha 4\gamma 2$  containing receptors, a reduction in  $\alpha 1\gamma 2$  containing receptors in the DG [76], and shift of  $\alpha 4$ -containing receptors from extrasynaptic to synaptic and perisynaptic locations, which is likely to be related to the appearance of  $\alpha 4\beta\gamma 2$  receptors [146, 166]. Changes in expression and localization of  $\alpha$ -subunits associated with changes in synaptic GABA<sub>A</sub>R composition result in a number of changes in synaptic inhibition in GCs, including diminished benzodiazepine sensitivity, enhanced zinc sensitivity, reduced neurosteroid modulation, and diminished phasic inhibition in dendrites [21, 30, 45, 146]. Preventing the reduction in GABA<sub>A</sub>R subunit  $\alpha 1$  expression after SE via viral-mediated transfer of an  $\alpha 1$  subunit transgene in adult rodents reduced subsequent epilepsy development, resulting in a three-fold increase in the mean time to the first spontaneous seizure, and a decrease to 39 % of AAV- $\alpha 1$ -injected rats developing spontaneous seizures in the first 4 weeks after SE compared to 100 % of rats receiving sham injections [99]. Together, these data support a role for GABA<sub>A</sub>R  $\alpha$ -subunit changes in the process of epileptogenesis.

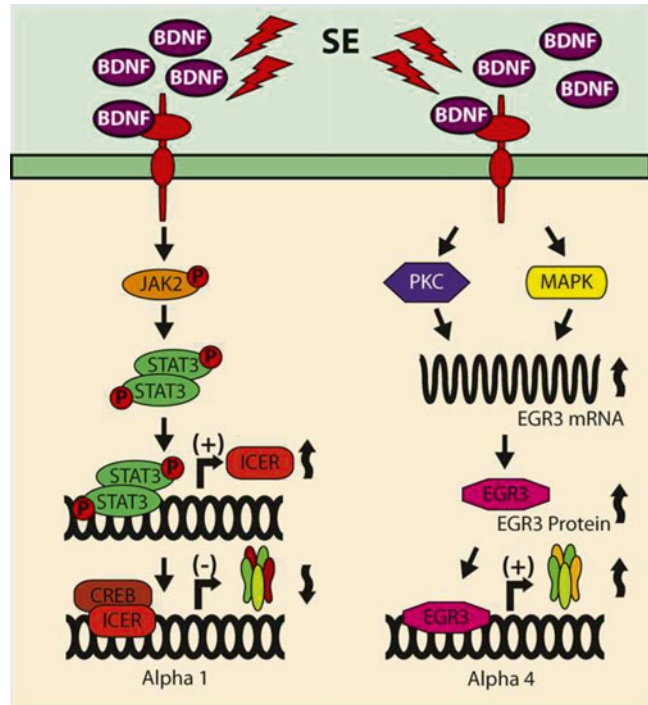
Receptors containing  $\alpha 4$  subunits are most often found with the  $\delta$  rather than the  $\gamma$  subunit in combination with  $\alpha\beta$ . These  $\alpha 4\beta\delta$  GABA<sub>A</sub>Rs are localized to extrasynaptic sites and contribute to tonic inhibition. Under physiological conditions, only a minor population of  $\alpha 4\beta\gamma 2$  GABA<sub>A</sub>Rs are found within GABAergic synapses on GCs, where they are proposed to affect both the rise time and decay of synaptic currents [71]. In parallel with the decrease in  $\alpha 1$  subunit expression in GCs after SE, there is a marked increase in  $\alpha 4$  subunit expression that results in an increase in the abundance of  $\alpha 4\gamma 2$ -containing receptors in synaptic and perisynaptic locations [146, 166] (see Fig. 11.2), along with the reduction in

$\alpha 1\gamma 2$ -containing receptors [76]. The  $\alpha 4\beta\gamma 2$  receptors may contribute to epileptogenesis, as  $\alpha 4$ -containing GABA<sub>A</sub>Rs have been shown to desensitize rapidly, especially when assembled with  $\beta 3$  subunits [71]. In addition, GABA<sub>A</sub>Rs containing the  $\alpha 4$  subunit are very sensitive to zinc blockade, as are GABA<sub>A</sub>Rs on GCs in the epileptic brain [21, 30]. Zinc containing mossy fiber terminals sprout from the granule cell layer of the hippocampus onto other GCs and into CA3, likely depositing zinc onto the newly formed  $\alpha 4\beta\gamma 2$  receptors causing a decreased response to GABA. Collectively these alterations may contribute to epilepsy development, pharmacoresistance and further epilepsy progression.

GABA<sub>A</sub>R subunit alterations after SE are regulated by increased synthesis of brain-derived neurotrophic factor (BDNF) and activation of its receptors (TrkB and p75) that control a number of down-stream pathways, including Janus kinase (JAK)/Signal Transducer and Activators of Transcription (STAT), protein kinase C, and mitogen activated protein kinase (MAPK; [76, 107, 108]). BDNF is known to enhance cAMP response element binding protein (CREB) phosphorylation through binding to TrkB receptors [105, 163], and is also a potent regulator of inducible cAMP response element repressor (ICER) synthesis [57]. Using chromatin immunoprecipitation (ChIP) and DNA pulldown studies, it has been determined that there is increased binding of pCREB and ICER to the GABA<sub>A</sub>R $\alpha 1$  gene promoter (*GABRA1-p*) in DG after SE [76]. BDNF regulation of ICER expression is mediated by JAK/STAT pathway activation, specifically activation of pJAK2 and pSTAT3 [76]. pSTAT3 association with the STAT-recognition site on the ICER promoter is enhanced after SE in DG and inhibition of JAK/STAT signaling pathway with pyridone 6 (P6) in primary hippocampal cultures and *in vivo* in DG prior to SE blocks both ICER induction and decreased transcription of *GABRA1* [76]. These findings suggest a specific signaling cascade involving BDNF, JAK/STAT, and CREB that is critical to the reported decreases in  $\alpha 1$  subunit levels following SE and may contribute to epileptogenesis. Increases in GABA<sub>A</sub>R $\alpha 4$  subunit are transcriptionally regulated by BDNF activation of the TrkB

**Fig. 11.3 Regulation of GABA<sub>A</sub> receptor expression after SE.**

BDNF regulates the final composition of GABA<sub>A</sub>Rs by differentially altering the expression of  $\alpha 1$  and  $\alpha 4$  subunits. Both in vivo and in vitro evidence suggest that increased levels of BDNF following SE activate at least two different signaling pathways: JAK/STAT and PKC/MAPK, resulting in the down-regulation of  $\alpha 1$  subunits and the up-regulation of  $\alpha 4$  subunits, respectively (Reproduced from Gonzalez and Brooks-Kayal [46])



receptor which leads to upregulation of the early growth response factor (Egr3) pathway via a PKC/MAPK-dependent pathway [107]. Egr3 association with the early-growth response-recognition (ERE) site on the *GABRA4* promoter is enhanced after SE in DG [107] (See Fig. 11.3).

### 11.3.3 Regulation of GABA in Early Development and Its Relevance to TLE

One of the themes in studies of animal models of TLE is the idea that the myriad of changes in hippocampal structure and function that have been described are associated with a recapitulation of development that is caused by the epileptogenic insult. A robust example is the dramatic increase in the rate of adult neurogenesis in the DG after epileptogenic insults like SE. First noted by Bengzon et al [10] using stimulus-evoked afterdischarges, and Parent et al. [90] after pilocarpine-induced SE, the increase in the rate of adult neurogenesis after seizures, and particularly SE (in adult rodents), has been reproduced by many laboratories in response to virtually all

epileptogenic insults: kindling, kainic acid or electrically-induced SE, or traumatic brain injury [121, 124].

Initially it was suggested that many of the neurons that are born after SE do not survive long-term [90] which has also been shown by others [96] but a substantial fraction of newborn neurons can survive in some animal models, and these migrate into the hilar region, where they are called hilar ectopic GCs (hEGCs; [119]). Other adult-born GCs migrate correctly but develop abnormal dendrites in the hilus, called hilar basal dendrites [104, 133]. These neurons also appear to survive long-term and can be generated for a long-time after SE [62]. Another subset of GC that develops after SE and is abnormal develops an enlarged cell body (hypertrophy; [98]). The abnormal GCs are potentially important because they contribute to mossy fiber sprouting, particularly hEGCs [69, 94, 119]. HEGCs participate in seizures in vivo [130] and their numbers are correlated with chronic seizure frequency [79]. Manipulations that reduce hEGC number reduce chronic seizure frequency after SE [63], although selective deletion of hEGCs is not yet possible. The hEGCs display a variety of

electrophysiological characteristics [61, 118, 164, 165] which are unlike normal GCs. For these reasons, the neurons that hypertrophy, and the hEGCs, have been suggested to contribute to seizure generation [63, 68, 98, 117, 119].

The plasticity of GABAergic mechanisms in animal models of TLE plays a potentially important role in the development of abnormal GCs, and therefore the role these GCs play in seizure generation. In a study that used experimental febrile seizures to induce epilepsy later in life, febrile seizures caused mismigration of immature GCs into the hilus by changing the normal regulation of migration by GABA acting at GABA<sub>A</sub>Rs. This study was important in showing that altering the normal effect of GABA by febrile seizures could cause aberrant circuitry that would persist long-term, potentially contributing to seizure generation. Interestingly, the way that GABA was altered was in the expression of GABA<sub>A</sub>Rs; more GABA<sub>A</sub>Rs were found by western blot after febrile seizures. In response to increased depolarization by GABA, immature GCs migrated opposite to their normal direction, into the hilus instead of the GC layer. Knockdown of NKCC1 could block the formation of hEGCs and reduce the long-term effects [68]. The studies of Koyama and colleagues and Swijsen et al. [149], who also studied febrile seizures, both found increased  $\beta$ 2/3 subunits occurred in newborn GCs after febrile seizures [149]. Changes in  $\alpha$ 3 subunits were also noted by Swijsen et al. [148]. The results suggest that febrile seizures lead to long-lasting changes in the expression of GABA<sub>A</sub>Rs in the DG, and in the GCs that were born after febrile seizures. These effects could lead to life-long reduction in limbic seizure threshold. They also may contribute to the comorbidities in TLE, such as depression [16, 66], a psychiatric condition where adult neurogenesis in the DG has been shown to play a critical role [112].

Another study of adult rodents is also relevant to the formation of aberrant GCs in TLE. This study used pilocarpine-induced SE in adult rodents to ask how KCC2 is altered immediately after SE. The investigators showed that there was a downregulation of KCC2 in the DG after

SE which would make GCs (both mature and immature GCs) depolarize more in response to GABA [91]. If the results of Koyama et al. [68] are correct, greater depolarization by GABA would be likely to foster mismigration of immature GCs. A similar phenomenon may explain why newborn neurons after SE, in the adult, mis-migrate for long distances—it has been described that they migrate from the subgranular zone to the border of the hilus and area CA3 [118]. Together the new information about  $[Cl^-]_i$  regulation are providing potential mechanisms underlying acquired epileptogenesis in the immature and mature brain. Although a great deal more information will be necessary before new treatments can be developed based on the new hypotheses, NKCC1 antagonists are already in clinical trial [64, 65].

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## 11.4 Summary

In the DG, the robust plasticity of GCs has been of avid interest because they upregulate numerous proteins and exhibit robust sprouting of their axons after seizures. Although extensive studies of GABA in the DG have been made in TLE, the remarkable plasticity of GABAergic mechanisms is often not considered as much as development of disinhibition or hyperinhibition. Here we suggest that there are numerous pre- and postsynaptic changes in GABAergic transmission, even if one only addresses GABAergic synapses on GCs and GABA<sub>A</sub> receptors. Taken together, this plasticity leads to more complexity of GABAergic transmission in the epileptic brain, not simply an increase or decrease. The idea that GABAergic inhibition is dramatically altered, rather than increased or decreased, is consistent with the diversity of results of past studies. Therefore, this perspective helps address some of the conflicts in the past. It also provides a different and potentially more accurate perspective that will facilitate antiseizure drug development.

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# Do Structural Changes in GABA Neurons Give Rise to the Epileptic State?

12

Carolyn R. Houser

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## Abstract

Identifying the role of GABA neurons in the development of an epileptic state has been particularly difficult in acquired epilepsy, in part because of the multiple changes that occur in such conditions. Although once questioned, there is now considerable evidence for loss of GABA neurons in multiple brain regions in models of acquired epilepsy. This loss can affect several cell types, including both somatostatin- and parvalbumin-expressing interneurons, and the cell type that is most severely affected can vary among brain regions and models. Because of the diversity of GABA neurons in the hippocampus and cerebral cortex, resulting functional deficits are unlikely to be compensated fully by remaining GABA neurons of other subtypes. The fundamental importance of GABA neuron loss in epilepsy is supported by findings in genetic mouse models in which GABA neurons appear to be decreased relatively selectively, and increased seizure susceptibility and spontaneous seizures develop. Alterations in remaining GABA neurons also occur in acquired epilepsy. These include alterations in inputs or receptors that could impair function, as well as morphological reorganization of GABAergic axons and their synaptic connections. Such axonal sprouting could be compensatory if normal circuits are reestablished, but the creation of aberrant circuitry could contribute to an epileptic condition. The functional effects of GABA neuron alterations

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thus may include not only reductions in GABAergic inhibition but also excessive neuronal synchrony and, potentially, depolarizing GABAergic influences. The combination of GABA neuron loss and alterations in remaining GABA neurons provides likely, though still unproven, substrates for the epileptic state.

### Keywords

Inhibition • Plasticity • Seizures • Sprouting • Somatostatin • Parvalbumin

## Abbreviations

CCK	Cholecystokinin
eGFP	Enhanced green fluorescent protein
eYFP	Enhanced yellow fluorescent protein
GABA	Gamma aminobutyric acid
GAD	Glutamic acid decarboxylase
NPY	Neuropeptide Y
PV	Parvalbumin
SOM	Somatostatin
s. oriens	Stratum oriens
TLE	Temporal lobe epilepsy
<i>uPAR</i>	Urokinase plasminogen activator receptor

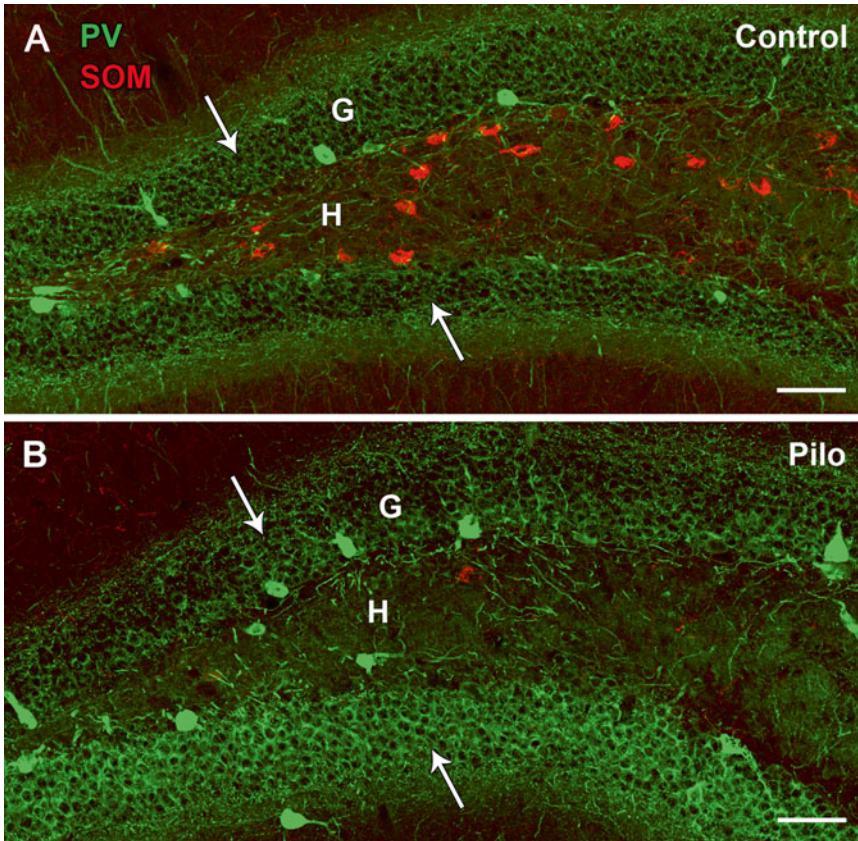
## 12.1 Introduction

The question of whether GABA neuron loss gives rise to the epileptic state in acquired epilepsy has persisted for many years. Indeed, even the occurrence of GABA neuron loss was questioned at one time. While progress has been made, and a loss of GABA neurons has been convincingly identified in humans with temporal lobe epilepsy (TLE) and in many related animal models, determining the functional consequences of GABA neuron loss in epilepsy remains a major challenge. This review will focus on some of the complexities associated with interneuron loss and their role in epilepsy and suggest that GABA neuron loss could indeed give rise to the epileptic state through both direct and indirect routes.

## 12.2 Loss of GABA Neurons Is a Consistent Finding in Models of Acquired Epilepsy

Interneuron loss is one of the most frequently observed alterations in models of TLE, and the consistency of GABA neuron loss provides a solid base for suggesting the potential importance of this alteration in creating an epileptic state. Although loss of GABA neurons has been identified in multiple brain regions, loss of somatostatin (SOM)-expressing GABA neurons in the hilus of the dentate gyrus remains one of the clearest and most consistent findings (Fig. 12.1a, b). Loss of these GABA neurons has been found in virtually all models of acquired epilepsy, including kindling, status epilepticus and traumatic brain injury models [7, 25, 32, 44, 45]. Importantly, loss of SOM/GABA neurons in the dentate hilus is also found in human TLE, as part of the broader loss of neurons in typical hippocampal sclerosis, as well as in pathological conditions with more limited cell loss such as end-foolium sclerosis [14, 35, 42, 48, 49].

SOM neurons in stratum oriens (s. oriens) of CA1 are also among the vulnerable interneurons in several models of acquired epilepsy [1, 11, 27, 36]. As SOM neurons in both the hilus and s. oriens provide GABAergic innervation of dendrites of granule cells and pyramidal cells, respectively, they are ideally positioned to control excitability of the principal cells directly at the sites of their major excitatory inputs. This pattern of GABA neuron loss has led to the suggestion that



**Fig. 12.1** Comparisons of somatostatin (SOM)- and parvalbumin (PV)-labeled neurons in the rostral dentate gyrus of control (a) and pilocarpine (Pilo)-treated (b) mice. (a) In the control dentate gyrus, cell bodies of SOM neurons are located primarily within the hilus (H) whereas those of PV neurons are positioned predominantly along the base of the granule cell layer (G). PV-labeled axon terminals are concentrated in

perisomatic locations within the granule cell layer (*arrows*) while SOM terminal fields are located in dendritic regions in the outer molecular layer (not shown). (b) In the pilocarpine-treated mouse at 2 months after status epilepticus, a severe loss of SOM neurons is evident in the hilus whereas many PV neurons and their axon terminals (*arrows*) in the granule cell layer are preserved. Scale bars, 100  $\mu\text{m}$

interneurons that provide dendritic innervation are more vulnerable to damage in epilepsy than those which provide primarily perisomatic innervation, such as basket cells and axo-axonic cells, many of which express the calcium binding protein parvalbumin (PV). Electrophysiological findings of decreased dendritic inhibition, with preservation of somatic inhibition, in pyramidal cells of CA1 in models of recurrent seizures support this idea [11].

While the distinction between dendritic and perisomatic innervation provides a useful framework for considering GABA neuron loss, the differences in vulnerability among the broad types

of interneurons are not clear-cut, and additional complexities exist. While PV-expressing neurons are a major source of perisomatic innervation of pyramidal cells in the hippocampus, cholecystokinin (CCK)-expressing interneurons also provide perisomatic innervation of these neurons in CA1 [3, 23]. In a mouse pilocarpine model of recurrent seizures, this CCK innervation appeared to be decreased while the PV innervation was preserved [51]. This could create an imbalance in perisomatic control that could favor synchronizing actions of PV basket cells, with loss of major modulatory inputs from CCK neurons.



Perisomatic innervation includes both basket cells and axo-axonic cells [22], and these cell types could be affected differentially in epilepsy. Decreased innervation of the axon initial segment by PV-expressing axo-axonic neurons has been found in the hippocampus and cerebral cortex in several epilepsy conditions, suggesting that axo-axonic cells could be more vulnerable than basket cells [13, 15, 41].

Thus loss of PV neurons can occur and has been described in the dentate gyrus in several animal models, without distinctions between basket cells and axo-axonic cells [1, 25, 29]. However, when both PV and SOM neurons have been studied in the same animals, the loss of PV-expressing neurons in the dentate gyrus is generally less severe than that of SOM interneurons [7] (Fig. 12.1a, b).

A decrease in numbers of PV-expressing interneurons has now been identified in several other regions of the hippocampal formation where their loss could be particularly important in regulating activity within the broader hippocampal circuit. A significant decrease in PV neurons has been identified in layer II of the entorhinal cortex where loss of these neurons could contribute to increased excitability of the perforant path input to the dentate gyrus [30]. Interestingly, lower densities of PV-containing neurons have also been identified in the subiculum, where their loss could lead to increased excitability of this major output region of the hippocampal formation, a region that is otherwise generally well preserved [2, 16].

Thus decreases in both SOM and PV neurons can occur in epilepsy, and the particular pattern of loss may vary among epilepsy models, species, brain regions and even rostral-caudal levels of the hippocampal formation. The types of GABA neurons that are affected remain important as they will determine the specific functional effects. However, loss of GABA neurons remains a unifying theme.

Despite strong evidence for GABA neuron loss in many brain regions, direct relationships to the epileptic state have been difficult to demonstrate. This could be in part because GABA neuron

loss does not occur in isolation in most forms of acquired or lesion-induced epilepsy. In human TLE and related epilepsy models, extensive cell loss can occur in many regions, including CA1 and CA3 as well as the dentate hilus and extra-hippocampal regions. This neuronal loss generally involves both principal cells and interneurons, making it more difficult to link GABA neuron loss directly to development of an epileptic state. However, findings in several genetic mouse models provide support for the importance of GABA neuron loss in the development of epilepsy, and these findings also have relevance for acquired epilepsy.

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### 12.3 Selective Loss of GABA Neurons Can Lead to an Epileptic State in Genetic Models

Some of the strongest evidence for loss of GABA neurons giving rise to the epileptic state has come from genetically-modified mice in which GABA neurons are selectively affected, and increased seizure susceptibility and spontaneous seizures occur. In mice with loss of the *Dlx1* gene, a transcription factor that regulates development of GABAergic interneurons originating in the medial ganglionic eminence, there is a time-dependent reduction in the number of interneurons in the cerebral cortex and hippocampus and development of an epilepsy phenotype [10]. SOM and calretinin-expressing neurons were reduced in number whereas PV-expressing neurons appeared to be unaffected. Because the loss of GABA neurons is apparently selective in these mice, the findings provide strong support for loss of GABA neurons giving rise to an epileptic state and also suggest that the loss of GABA neurons does not need to be extensive. In these mice, behavioral seizures were selectively induced by mild stressors by 2 months of age when there was an approximately 22 % reduction in GAD67-labeled neurons in the cerebral cortex and 24 and 29 % reduction in the dentate gyrus and CA1 respectively. Comparable or even greater GABA neuron loss

has been observed in the hippocampal formation in models of acquired epilepsy [37, 50].

Similarly, in mice with mutation of the gene encoding urokinase plasminogen activator receptor (*uPAR*), a key component in hepatocyte growth factor activation and function, interneuron migration is altered, and the mice have a nearly complete loss of PV neurons in the anterior cingulate and parietal cortex [40]. These mice also developed spontaneous myoclonic seizures and increased susceptibility to pharmacologically-induced convulsions. Thus the apparently selective loss of either of two major groups of interneurons supports the importance of GABA neuron loss in the development of epilepsy.

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## 12.4 Remaining GABA Neurons Could Play a Critical Role in Development of Epilepsy

Despite clear evidence for loss of GABA neurons in virtually all models of acquired epilepsy and human TLE, some GABA neurons invariably remain, and alterations in these neurons could contribute to the creation of an epileptic condition. Indeed, it may be difficult to separate the effects of loss of GABA neurons from altered function of remaining neurons as the initial loss of GABA neurons may be a stimulus for the subsequent changes in remaining GABA neurons. Critical changes may include impaired function of remaining GABA neurons and morphological reorganization of remaining interneurons that could lead to altered or aberrant circuitry.

### 12.4.1 Impaired Function of Remaining Interneurons

Remaining GABA neurons often appear particularly prominent in tissue from animal models of epilepsy, and the preservation of some GABA neurons has suggested that GABA neuron loss may be of limited importance in establishing the epileptic state. However, the function of remaining GABA neurons could be altered, leading to

inadequate control of principal cell activity. In the dentate gyrus and hippocampus, the functional state of remaining basket cells has been debated for many years. Specific details of the “dormant basket cell” hypothesis [46, 47] have been questioned, including the role of hilar mossy cell loss in reducing basket cell activity [5, 18]. However, the broad suggestion that basket cells and other GABAergic neurons might be functioning sub-optimally due to decreased or impaired excitatory input remains plausible. Recent studies have identified deficits in basket cell function in the dentate gyrus that could indicate a decrease in excitatory afferent input or reduction of the readily releasable pool of synaptic vesicles, in association with an increased failure rate at basket cell to granule cell synapses [53].

Alterations in the receptors and channels of remaining GABA neurons also could reduce the activity of these neurons. In both the rat and mouse pilocarpine model, expression of the  $\delta$  subunit of the GABA<sub>A</sub> receptor is increased in subgroups of GABA neurons in the dentate gyrus [38, 52]. As GABA<sub>A</sub> receptors expressing the  $\delta$  subunit are responsible for the majority of tonic inhibition in these neurons [24], an increase in  $\delta$  subunit expression in interneurons could reduce their excitability and impair inhibitory control of the network [38]. Recent studies have demonstrated that tonic inhibition is indeed enhanced in fast-spiking basket cells of the dentate gyrus at 1 week after pilocarpine-induced status epilepticus [52]. However, additional changes, including decreased KCC2 expression in the basket cells, appeared to compensate partially for the increased tonic inhibition of the basket cells, and dentate excitability was not increased. Nevertheless, simulation studies suggested that the changes in tonic inhibition, in combination with other recognized alterations in dentate gyrus circuitry in epilepsy models, could lead to increased granule cell firing and self-sustained seizure-like activity in a subset of simulated networks [52]. Thus occasional alterations in interneuron activity, when combined with other changes in the network, may be sufficient to overrule compensatory changes and lead to sporadic seizure activity.

While regulation of  $\delta$  subunit-containing GABA<sub>A</sub> receptors by neurosteroids and other endogenous modulators could play important roles [21], changes in numerous other channels and receptors in remaining GABAergic interneurons could reduce their effectiveness and contribute to the epileptic state.

Functional alterations in interneurons and their relationship to seizure activity are demonstrated convincingly in genetic mouse models in which specific channels have been deleted relatively selectively in interneurons. As a key example, loss of the alpha subunit of the Na<sub>v</sub>1.1 sodium channel, that is encoded by the *SCN1A* gene, impairs sodium currents more severely in GABAergic neurons than in pyramidal cells [8, 17, 34]. Such changes limit the ability of the inhibitory interneurons, including PV neurons, to fire action potentials at high frequency, and the animals develop spontaneous generalized seizures.

Similarly, loss of function of the Ca<sub>v</sub>2.1 voltage-gated Ca<sup>2+</sup> channel reduces GABA release from cortical PV neurons, and generalized seizures occur in mice with such loss [43]. While decreased expression of this calcium channel was found in both PV and SOM neurons, only the loss in fast spiking, presumably PV, interneurons led to spontaneous seizures. Compensation by N-type Ca<sup>2+</sup> channels appeared to maintain function of the SOM interneurons but was insufficient for adequate function of the PV neurons.

Finally, elimination of the voltage-gated potassium channels of the Kv3 subfamily, that are particularly prominent in fast-spiking interneurons in the deep layers of the neocortex, led to an inability of these interneurons to fire at their normal high frequency and an increased susceptibility to seizures [31].

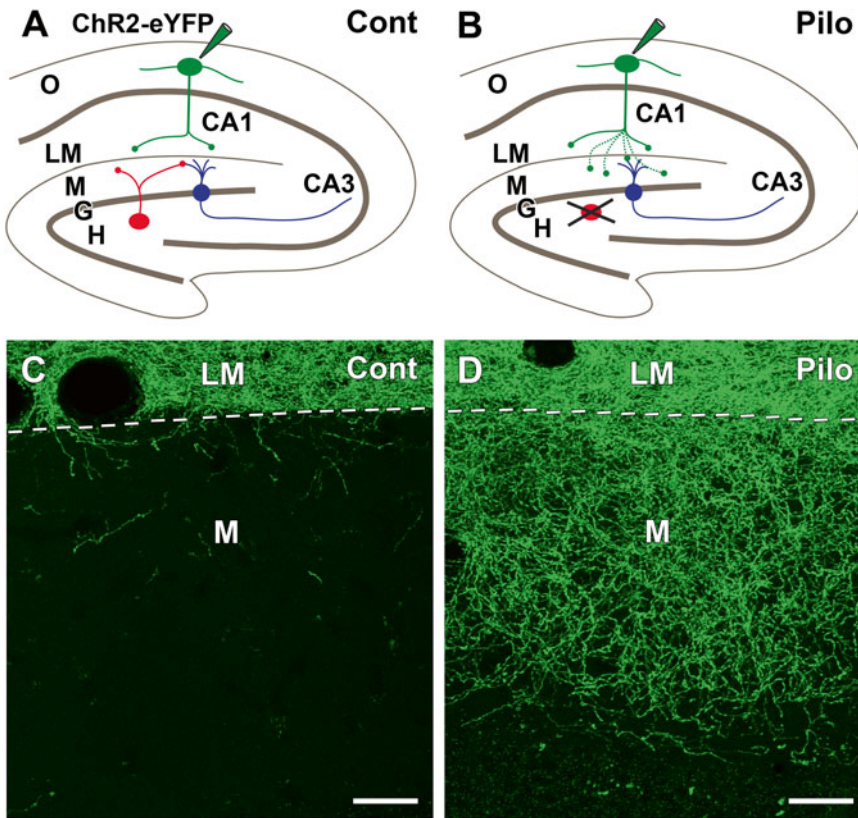
Thus in several genetic models, impairment of fast-spiking PV neurons, particularly a reduction in their ability to fire action potentials at high frequency, can lead to increased seizure susceptibility. Although these functional deficits are induced by specific genetic modifications, similar alterations in remaining GABA neurons may occur in acquired epilepsy, and even small functional impairment in remaining neurons could tip the balance toward seizure activity.

## 12.4.2 Morphological Reorganization of Remaining Interneurons

Clear demonstrations of loss of GABA neurons in acquired epilepsy have often been obscured by the plasticity of remaining interneurons. Remaining GABA neurons frequently express increased levels of GABA neuron markers, including the mRNA and protein of two isoforms of the GABA synthesizing enzyme, glutamic acid decarboxylase 65 and 67 (GAD65 and GAD67), as well as GABA [9, 19, 20]. Similarly the expression of peptides such as SOM and neuropeptide Y (NPY) within specific subclasses of GABA neurons are frequently upregulated [6, 33, 44]. These changes can be substantial and, during the chronic period, labeling of remaining GABA neurons can be quite strong and can suggest that either little loss of GABA neurons has occurred or that axons of remaining GABA neurons have sprouted [12]. It has remained particularly difficult to distinguish morphological growth and reorganization of GABAergic axons from an increase in GABAergic markers within remaining neurons [4, 27].

Additional support for sprouting of existing SOM neurons in the dentate gyrus has been obtained from mice that express enhanced green fluorescent protein (eGFP) in a subgroup of SOM neurons [54]. By studying the labeled interneurons in pilocarpine-treated mice, Buckmaster and colleagues demonstrated that SOM neurons that survive in the ventral (caudal) dentate gyrus can re-innervate the outer half of the dentate gyrus that was partially deafferented by loss of hilar SOM neurons. Such reorganization has generally been presumed to be compensatory as remaining GABA neurons are replacing the innervation of neurons of a similar type and function [26, 54].

Axonal reorganization of remaining GABA neurons can also create aberrant GABAergic circuitry as has been observed in the rostral dentate gyrus in the pilocarpine mouse model [39]. Apparent reinnervation of the dentate molecular layer was observed during the chronic period, but, in contrast to the previous study, few remaining SOM neurons were found in the rostral hilus. To determine if the innervation could be derived



**Fig. 12.2** Axonal reorganization of remaining somatostatin (SOM) neurons in pilocarpine (Pilo)-treated mice at 2 months after status epilepticus, illustrated schematically in (a, b) and in confocal images in (c, d). (a) This schematic illustrates the normal circuitry of SOM neurons in the hilus (red) and s. oriens (green) and the labeling protocol. In a control SOM-Cre mouse, selective labeling of SOM neurons in s. oriens (O) of CA1, by Cre-dependent AAV transfection of Chr2-eYFP, leads to labeling of their axon terminals that are confined to s. lacunosum-moleculare (LM). SOM neurons (red) in the hilus (H) innervate the outer molecular layer (M) of the dentate gyrus where they form synapses with dentate granule cells (G, blue). These hilar SOM neurons are not labeled by the injection in s. oriens. (b) In pilocarpine-treated mice,

similar labeling of SOM neurons in s. oriens leads to axonal labeling not only in s. lacunosum-moleculare of CA1 but also in the molecular layer of the dentate gyrus, a region that was previously innervated by vulnerable SOM neurons (red) in the hilus. (c) In a control SOM-Cre mouse, eYFP-labeled axons are concentrated in s. lacunosum-moleculare (LM), and only a limited number of labeled fibers cross the hippocampal fissure (dashed line) to enter the molecular layer (M) of the dentate gyrus. (d) In a similarly transfected pilocarpine-treated mouse, numerous labeled fibers cross the hippocampal fissure and form an extensive plexus in the outer two-thirds of the dentate molecular layer, where they innervate dentate granule cells. Scale bars, 20  $\mu\text{m}$  (Adapted from data in Peng et al. [39])

from other sources, SOM neurons in s. oriens of control and pilocarpine-treated SOM-Cre recombinase mice were selectively labeled with a viral vector containing Cre-dependent channel-rhodopsin2 (Chr2) fused to enhanced yellow fluorescent protein (eYFP). In control mice, the axons of many labeled SOM neurons in s. oriens formed a dense plexus of fibers in s. lacunosum-moleculare of CA1 (Fig. 12.2a, c). This plexus

was sharply delineated by the hippocampal fissure, and relatively few fibers crossed the fissure to enter the adjacent molecular layer of the dentate gyrus (Fig. 12.2c). In contrast, in pilocarpine-treated mice, an extensive axonal plexus of eYFP-labeled fibers was evident in the outer two-thirds of the dentate gyrus during the chronic period (Fig. 12.2b, d). Thus SOM neurons in s. oriens exhibited an unexpected capacity for

morphological growth and reorganization, and created an aberrant circuit between hippocampal interneurons in s. oriens and granule cells of the dentate gyrus. The reorganized axons formed symmetric synaptic contacts with presumptive granule cell dendrites and spines, and optogenetic stimulation demonstrated that activation of the reorganized neurons produced GABAergic inhibition in dentate granule cells [39].

The *in vivo* effects of the altered circuit are not known, but it is unlikely to provide normal control of granule cell activity. Because the reorganized fibers originated from GABA neurons in s. oriens of CA1, they would not receive the normal input from dentate granule cells that would be required for efficient feedback inhibition. However, strong activity of CA1 pyramidal cells could potentially activate these SOM neurons and produce inhibitory responses in the granule cells, although through an indirect circuit with presumably altered timing.

These results emphasize that the reemergence of a GABAergic axonal plexus does not necessarily indicate establishment of normal circuitry, and the reorganized circuit could be ineffective in controlling activity of principal cells. Such findings demonstrate yet another way in which GABA neuron loss could lead to altered inhibitory control and thus contribute to an epileptic state.

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## 12.5 Replacement of GABA Neurons Supports Their Functional Importance in Epilepsy

Recent studies of transplantation of GABA neurons in the hippocampal formation of pilocarpine-treated mice support contributions of GABA neuron loss to the epileptic state [28]. After GABA neuron transplantation in the hippocampal formation, the number of spontaneous seizures in these mice was reduced, despite the maintained presence of mossy fiber sprouting in

the inner molecular layer and, presumably, loss of mossy cells in the dentate hilus. While these findings are consistent with a loss of GABA neurons leading to the epileptic state, it remains possible that the transplanted GABA neurons could be counteracting other fundamental epilepsy-producing alterations through compensatory increases in inhibition.

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## 12.6 GABA Neuron Loss Has Multiple Effects in Epilepsy

Loss of even a small fraction of GABA neurons can have profound functional effects due to the innervation of numerous principal cells by the expansive axonal plexus of many interneurons. However the effects of an initial loss of GABA neurons could be enhanced further by alterations of remaining GABA neurons. Despite having some basic compensatory effects, the remaining GABA neurons could contribute periodically to the epileptic state through multiple mechanisms. These could include creation of excessive synchronous activity within the network and an inability of aberrant GABAergic circuitry to respond appropriately when increased inhibitory control is required. While still speculative, there is increasing evidence that GABA neuron loss, through both direct and indirect mechanisms, could give rise to the epileptic state.

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# Does Mossy Fiber Sprouting Give Rise to the Epileptic State?

# 13

Paul S. Buckmaster

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## Abstract

Many patients with temporal lobe epilepsy display structural changes in the seizure initiating zone, which includes the hippocampus. Structural changes in the hippocampus include granule cell axon (mossy fiber) sprouting. The role of mossy fiber sprouting in epileptogenesis is controversial. A popular view of temporal lobe epileptogenesis contends that precipitating brain insults trigger transient cascades of molecular and cellular events that permanently enhance excitability of neuronal networks through mechanisms including mossy fiber sprouting. However, recent evidence suggests there is no critical period for mossy fiber sprouting after an epileptogenic brain injury. Instead, findings from stereological electron microscopy and rapamycin-delayed mossy fiber sprouting in rodent models of temporal lobe epilepsy suggest a persistent, homeostatic mechanism exists to maintain a set level of excitatory synaptic input to granule cells. If so, a target level of mossy fiber sprouting might be determined shortly after a brain injury and then remain constant. Despite the static appearance of synaptic reorganization after its development, work by other investigators suggests there might be continual turnover of sprouted mossy fibers in epileptic patients and animal models. If so, there may be opportunities to reverse established mossy fiber sprouting. However, reversal of mossy fiber sprouting is unlikely to be antiepileptogenic, because blocking its development does not reduce seizure frequency in pilocarpine-treated mice. The challenge remains to identify which, if any, of the many other structural changes in the hippocampus are epileptogenic.

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## Keywords

Dentate gyrus • Granule cell • Epilepsy • Epileptogenesis • Hilus • Seizure • Pilocarpine

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## 13.1 Introduction

Temporal lobe epilepsy is common and its underlying mechanisms remain unclear [12]. Many patients have a history of an initial brain insult followed by a latent period [30]. A common view of temporal lobe epileptogenesis is that during the latent period, cascades of molecular and cellular events together alter the excitability of neuronal networks, ultimately causing spontaneous seizures [38]. According to this view, following an injury there is a critical period when temporary treatment might permanently prevent network reorganization. Substantial network reorganization occurs in the hippocampus of many patients with temporal lobe epilepsy [29]. The hippocampus is prone to epileptic activity [40] and is a site of seizure initiation in patients [37, 45]. Therefore, it is logical to ask whether structural changes in the hippocampus give rise to the epileptic state and whether blocking the development of structural changes during a critical period would prevent epileptogenesis. Structural changes in the hippocampus of patients with temporal lobe epilepsy include specific patterns of neuron loss [29], including inhibitory interneurons [11], hypertrophy of some surviving interneurons [28], GABAergic axon sprouting [1], dispersion of granule cells to ectopic locations [19], excessive development of hilar basal dendrites on granule cells [53], and mossy fiber sprouting (reviewed in [2]).

Philip Schwartzkroin's research included work on mossy fiber sprouting. His laboratory's slice experiments on tissue resected to treat patients revealed a general correlation between mossy fiber sprouting and hyperexcitability [13]. In those experiments, intracellular labeling and electron microscopy showed sprouted mossy fibers synapsing with dendrites of granule cells and interneurons. Those findings were supported and extended by experiments with kainate-treated rats that included evidence of autaptic synapses by sprouted mossy fibers [58]. However, Phil and colleagues cautioned that the results provided no direct evidence that mossy fiber sprouting was either necessary or sufficient for hyperexcitability [13].

Phil and colleagues characterized mossy fiber sprouting in other animal models, including p35-deficient mice with cortical dysplasia [57], different mouse strains treated with kainic acid [31], and infant monkeys after limbic status epilepticus [16, 56]. Phil's laboratory also helped localize the zinc transporter-3 to mossy fiber synaptic vesicle membranes [55], which has become a useful marker for visualizing mossy fiber sprouting.

Despite much investigation, the role of mossy fiber sprouting in epileptogenesis remains unclear and controversial. It has been proposed to be proepileptogenic [48], antiepileptogenic [44], and an epiphenomenon [15]. Recent evidence reviewed here raises questions about whether there is a critical period for mossy fiber sprouting after epileptogenic injuries and whether mossy fiber sprouting contributes to the generation of spontaneous seizures.

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## 13.2 Is Mossy Fiber Sprouting a Homeostatic Mechanism?

Neuron loss in the hilus of the dentate gyrus is a common structural change in patients with temporal lobe epilepsy [29] that is replicated in animal models. Nadler et al. (1980) first showed that the excitotoxin kainic acid kills hilar neurons, whose axons degenerate in the inner third of the molecular layer into which mossy fibers later sprout. To quantify the initial loss of synapses onto granule cell proximal dendrites in the inner molecular layer and the later restoration of excitatory synaptic input from sprouted mossy fibers, we used stereological electron microscopy to evaluate a rat model of temporal lobe epilepsy [49]. Tissue was obtained: (1) from rats 5 days after pilocarpine-induced status epilepticus to measure loss of synaptic input to granule cells before axon sprouting had occurred and (2) after mossy fiber sprouting was well established 3–6 months after status epilepticus. Numbers of granule cells were estimated from Nissl stained sections. Numbers of excitatory synapses in the molecular layer, where granule cell dendrites extend, were estimated in serial electron micrographs

that had been processed by post-embedding immunocytochemistry for GABA to avoid counting inhibitory synapses. Subsequently, numbers of excitatory synapses per granule cell were calculated for each rat (Fig. 13.1a). Analysis of the inner third of the molecular layer revealed that the number of excitatory synapses per granule cell decreased to only 38 % of controls by 5 days after pilocarpine-induced status epilepticus (Fig. 13.1b). This substantial loss of synapses probably is attributable primarily to loss of hilar mossy cells. Mossy cells, which were first characterized electrophysiologically by intracellular recording and anatomical labeling techniques in Phil's laboratory [39], project most of their axon collaterals to the inner molecular layer of the dentate gyrus where they form glutamatergic synapses with proximal dendrites of granule cells [6, 8, 54]. Epileptogenic injuries, like status epilepticus, kill mossy cells [4] and thereby denervate proximal dendrites of granule cells. The extent of mossy cell loss correlates with the extent of mossy fiber sprouting [23]. However, mossy cell loss alone is insufficient to cause mossy fiber sprouting [24], and the molecular signals necessary for triggering mossy fiber sprouting are not yet known.

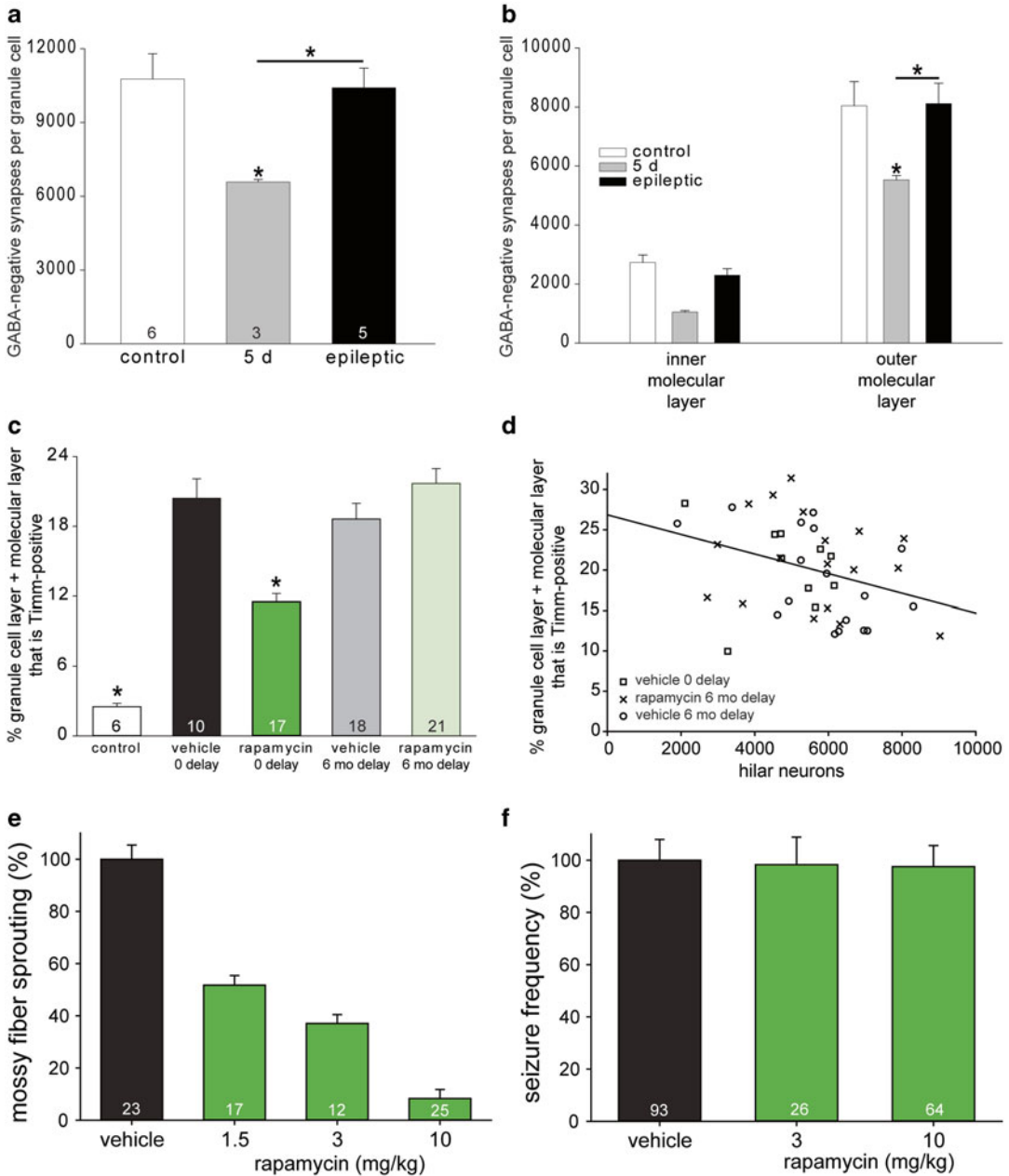
After initial loss, numbers of excitatory synapses per granule cell in the inner molecular layer partially rebound to 84 % of controls in rats 3–6 months after status epilepticus (Fig. 13.1b). This recovery probably is attributable primarily to mossy fiber sprouting [9], but other sources of excitatory synaptic input to the proximal dendrites of granule cells include surviving mossy cells and proximal CA3 pyramidal cells [60]. Synapses with proximal dendrites of granule cells at 3–6 months after status epilepticus are 1.3-times larger than in controls and twice as likely to be perforated [49]. Large, perforated synapses are likely to be functionally stronger than small, nonperforated synapses [14, 33, 35]. To maintain functional stability in the face of change, brains use an array of homeostatic mechanisms, including synaptic scaling [50]. Larger, stronger mossy fiber synapses in the inner molecular layer of epileptic rats might be a homeostatic mechanism to compensate for fewer synapses (84 % of controls, in this case).

Similarly, on granule cell distal dendrites in the outer two-thirds of the molecular layer, numbers of excitatory synapses decrease to 69 % of controls by 5 days after status epilepticus, but rebound to 101 % of controls by 3–6 months (Fig. 13.1b). With more complete recovery of synapse numbers, synapse size in the outer molecular does not change significantly [49]. Initial loss of synapses with distal dendrites of granule cells probably is attributable to partial loss of layer II entorhinal cortical neurons caused by status epilepticus [26]. And recovery of synapses probably is attributable to sprouting of axons of surviving layer II neurons [46]. Together, findings from the inner and outer molecular layer suggest a homeostatic mechanism maintains excitatory synaptic input to granule cells in response to synapse loss after an epileptogenic injury.

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### 13.3 No Critical Period for Mossy Fiber Sprouting

If a homeostatic mechanism controls the number of excitatory synapses with granule cells, signals underlying that control might persist as long as a synaptic deficit continues. Persistent signals contrast with the view of a transient cascade of molecular and cellular events that peak and then diminish after a critical period following a brain injury. To address these issues, we determined whether mossy fiber sprouting would occur after a 2 month delay [27]. Rapamycin, which inhibits mossy fiber sprouting [3], was administered to mice daily beginning 24 h after pilocarpine-induced status epilepticus. After 2 months, mossy fiber sprouting was suppressed almost by half in the rapamycin group compared to vehicle-treated controls (Fig. 13.1c). Another cohort was evaluated 6 months after the end of treatment, which was 8 months after status epilepticus. Mossy fiber sprouting was well developed in both vehicle- and rapamycin-treated mice, indicating that signals stimulating mossy fiber sprouting must have persisted for more than 2 months. These findings suggest there is no transient critical period for mossy fiber sprouting after an epileptogenic



**Fig. 13.1** Excitatory synapse loss and mossy fiber sprouting of granule cells in pilocarpine-treated rodent models of temporal lobe epilepsy. **(a)** Number of putative excitatory synapses per granule cell in control rats and in rats 5 days and 3–6 months (epileptic) after pilocarpine-induced status epilepticus. **(b)** Number of synapses with granule cell dendrites in the inner one-third and outer two-thirds of the molecular layer. Values represent mean  $\pm$  sem. Sample size indicated at base of bars. Asterisks indicate differences from the control value unless specified by a horizontal line ( $p < 0.05$ , ANOVA, Student-Newman Keuls method). **(a)** and **(b)** from Thind et al. [49]. **(c)** Extent of mossy fiber sprouting in control mice and mice that experienced status epilepticus and were treated with vehicle or 1.5 mg/kg rapamycin every day for 2 months and then were perfused with no delay (0 delay) or after a 6 month delay (6 month delay) (From Lew and Buckmaster [27]). **(d)** Number of

large hilar neurons ( $>12 \mu\text{m}$  soma diameter) per hippocampus versus extent of mossy fiber sprouting in mice that experienced pilocarpine-induced status epilepticus and were treated with vehicle or rapamycin for 2 months and then evaluated immediately (vehicle 0 months) or after another 6 months (vehicle or rapamycin 6 months). A linear regression line is plotted ( $R = 0.34$ ,  $p = 0.021$ , ANOVA). **(e)** Percent mossy fiber sprouting was calculated by subtracting the average percentage of the molecular layer plus granule cell layer that was Timm-positive in control mice and normalizing by the average value of mice that had experienced status epilepticus and were treated with vehicle for 2 months. Averages of all groups are significantly different from others ( $p < 0.05$ , ANOVA, Student-Newman-Keuls method). **(f)** Percent seizure frequency was calculated by normalizing by the average of the vehicle-treated group. **(e)** and **(f)** from Heng et al. [18]

brain injury. Instead, preventing mossy fiber sprouting might require long-term or continuous treatment. This scenario challenges the view of transient signaling cascades whose consequences could be permanently blocked by temporary treatment during a critical period.

One might question whether the precipitating injury in the mouse model was so severe that it maximally stimulated mossy fiber sprouting toward saturation levels despite the delay caused by rapamycin. However, an all-or-none “toggle-like” signal and saturation effect is inconsistent with the graded degree of mossy fiber sprouting among individual mice, which ranged over a factor of three and was correlated with the extent of hilar neuron loss (Fig. 13.1d). Wide ranges in mossy fiber sprouting between individuals were evident in vehicle-treated mice 2 months after status epilepticus and vehicle- and rapamycin-treated mice 8 months after status epilepticus, indicating that sprouting did not progressively develop toward saturated levels. These findings suggest that a target level of mossy fiber sprouting in an individual might be determined shortly after a brain injury and then remain constant.

Together, findings from stereological electron microscopy and rapamycin-delayed mossy fiber sprouting suggest a persistent, homeostatic mechanism exists to maintain a set level of excitatory synaptic input to granule cells. If mossy fiber sprouting were epileptogenic, this might be an example of a normally adaptive homeostatic mechanism that became pathogenic in response to an injury, which has been proposed previously as a theoretical possibility [10]. More generally, epileptogenesis might be an unintended side-effect of the brain’s homeostatic mechanisms, which evolved to maintain function in the face of plasticity. Epileptogenic injuries might trigger changes so much more extensive than normal plasticity that they push homeostatic responses into a range that creates a network that generates spontaneous seizures. Phil proposed a similar idea [41]: “I believe that the brain has been designed to operate at a knife’s edge. The evolutionary demand for plasticity – a key attribute of higher order learning, memory, and all those complex functions that are characteristic of the

mammalian CNS – has necessitated a sacrifice in stability of neuronal function.”

On the other hand, if epileptogenesis is maintained by homeostatic mechanisms gone awry, there may be opportunities to reverse established epilepsy-related structural abnormalities. Although mossy fiber sprouting appears to develop gradually, plateau, and then cease, there might instead be continual turnover. Mossy fibers in tissue from patients with temporal lobe epilepsy display evidence of continuing synaptic reorganization years after precipitating injuries [21, 32, 36]. At least some sprouted mossy fibers arise from adult generated granule cells [22, 25], which might continue to be generated long after precipitating injuries (but see [17]). To test whether mossy fiber sprouting could be reversed after it had established, we infused rapamycin focally into the dentate gyrus for 1 month beginning 2 months after pilocarpine-induced status epilepticus in rats, but there was no effect [3]. However, Huang et al. [20] reported that in chronically epileptic pilocarpine-treated rats, systemically administered rapamycin partially reversed already established mossy fiber sprouting. Moreover, grafts of CA3 pyramidal cells reduce mossy fiber sprouting even when implanted 45 days after kainate-treatment, during which time considerable mossy fiber sprouting is likely to have developed [42]. In addition, mild mossy sprouting generated by electroconvulsive shock was reported to decline over time [51]. Thus, more work is needed to test the reversibility of mossy fiber sprouting.

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### 13.4 Mossy Fiber Sprouting Is Not Epileptogenic

Rapamycin also was used to test whether mossy fiber sprouting was epileptogenic. Systemic treatment with rapamycin at increasing doses to inhibit mossy fiber sprouting to increasing degrees had no effect on the frequency of spontaneous seizures in mice that had experienced pilocarpine-induced status epilepticus (Fig. 13.1e, f) [5, 18]. These findings suggest that mossy fiber sprouting is neither pro- nor antiepileptogenic, but instead is

an epiphenomenon unrelated to seizure genesis. There are caveats with this conclusion, because rapamycin has side-effects [47], including suppression of axon sprouting by inhibitory GABAergic interneurons [7]. And rapamycin reduces seizure frequency in some rat models of temporal lobe epilepsy [20, 52, 59] but not all [43]. It remains unclear whether rapamycin's action in rats is antiseizure or antiepileptogenic. Nevertheless, the findings from the mouse studies suggest mossy fiber sprouting is not epileptogenic.

### 13.5 Conclusions

Patients with temporal lobe epilepsy display many structural changes, especially in the hippocampus. One possibility is that together numerous structural changes and other abnormalities all contribute partially to seizure generation. In that scenario, blocking the development of any one change, like mossy fiber sprouting, might have negligible effects on epileptogenesis. Another possibility is that some or perhaps even many epilepsy-related structural changes are not epileptogenic, including mossy fiber sprouting, and that seizure generation is attributable to one or two critical abnormalities whose importance has not yet been recognized. These alternate possibilities – many abnormalities each contributing partially versus one or two abnormalities primarily responsible for seizure generation—might require different therapeutic approaches, so it is important to distinguish between them. To do so, it will be useful to tap the ever-increasing knowledge base of molecular and cellular mechanisms underlying brain developmental processes and responses to injury. Creative application of ideas and reagents (for example, rapamycin), even from fields outside of epilepsy research, might yield useful approaches for specifically inhibiting or exacerbating individual epilepsy-related structural changes. Experimental manipulation of specific structural changes, one at a time, and rigorous measurement of effects on spontaneous seizures, might eventually reveal which, if any, are epileptogenic. If no single change alone appears to be responsible, then blockade of many or all could be

used to test whether together they make the brain epileptic or if the cause of seizures is unrelated to structural changes in the hippocampus.

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# Does Brain Inflammation Mediate Pathological Outcomes in Epilepsy?

# 14

Karen S. Wilcox and Annamaria Vezzani

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## Abstract

Inflammation in the central nervous system (CNS) is associated with epilepsy and is characterized by the increased levels of a complex set of soluble molecules and their receptors in epileptogenic foci with profound neuromodulatory effects. These molecules activate receptor-mediated pathways in glia and neurons that contribute to hyperexcitability in neural networks that underlie seizure generation. As a consequence, exciting new opportunities now exist for novel therapies targeting the various components of the immune system and the associated inflammatory mediators, especially the IL-1 $\beta$  system. This review summarizes recent findings that increased our understanding of the role of inflammation in reducing seizure threshold, contributing to seizure generation, and participating in epileptogenesis. We will discuss preclinical studies supporting the hypothesis that pharmacological inhibition of specific proinflammatory signalings may be useful to treat drug-resistant seizures in human epilepsy, and possibly delay or arrest epileptogenesis.

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## Keywords

Inflammation • IL-1 $\beta$  • TNF- $\alpha$  • IL-6 • Reactive astrogliosis

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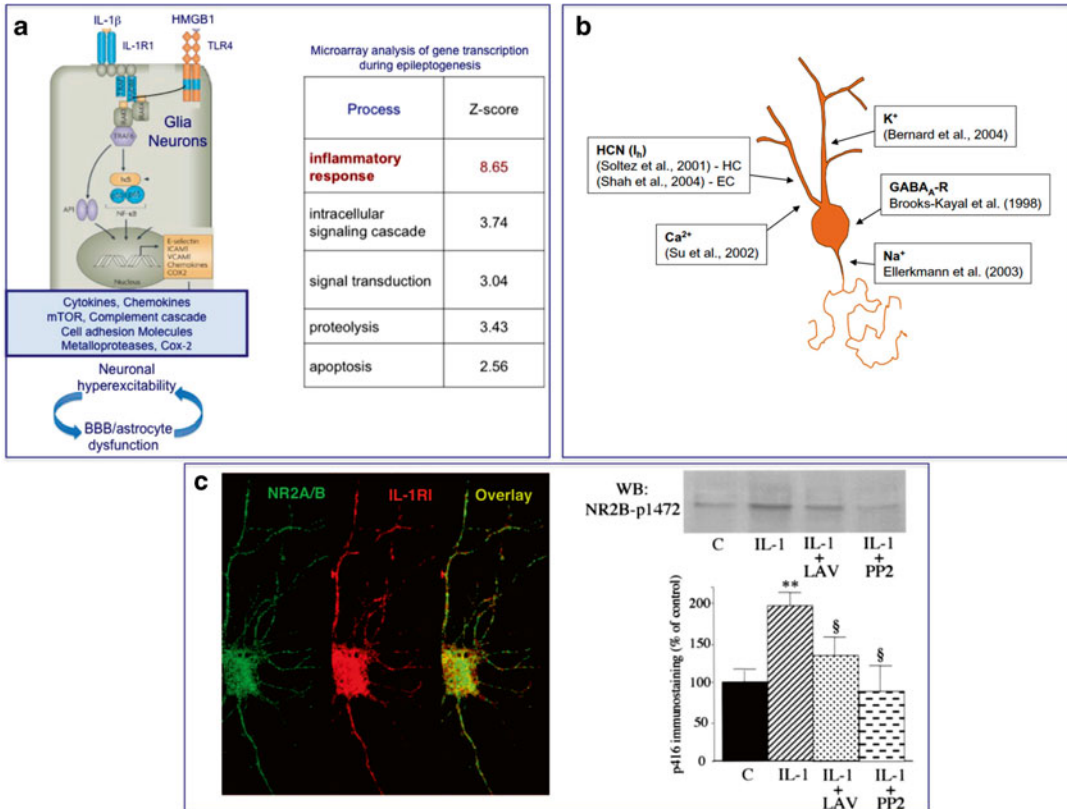
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## 14.1 Introduction

The state-of-the-art knowledge acquired in the last decade of experimental and clinical work indicates that cytokines and related molecules are increased in brain tissue after epileptogenic injuries or during seizures. In the experimental setting, these molecules, endowed with proinflammatory properties, contribute significantly to the generation





**Fig. 14.1** Schematic representation of the pathophysiologic outcomes of innate immunity activation in epilepsy. Activation of innate immune signaling occurs in epilepsy also in the absence of infection, thus triggering the so-called “sterile” inflammatory cascade (a). Endogenous molecules (damage associated molecular patterns, DAMPs) such as IL-1 $\beta$  and the High Mobility Group Box 1 (HMGB1) protein are released by neurons and glia following epileptogenic inciting events, or during recurrent seizures. The activation of their cognate receptors (IL-1R type 1 and TLR4, respectively) upregulated in astrocytes triggers the NF $\kappa$ B-dependent inflammatory genes cascade, thus

inducing various molecules with proinflammatory and neuromodulatory properties. The signaling activation in neurons increases excitability by provoking acquired channelopathies involving voltage-gated channels (HCN1) or AMPA and GABA-A receptor complexes (b), as well as by rapid activation of Src kinase inducing the phosphorylation of the NR2B subunit of the NMDA receptor thereby promoting neuronal Ca<sup>2+</sup> influx (c). This chain of event contributes to the generation and establishment of an hyperexcitable neuronal network by direct receptor-mediated neuronal effects or indirectly by inducing astrocytes and BBB dysfunctions

and maintenance of a hyperexcitable neuronal network, thus decreasing seizure threshold (Fig. 14.1) and making the occurrence of a seizure more likely.

A key question that basic science has been addressing is how these proinflammatory molecules affect neuronal and glial functions. Answers to this question will increase our knowledge of the complex mechanistic aspects of hyperexcitability following inflammation and will be instrumental in highlighting novel targets for

developing drugs and therapies that raise seizure threshold, prevent seizure generation after an inciting event, and inhibit their recurrence in chronic epilepsy.

### 14.1.1 Inflammatory Molecules as Neuromodulators

The presence of molecules with proinflammatory properties in brain specimens obtained from

**Table 14.1** Inflammatory mediators in human epilepsies and experimental models

<i>Clinical evidence</i>
Inflammatory mediators are overexpressed in epileptogenic foci in human pharmacoresistant epilepsy of differing etiologies (e.g. RE, LE, MCD, mTLE)
Microglia and astrocytes are main sources of inflammatory mediators in brain tissue; neurons and endothelial cells of the blood brain barrier (BBB) also contribute to the generation of brain inflammation
Leukocyte extravasation in brain depends on the etiology of epilepsy
BBB damage is often detected together with brain inflammation
<i>Experimental evidence</i>
Recurrent seizures and epileptogenic brain injuries induce inflammatory mediators in astrocytes, microglia, neurons, and microvessels in brain areas involved in seizure onset and generalization
This phenomenon is long lasting and may exceed the initial precipitating event by days or weeks depending on the epilepsy model. It is inadequately controlled by anti-inflammatory mechanisms
In models of epileptogenesis, inflammation initiates before the development of epilepsy
Specific anti-inflammatory treatments reduce acute and chronic seizures and delay their time of onset
Transgenic mice with perturbed cytokine signaling show altered seizure susceptibility
Proinflammatory insults decrease seizure threshold ( <i>acutely</i> and <i>long-term</i> )

patients with epilepsy has been described as “brain inflammation” (Table 14.1). However, there is emerging evidence that these molecules have neuromodulatory functions that activate signaling in neurons and glia that are different from those induced by the same molecules in leukocytes in the frame of a classical inflammatory response to infection. During infection, proinflammatory cytokines and related molecules are released during innate immunity activation by immunocompetent cells following “pathogen associated molecular patterns” (PAMPs) activation of toll-like receptors (TLRs) or nucleotide-binding oligomerization domain (NOD-like) receptors. Cytokine release activates inflammatory programs for pathogen removal and the subsequent induction of homeostatic tissue repair mechanisms. Notably, in humans affected by various forms of pharmacoresistant epilepsy of differing etiolo-

gies (e.g. Rasmussen’s (RE) and limbic encephalitis (LE), malformations of cortical development, and mesial temporal lobe epilepsy (mTLE)) increased inflammatory mediators are measured in epileptogenic foci in the absence of an identifiable active infectious process. However, it is also important to note that CNS infection, which is a common cause of TLE, can also result in a cytokine storm that affects excitability. In this context, evidence of HHV6 infected astrocytes and neurons has been reported in about 2/3 of patients with mTLE [108]. Moreover, recent work has shown the presence of Human Papilloma virus in human focal cortical dysplasia type II which might be responsible for focal epileptogenic malformations during fetal brain development in association with enhanced mTORC1 signaling [18].

The so-called *sterile inflammation* in the brain can be induced when TLRs are activated by endogenous molecules released by injured brain cells, named “danger signals” or “damage-associated molecular patterns” (DAMPs). In particular, the activation of TLR4, which can also be activated by gram-negative bacteria, is induced by the ubiquitous nuclear protein High Mobility Group Box 1 (HMGB1) which is released, upon its cytoplasmic translocation, by neurons and glial cells. In concert with IL-1 $\beta$  released by glia, thereafter activating IL-1 receptor type 1 (IL-1R1), HMGB1 induces the transcriptional up-regulation of various inflammatory genes, therefore promoting the generation of the brain inflammatory cascade in glia and endothelial cells of the BBB (Fig. 14.1). In the context of malformations of cortical development, the inflammatory cascade is also induced in aberrant neuronal cells [3]. The activation of the IL-1R1/TLR4 signaling in neurons, which overexpress these receptors in pathologic conditions, in concert with pathways induced by other cytokines such as TNF- $\alpha$ , IL-6, the complement system and some prostaglandins, alters neuronal excitability by modifying either glutamate or GABA receptor subunit composition, or trafficking of receptors, or the function of voltage-gated ion channels via rapid onset post-translational mechanisms [118, 123]. Furthermore, initiation of the JAK/STAT and other signaling pathways through these mechanisms can also result in activation of

**Table 14.2** Antagonism of IL-1R1/TLR4 in rodent models of seizures

<i>Seizure reduction in rodents exposed to an acute challenge</i>
Kainic acid (lesional model), bicuculline and febrile seizures (non lesional models) [28, 87, 114, 119]
Status epilepticus [24, 64]
Electrical rapid kindling [88, 5, 6]
<i>Chronic recurrent seizures reduced in</i>
mTLE mouse model [66, 67]
SWD in GAERS & WAG/Rij (absence seizures models) [1, 49]
<i>Other inflammatory signaling contributing to seizures are mediated by</i>
TNF- $\alpha$ , IL-6, COX-2 & complement system (reviewed in [50, 115, 3])

glial cells, inducing a cascade of events that alters their structure and function in a variety of ways that can also contribute to aberrant excitability [99].

In animal models, pharmacological intervention to block or activate specific inflammatory pathways induced in human epilepsy brain specimens has shown that: (i) cytokines such as IL-1 $\beta$ , TNF- $\alpha$ , and IL-6, and danger signals such as HMGB1 and S100 $\beta$ , contribute to seizures in a receptor-dependent manner; (ii) the complement system contributes to seizure generation and cell loss; and (iii) PGE2 contributes to cell loss by activating EP2 receptors in neurons (Table 14.2). This set of evidence is corroborated by the assessment of susceptibility to seizures and cell loss in transgenic mouse models with impaired or overexpressed inflammatory signalings [118].

### 14.1.2 IL-1 $\beta$ , HMGB1 and the NMDA and GABA Receptors

IL-1 $\beta$  and HMGB1 both potentiate NMDA receptor function in cultured hippocampal neurons using post-translational mechanisms mediated by activation of IL-1R1 and TLR4, respectively [8, 53, 121]. In particular, these cytokines enhance NMDA-mediated Ca<sup>2+</sup> influx by activating Src kinases-dependent NR2B phosphorylation (Fig. 14.2). This signaling has been demonstrated

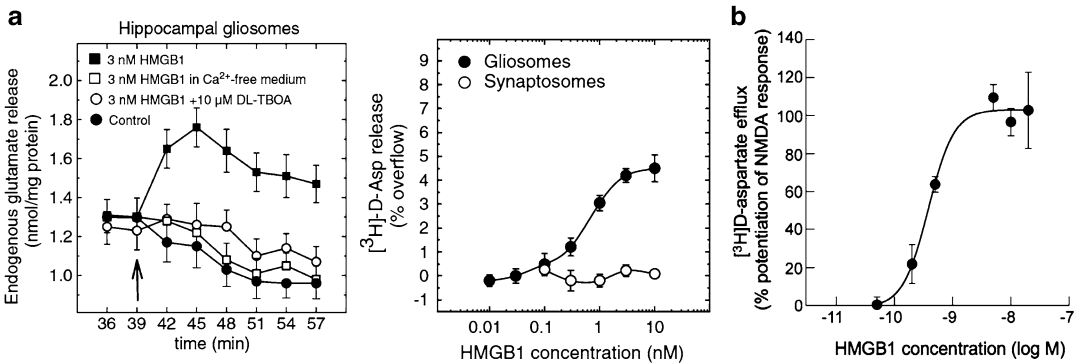
to underlie the proictogenic and proneurotoxic properties of these cytokines [7, 8, 40, 121].

This rapid onset (within 2 min) mechanism is reminiscent of that induced by IL-1 $\beta$  in hypothalamic neurons, which underlies the initial rise in body temperature induced by this cytokine [23, 91, 105], and it involves MyD88-dependent and ceramide-mediated activation of Src kinases. IL-1 $\beta$  also down-regulates AMPA receptor expression and their phosphorylation state in a Ca<sup>2+</sup>- and NMDA-dependent manner in hippocampal neurons [53]. Recent evidence shows that HMGB1 effects on neuronal excitability may also include a physical, receptor unrelated, interaction with presynaptic NMDA receptors resulting in enhanced Ca<sup>2+</sup>-dependent glutamate release from presynaptic terminals evoked upon NMDAR stimulation [80]. Notably, HMGB1 per se can also induce glutamate release from hippocampal gliosome preparations implying that this molecule may increase gliotransmission [81]. While the effect of IL-1 $\beta$  and HMGB1 on NMDA-induced Ca<sup>2+</sup>-influx in neuronal cell soma and dendrites mediates cell loss and increases seizures [7, 8, 121], whether the effect of HMGB1 on presynaptic or glial glutamate release results in pathologic outcomes has not been yet investigated.

Excitatory actions of IL-1 $\beta$  have been reported in hippocampal slices or cultured pyramidal neurons where the cytokine reduces synaptically-mediated GABA inhibition in CA3 hippocampal region via still unidentified kinases [123, 129], and increases CA1 neurons excitability by reducing NMDA-induced outward current. This latter action involves activation of cytoplasmatic P38 MAPK phosphorylating large-conductance Ca<sup>2+</sup>-dependent K channels [131].

### 14.1.3 Cytokines, Synaptic Transmission/Plasticity and Seizures

Cytokine receptors are expressed by the same resident CNS cells that express their cognate cytokines, namely neurons, microglia, and astrocytes. Binding of ligands to these receptors set



**Fig. 14.2** Presynaptic and postsynaptic effects of HMGB1 on glutamatergic transmission. HMGB1 protein evokes (<sup>3</sup>H)D-aspartate and glutamate release from re-sealed glial (gliosomes) and neuronal (synaptosomes) subcellular particles isolated from the mouse hippocampus (a). This protein per se augments the calcium-independent neurotransmitter outflow from gliosomes, but not from synaptosomes, in a concentration-dependent manner. This outflow is

likely mediated by reversal of glutamate transporter (GLAST) since it is blocked by DL-threo-b-benzyloxyaspartate (TBOA) [81]. HMGB1 augments the NMDA-induced (<sup>3</sup>H)D-aspartate calcium-dependent release from synaptosomes (b). This enhancing effect is mediated by increased intracellular calcium via the MK-801 sensitive channel. This HMGB1-NMDA receptor interaction involves the NR2B subunit [80]

into motion a variety of signaling pathways that activate glial cells and can also lead to enhanced excitability of neurons.

**IL-1 $\beta$ .** In the hippocampus, IL-1 $\beta$  was reported to induce rapid changes in synaptic transmission, and to inhibit LTP via activation of MAPK and PKC [12, 75, 84, 96]. Fast neuronal actions of IL-1 $\beta$  were described in the preoptic/anterior hypothalamic neurons involving A-type K<sup>+</sup> currents and the consequent reduced synaptic release of GABA [105].

**TNF- $\alpha$ .** Work by Stellwagen et al. demonstrated that TNF- $\alpha$  released by astrocytes binds to the TNF- $\alpha$  receptors (TNFR) on neurons and induces an increase in AMPA-type glutamate receptors and a concomitant decrease of GABA<sub>A</sub> receptors at synapses [102]. Specifically, TNF- $\alpha$  has been shown to increase trafficking of GluR2-lacking AMPA receptors to synaptic membranes in both hippocampal and motor neurons [11, 55, 56, 102, 103, 126]. In hippocampal neurons, this trafficking has been shown to depend on the PI3K–Akt pathway [102]. GluR2-lacking receptors are permeable to Ca<sup>2+</sup> and activation of these receptors could dramatically alter synaptic strengths at these synapses or contribute to excitotoxicity. While TNFR knock out mice do not appear to have impaired long term potentiation (LTP) or long term depression (LTD), synaptic

scaling may be modulated by TNF- $\alpha$  [101, 103]. While it is currently unclear what role TNF- $\alpha$  signaling may be playing in receptor trafficking in epilepsy, recent work using the Theiler's Murine Encephalomyelitis Virus (TMEV) model of TLE has demonstrated that there is over a 120-fold increase in whole brain TNF- $\alpha$  mRNA soon after infection in C57Bl/6 mice [47]. This dramatic increase in TNF- $\alpha$  expression is associated with acute seizures and changes in mEPSC amplitudes and decay times in hippocampal brain slices prepared from animals acutely infected with TMEV [57, 98, 104]. In addition, TNFR1 knockout mice are much less likely to exhibit seizures during the acute infection period. Taken together, the evidence suggests an important role of TNF- $\alpha$  in modulating excitatory circuits and excessive amounts of TNF- $\alpha$  may contribute to seizure activity. Accordingly, a proictogenic role of TNF- $\alpha$  mediated by TNFR1, and an opposite anti-ictogenic role of this cytokine mediated by TNFR2 have been reported in chemoconvulsant models of seizures [7–9, 124]. Molecular and functional interactions between TNFR and the glutamatergic system in the hippocampus appear to be implicated in the effect of this cytokine in seizure susceptibility [8].

In addition to modifying synaptic transmission, TNF- $\alpha$  is also known to stimulate the release of

glutamate from microglia [17, 107] and astrocytes [92, 93], and these additional sources of extracellular glutamate likely contribute to excitotoxicity in injured brain regions. Activation of TNFR in cultured microglia results in an increased expression of glutaminase, which converts glutamine to glutamate. This excess intracellular glutamate is then released through connexin 36 hemi-channels and can be blocked by the gap junction inhibitor, carbenoxolone [107]. It is thought that this mechanism can contribute to neuronal cell death that often accompanies chronic or prolonged tissue inflammation.

*IL-6.* Recent work has demonstrated that IL-6, another cytokine that is increased in response to epileptogenic insults, decreases GABA and glycine-mediated inhibitory synaptic currents following bath application to spinal cord slices [46]. Such changes in synaptic neurotransmitter receptor function can result in tipping the balance of excitation and inhibition towards hyperexcitability. Binding of IL-6 to its receptor results in the activation of the JAK/STAT pathway and this pathway is known to regulate the expression of many different receptor gated ion channel subunits [60] and underlies NMDA-dependent LTD in the hippocampus [72]. Therefore, changes in IL-6 expression levels could dramatically influence excitability of neural circuits responsible for seizure generation. Recent work with the TMEV mouse model of TLE, demonstrated that IL-6 mRNA expression increases significantly during the acute infection period and this increase parallels the onset of seizures in this model. Furthermore, IL-6 receptor knockout mice have a reduced incidence of seizures following TMEV infection, suggesting that this cytokine, which is largely expressed in this animal model by infiltrating macrophages, contributes to lowering seizure thresholds [21, 47]. Finally, treatment of TMEV infected mice with either minocycline or wogonin, were both found to dramatically reduce concomitantly the number of infiltrating macrophages in the brain and seizure incidence [21]. These results suggest that IL-6 may be an important regulator, possibly through the JAK/STAT pathway, of synaptic plasticity and seizure activity.

#### 14.1.4 Cytokines and Voltage-Gated Ion Channels

While cytokines have been extensively studied in neuropathic pain and in epilepsy, very few studies have examined the effects of the prominent cytokines on voltage gated ion channels (see [122]). Nevertheless, the limited available literature demonstrates that cytokines can modulate a variety of voltage gated ion channels through multiple mechanisms [95]. For example, TNF- $\alpha$  has been shown to increase expression of TTX resistant sodium channels in isolated dorsal root ganglion cells, increase Ca<sup>2+</sup> currents in cultured hippocampal neurons and decrease inwardly rectifying K<sup>+</sup> currents in cultured cortical astrocytes [35, 44, 48]. IL-1 $\beta$  has been shown to decrease Ca<sup>2+</sup> currents in cultured hippocampal and cortical neurons [83, 84, 132, 133] as well as Na<sup>+</sup> and K<sup>+</sup> currents in dissociated retinal ganglion cells [26].

The effect of cytokines on ion channel function is an area where clearly further work is necessary so as to inform hypotheses about the full range of activity of cytokines in epilepsy, particularly in view of the plethora of differing effects on neuronal functions that cytokines may have depending on their concentration, timing of tissue exposure, the type of neuronal cells expressing the relevant receptors, and the concomitant presence of other neuromodulatory molecules.

#### 14.1.5 Prostaglandins, Synaptic Plasticity and Seizure Activity

Arachidonic acid (AA) is converted to prostanoids via activity of the enzyme cyclooxygenase (COX). COX-2 is constitutively active at low levels in the hippocampus, its expression rapidly increases as a consequence of neural activity, and is necessary for some forms of synaptic plasticity, such as LTP in the dentate gyrus [42]. Prostaglandin E2 (PGE2), one of the most common of the prostanoids to be formed in the hippocampus, binds to the G-protein coupled EP2 receptor on neurons, activates cAMP and mediates synaptic plasticity via the cAMP-protein kinase A (PKA)-cAMP-responsive

element binding protein (CREB) pathway [42, 116]. Following status epilepticus (SE), COX-2 expression is increased in the hippocampus and prostaglandins, including PGE<sub>2</sub>, are also subsequently increased and hypothesized to be involved in mediating neurodegeneration that occurs in multiple brain regions following SE. This neurotoxic effect may be due to excessive stimulation of EP2 receptors expressed by microglia and the consequent activation of an alternative pathway, the cAMP-Epac signaling pathway promoting upregulation of various inflammatory mediators and oxidative stress [42]. Whereas pharmacological inhibition of COX-2 can be neuroprotective following CNS insults, this approach has not yielded great success in preventing the development of epilepsy following SE although disease-modifying effects have been reported [45, 51, 61, 85]. Depending on the drug used to inhibit COX-2 and the trigger of SE, adverse events have also been described in epileptic rats [39, 85]. Therefore, the search is on for drugs that can selectively interfere with downstream pathways of COX-2 in an effort to mitigate the detrimental inflammatory actions that can occur in the CNS following SE. Recently, Jiang et al. evaluated the ability of a novel small molecule and brain permeable EP2 antagonist, TG6-10-1, to confer neuroprotection and prevent the development of epilepsy in mice treated with pilocarpine [43]. Encouragingly, there was significant neuroprotection and decreased mortality following SE in the treated mice. However, there were no differences with vehicle-treated mice in spontaneous seizure frequency, suggesting that epileptogenesis was not interrupted with this treatment [43]. This suggests that adjunctive therapy with an EP2 antagonist may be important for attaining neuroprotection in patients experiencing SE, but additional approaches will be necessary to prevent the development of epilepsy. In this context, a recent study reported that co-treatment with IL-1 receptor antagonist (IL-1Ra, anakinra) and a COX-2 inhibitor given at the time of SE induction were required to reduce both cell loss and epileptogenesis in rats [52]. Similarly, combined treatment with IL-1Ra and VX-765, an inhibitor of IL-1 $\beta$  biosynthesis, given systemically to rats after 3 h of uninterrupted

SE, afforded significant neuroprotection although not inhibiting epilepsy development [74]. This evidence highlights the need of both early intervention and combined anti-inflammatory treatments for optimizing beneficial clinical outcomes.

Another strategy to be investigated is a combination of specific anti-inflammatory drugs with classical antiepileptic drugs (AED) targeting complementary mechanisms. Indeed, some AEDs afford neuroprotection or decrease the severity of spontaneous seizures induced in SE models [71].

#### 14.1.6 TLR4 and Neuronal Excitability

Out of 11 members of the TLRs family, TLR4 is the most extensively studied in CNS for its involvement in increasing brain excitability and cell loss, and for reducing neurogenesis.

Rat cortical application of lipopolysaccharide (LPS), a PAMP component of gram-negative bacteria wall and prototypical activator of TLR4, has been reported to rapidly increase the excitability of local neurons as assessed by measuring amplitudes of sensory evoked field potentials following rat forepaw stimulation and spontaneous activity [90]. A ten-fold higher LPS concentration could evoke epileptiform activity which was prevented by pre-application of IL-1Ra, implicating a role of IL-1 $\beta$  released from LPS-activated microglia [90].

We recently discovered that intracerebral LPS application reduces hyperpolarization-activated ion channel (HCN1) protein in hippocampal tissue, an effect associated with a reduction in I<sub>h</sub> current as assessed in whole-cell patch recording of CA1 pyramidal neurons. This effect is long-lasting but reversible upon resolution of both microglia activation and induction of proinflammatory cytokines in these cells. The activation of IL-1R1/TLR4 signaling is responsible for this effect since it was precluded in TLR4 or IL-1R1 knock-out mice, and by pharmacological blockade of these receptors with selective antagonists (Bernard et al., 2013, personal communication).

The reported LTP and LTD impairment induced by TLR4 stimulation is compatible with neurological dysfunction and cognitive deficits induced by early life exposure to LPS which are associated with specific and persistent changes in NMDA receptor subunits expression in the cortex and hippocampus, predicting modifications in CNS excitability (for review see [89, 127]).

### 14.1.7 Inflammation-Induced Functional Changes in Astrocytes

Reactive astrogliosis occurs as a consequence of cytokine activation of the IL-1R/TLR and JAK/STAT pathway and other signaling pathways following CNS insults such as traumatic brain injury (TBI), SE, and infection [99]. Astrogliosis is a graded process and is characterized by hypertrophy of primary processes, dramatic increases in the expression of intermediate filament proteins such as glial fibrillary acidic protein (GFAP), a decrease and cell redistribution in glutamine synthetase [20, 29, 78, 125], an increase in expression of adenosine kinase, and, in some cases, a disruption in domain organization of glial processes [76, 99]. There is also a dramatic increase in gap junction coupling between astrocytes in animal models [106] and resected human tissue [19, 32, 70], and a number of specific subunits of kainate receptors (KAR) were recently found to be expressed in reactive astrocytes following chemoconvulsant-induced SE in rodents [112]. There are, therefore, a multitude of changes in astrocytes following seizure-inducing insults and these changes may have a dramatic impact on the circuit dynamics underlying seizure generation [25, 36].

As astrocytes are intricately involved in regulating neuronal activity at the tri-partite synapse (review [2]), some of the changes in glial function that are observed in rodent models and human epilepsy could easily lead to hyperexcitability in neural circuits and contribute to seizure generation. For example, decreases in the endogenous anticonvulsant adenosine as a consequence of increased expression of adenosine kinase can lead

to hyperexcitability and seizure activity [4, 15] and, while early after SE, glutamate uptake by astrocytes seems to be functioning well [106], there are numerous reports of cytokine-mediated decreases in glutamate transporter function in epilepsy and other disorders which could readily lead to excess excitation and cell death in vulnerable neurons [62, 68, 86, 94]. Reactive astrocytes have also been reported to have a decrease in the inward rectifier potassium channel ( $K_{IR}$ ), namely Kv4.1, a critical ion channel that aids in the buffering of extracellular potassium concentrations, and this altered expression may be mediated by IL-1 $\beta$  [134]. Electrophysiological recordings in acute brain slices obtained from surgical specimens of patients with mTLE, have revealed a reduced  $K_{IR}$  conductance in reactive astrocytes [38]. However, we recently demonstrated that  $K_{IR}$  mediated currents were not altered in astrocytes during the latent period up to 2 weeks following SE in the KA-treated rat [106], and this is consistent with a recent report demonstrating that initial decreases in Kv4.1 mRNA and protein return to control levels by day 7 after SE [134]. Therefore, reactive astrocyte function may change over time as epilepsy develops.

While many of the observed changes in astrocytes that occur as a consequence of inflammation may actively contribute to network hyperexcitability, other components of reactive astrogliosis, such as increased gap junctional coupling, or increased neurotrophins may be critical compensatory mechanisms following injury, and may act to dampen excitability and protect neurons [36]. Thus, simply blocking the inflammatory response in glial cells may be too global an approach for disease modification during epileptogenesis, while targeting specific processes, such as maintaining  $K_{IR}$  function, might prove to be a more useful approach.

### 14.1.8 Cytokines Effects on BBB: Consequences for Neuronal Excitability

Evidence obtained using in vitro models of the BBB [31, 130] or epilepsy models [58, 77, 111, 116]

demonstrated that cytokines and prostaglandins compromise the permeability properties of the BBB, and that such alteration in brain vessels is a common feature of drug-resistant epileptogenic foci in humans and experimental models. In particular, there is evidence of the presence of IL-1 $\beta$  in perivascular glia and astrocytic endfeet impinging on brain vessels in epilepsy tissue where the BBB is altered, as shown by the parenchymal extravasation of serum macromolecules such as albumin and IgG. One mechanism of BBB damage induced by cytokines involves breakdown of tight-junction proteins in brain vessels [58, 59, 69, 73] induced by activation of Src kinases. This evidence highlights that key molecular pathways activated by cytokines in epilepsy result in different outcomes depending on the target cell population (expressing the relevant receptors), i.e. BBB permeability function is compromised in vessels, hyperexcitability is induced in neurons, and astrocyte function is greatly modified.

BBB damage leads to albumin extravasation which induces TGF- $\beta$  signaling in astrocytes by activating the TGF- $\beta$  receptor type 2 [33]. This signaling mediates transcriptional up-regulation of IL-1 $\beta$  and other inflammatory genes in astrocytes [16, 34] while glutamate transporter and Kir4.1 channels are down-regulated. These pathologic changes have been shown to establish a hyperexcitable milieu in surrounding neurons due to increased extracellular K<sup>+</sup> and glutamate [97] which decreases seizure threshold and may induce per se epileptiform activity [22, 34].

#### 14.1.9 Leukocytes, Autoantibodies and Neuronal Excitability

There is evidence of adaptive immunity activation in rare disorders such as Rasmussen's encephalitis (RE), viral and limbic encephalitis and neurological or systemic autoimmune disorders. These conditions are often associated with seizures and epilepsy development. In RE brain tissue, cytotoxic CD8<sup>+</sup> T lymphocytes have been demonstrated in close apposition to neurons and astrocytes, then provoking their apoptosis by releasing granzyme

B [10, 79]. The presence of these cells, and more in general CD3<sup>+</sup> leukocytes, appears to be much less prominent in more common forms of epilepsy. For example, in focal cortical dysplasia (FCD) type 2, scattered lymphocytes have been described in brain tissue while this phenomenon occurs at a minor extent in FCD type 1, and is almost undetectable in mTLE [41, 65, 110]. Others have detected leukocytes in brain parenchyma surrounding brain vessels also in mTLE [30, 128]. In animal models of epilepsy the role of these cells is still uncertain since they were reported to mediate anti-epileptogenic and neuroprotective effects in KA-treated rats [128] whereas they contribute to the pathology in pilocarpine-treated mice [30]. Notably, in this latter instance the effects of leukocytes may be ascribed to the peculiar mechanisms mediating seizures caused by pilocarpine and which are not shared by other chemoconvulsants [64, 109, 117].

A recent randomized clinical study using tacrolimus, which impedes T cell proliferation and activation, in recent onset RE patients showed delayed deterioration of neurological deficits but the treatment did not ameliorate drug resistant seizures [13]. However, case reports have shown decreased seizure frequency in one RE patient treated with natalizumab, a blocker of T cell entry into the CNS [14] and in a patient with multiple sclerosis and refractory epilepsy [101]. The authors discussed that interpretation of data was limited by an additional coadministration of varying antiepileptic medications.

In limbic encephalitis and autoimmune disorders, circulating autoantibodies against various neuronal proteins have been detected (for review, see [120]). These antibodies recognizing membrane neuronal proteins may have a pathologic role, in addition to their diagnostic value. In particular, antibodies against NR1/NR2 subunits obtained from serum of affected patients can increase extracellular hippocampal glutamate levels when intracerebrally infused in rats. Increased sensitivity to AMPA receptor-mediated neuronal excitability and GABAergic dysfunction have also been reported [63]. Antibodies directed against voltage-gated K<sup>+</sup> channel complex increase excitability of hippocampal CA3 pyramidal



cells by reducing channel function at mossy fiber-CA3 synapses [54]. AMPA receptor antibodies alter synaptic receptor location and number by reducing those receptors containing the GLUR2 subunit, therefore increasing the relative abundance of Ca<sup>2+</sup>-permeable receptors [53].

## 14.2 Conclusions

While understanding of the role of the innate immune system and the associated molecules with inflammatory properties in epilepsy and seizure threshold changes has advanced tremendously over the last decade, there are still a number of questions that yet remain open and require further investigation. For example, it is not yet clear which molecules and inflammatory pathways activated following epileptogenic brain insults will make the most appropriate targets for intervening to prevent seizure occurrence and/or the process of epileptogenesis. The complex network changes that occur in a number of cell types in the CNS, including neurons, microglia and astrocytes, in response to increases in a myriad of neuromodulatory and inflammatory molecules such as IL-1 $\beta$ , TNF- $\alpha$ , IL-6 and interferon- $\gamma$  to name but a few, are difficult to decipher. Moreover, it has still to be determined which are the master regulators of the inflammatory cascade, and when and how to prevent the induction of brain inflammation or rather promote its resolution by implementing the effects of the endogenous antiinflammatory molecules, which are defective in epilepsy [82, 87].

Nevertheless, the increasing recognition that the innate immune system is tightly coupled to epileptogenesis and seizure threshold changes is encouraging as it opens up many potential novel molecular targets for therapeutics. Most AEDs are mainly antiseizure, symptomatic drugs that target neuronal proteins such as sodium channels or glutamate receptors. Their adverse effects on cognition and induction of sedation, coupled with the knowledge that nearly 30 % of patients with epilepsy do not have their seizures adequately controlled with current AEDs, suggest that targeting the neuromodulatory inflammatory pathways is a promising novel strategy with disease-modifying

potential. Considering that prolonged administration in epilepsy is likely to be required, and the constraints imposed by the BBB, both the efficacy and the safety of drugs that preclude or reverse the over-activation of specific innate immune mechanisms should be carefully considered. Importantly, some of these antiinflammatory drugs are already in clinical use showing therapeutic effects in peripheral inflammatory conditions [27, 37, 113]. These drugs might be considered to complement the symptomatic treatment provided by available AEDs for resolving the inflammatory processes in the brain, therefore raising seizure threshold and decreasing the likelihood of seizure recurrence. In this context, a phase 2 clinical study with VX765 has given promising results in adult patients with drug resistant partial onset seizures (<http://clinicaltrials.gov/ct2/show/NCT01048255>; [www.epilepsy.com/files/Pipeline2012/6-7](http://www.epilepsy.com/files/Pipeline2012/6-7)).

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# Are Changes in Synaptic Function That Underlie Hyperexcitability Responsible for Seizure Activity?

# 15

John G.R. Jefferys

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## Abstract

The synaptic and intrinsic mechanisms responsible for epileptic seizures and briefer interictal epileptic discharges have been characterized in some detail. This chapter will outline some aspects of this work in the context of focal epilepsies, particularly in the temporal lobe, and will identify some of the major questions that remain. Early work, mainly using the actions of convulsant treatments on brain slices *in vitro*, revealed synaptic circuitry that could recruit populations of neurons into synchronous epileptic discharges. Subsequent investigations into cellular mechanisms of chronic experimental and clinical foci, again often *in vitro*, have revealed complex changes in synaptic properties, synaptic connectivity, intrinsic neuronal properties and selective losses of neurons: unraveling their roles in generating seizures, interictal discharges and interictal dysfunctions/comorbidities remains a significant challenge. *In vivo* recordings have revealed aspects of the pathophysiology of epileptic foci that have practical implications, for instance high-frequency oscillations, and potentially high-frequency hypersynchronous neuronal firing, which have been useful in localizing the epileptogenic zone for surgical resection.

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## Keywords

Temporal lobe epilepsy • Disease models • Cellular electrophysiology • Synaptic transmission • Chronic models • Hippocampus

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## 15.1 Introduction

Phil Schwartzkroin pioneered cellular electrophysiology and basic research on epilepsy. I started my research career a couple of years after he did. His work impressed and inspired me from the start. His impact goes far wider than his considerable innovations and insights into the basic mechanisms

of epilepsy, notably through his leadership roles in epilepsy societies and his development of influential monographs such as the “Encyclopedia of Basic Epilepsy Research” and “Models of Seizures and Epilepsy” [53, 62]. In summary Phil has made major contributions to the development of the basic science of epilepsy, which thoroughly justify this volume celebrating his career.

Seizures are the diagnostic feature of epilepsy and are classically considered as hypersynchronous electrophysiological activity [52]. The idea that seizures are hypersynchronous has been challenged recently, a point I will return to at the end of this chapter [36]. While the remit of this chapter is on the role of synaptic function in hyperexcitability and seizure generation, I will address broader issues on the pathophysiology of focal epilepsy. A quick definition of the terms of the title: hyperexcitability is a condition of neurons or neuronal networks in which they respond more intensely or more readily to normally innocuous activity, or may become spontaneously active; such responses or spontaneous activity may lead to the generation of seizures or other (briefer) epileptic discharges. Hyperexcitability can lead to seizures but the two are distinct concepts.

Normal brain tissue can generate seizures when exposed to convulsant conditions, as in acute models of epilepsy. Such acute models laid much of the groundwork for our understanding of basic mechanisms of clinical epilepsy, as well as being directly relevant to clinical symptomatic seizures. However, epilepsy is by definition a chronic condition where seizures occur spontaneously under physiological conditions; understanding why they do is a major challenge for both clinical and basic research.

Phil Schwartzkroin pioneered many of the models, preparations and concepts involved in understanding seizures and hyperexcitability, including: acute and chronic models of epilepsy, *in vitro* brain slices, synaptic properties, synaptic connectivity, intrinsic neuronal properties, glial properties ([39, 40]; for example: [60, 61, 63–68]). This chapter will outline work on basic mechanisms

of epilepsy, particularly focal epilepsy of the medial temporal lobe, which has, in large part, developed from these innovations.

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## 15.2 Acute Epilepsy and Hyperexcitability

Some of the earliest epilepsy research used acute pharmacological block of inhibition (e.g. with penicillin, picrotoxin or bicuculline) to produce epileptic discharges, initially *in vivo* and then in hippocampal and neocortical slices *in vitro*. The introduction of brain slices *in vitro* into epilepsy research was a major step towards developing detailed cellular models of epileptic activity. Early advances included the discovery that the intrinsic electrical properties of central neurons were rather complex, and could look a lot like epileptic bursts [67, 75]. Another early discovery was that excitatory pyramidal cells were interconnected to form a recurrent excitatory network that was held in check by networks of inhibitory neurons [40, 46]. It is hard to think back to the state of the field 40+ years ago, but the idea that excitatory neurons within a brain structure made connections with each other seemed novel at that time. Changing that mindset is an example of how epilepsy research can make major contributions to our understanding of normal brain mechanisms [44]. Essentially the idea that emerged is that excitatory neurons in regions such as the hippocampus and neocortex form interconnected excitatory synaptic networks which present the risk of a chain reaction of positive feedback [77]. Normally negative feedback, mediated by some types of inhibitory neuron, prevents the build-up of a chain reaction, but blocking or depressing GABAergic transmission clearly disrupts this control. Other convulsant treatments include changes in extracellular ions (e.g.  $K^+$  or  $Mg^{2+}$  or  $Cl^-$ ) or channel blockers (e.g. 4-aminopyridine) have also been investigated in some depth (for review see [31]). Arguably all of these treatments make neurons and/or neuronal networks hyperexcitable.



In practice the ability of isolated hippocampal slices *in vitro* to generate synchronous epileptiform discharges depends on both synaptic and intrinsic neuronal properties [76]. Synaptic properties provide the most obvious mechanism for synchronization, although non-synaptic mechanisms can play a role [30], particularly when neurons are firing spontaneously as is discussed in the section on high-frequency oscillations. Intrinsic neuronal properties determine cellular excitability and amplify feedback excitation, as is the case with voltage-gated  $\text{Ca}^{2+}$  currents in CA3 pyramidal cells in disinhibited hippocampal slices [77]. Intrinsic neuronal properties also can shape the morphology of epileptic discharges, for instance where voltage-gated  $\text{Ca}^{2+}$  currents and  $\text{Ca}^{2+}$ -dependent  $\text{K}^{+}$  currents generate rhythmic afterdischarges [76]. The importance of intrinsic neuronal properties is underlined by the many mutations in the genes for voltage-gated ion channels and their accessory subunits that have been associated with certain clinical epilepsies [5], and experimental evidence of “acquired channelopathies” following induction of epileptic foci in rodents [7, 54].

A great deal of progress was made on the cellular pathophysiology of acute epilepsy models in hippocampal and neocortical slices during the 1980s and 1990s. As I will outline in the next section, similar work on chronic models has developed rapidly since then. However, acute models *in vitro* continue to play important roles in epilepsy research. One example is the concept of clustering of neuronal firing during high-frequency oscillations, where subsets of neurons firing at lower rates combine to generate a collective high-frequency rhythm [8, 24, 35]. Another example comes from work on neocortical slices exposed to low concentrations of extracellular  $\text{Mg}^{2+}$ : bursts of rapid neuronal firing were spatially restricted and propagated relatively slowly across the slices. This study exploited particular advantages of brain slices *in vitro* for integrating optical and electrical recordings of neuronal activity, concluding that the rate of propagation was controlled by feed-forward or surround

inhibition which balanced excitatory synaptic outputs from the discharging neurons [78].

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## 15.3 Chronic Models and Hyperexcitability

Epilepsy is a chronic disease and the most realistic models also are chronic and characterized by spontaneous seizures. Again Phil Schwartzkroin made pioneering contributions, particularly with the alumina gel model and with human recordings *in vitro* [68]. The idea of using the precision of *in vitro* methods to investigate chronic models and clinical conditions is fundamentally important, particularly in epilepsy. I will outline some recent progress on chronic models, mainly of temporal lobe epilepsy.

### 15.3.1 Cellular Pathophysiology in Chronic Foci

Perhaps the most common methods to model medial temporal lobe epilepsy in rodents rely on inducing an initial status epilepticus, by injection of pilocarpine (usually systemically) [19], injection of kainic acid (systemic or intrahippocampal) [9, 85] or prolonged electrical stimulation [50, 80, 81]. Chapters on all these models can be found in a major monograph on epilepsy models which was co-edited by Phil Schwartzkroin [53]. Intrahippocampal tetanus toxin does not cause status epilepticus and causes little or no histopathology in the short term [32, 45], but can cause spatially limited hippocampal sclerosis in a minority of cases as well as a late loss of somatostatin-containing interneurons [47]. All these models have a “latent period” during which epileptogenesis transforms normal into epileptic brain tissue and results in spontaneous seizures, which are electrographically similar to those seen clinically.

One productive approach uses *in vivo* (or perhaps more accurately, *ex vivo*) brain slices to investigate cellular mechanisms of chronic epileptic

foci, both experimental and clinical [17, 29, 42, 68]. Such studies have revealed diverse cellular changes, usually several coexisting in specific kinds of epileptic foci. These changes include synaptic, intrinsic neuronal, glial and structural. The problem gets more complex because while some changes increase excitability and probably promote epileptic activity, others may be adaptive, reducing excitability and tending to control epileptic activity, and some may even be epiphenomena with no direct consequence for generation of epileptic seizures.

Many synaptic changes have been found in chronic epileptic foci, e.g. affecting transmitter release, receptor expression and synaptic modulation (reviewed in [16]). I will outline a few examples here. Several of these epilepsy-related changes affect GABA-ergic inhibition. Selective losses of inhibitory interneurons provide an attractive mechanism for parallel increases in excitability and propensity to seizures. Histopathological evidence has been variable and a review of the substantial clinical and experimental evidence is beyond the scope of this chapter. Several studies have shown losses of somatostatin containing interneurons in chronic models of temporal lobe epilepsy, both in the hilus of the dentate area [15, 47] and in CA1 [18, 22]. There also is evidence of loss of axon- and soma-targeting interneurons in clinical [21] and experimental material [41, 58] although this is less consistent and is complicated by changes in expression of markers, including parvalbumin, used to identify specific classes of interneurons [71] although evidence on soma-targeting interneurons is contradictory [22, 71]. Survival of normal numbers of interneurons does not necessarily mean that inhibitory function remains intact. Inhibition may be dysfunctional, as in the original dormant basket cell hypothesis or other conditions where excitation of interneurons is impaired [69–71, 82, 88]. Inhibition can also be weakened indirectly by changes in chloride homeostasis in the postsynaptic neurons, as shown in subicular slices from humans with medically intractable epilepsy undergoing surgical resection [27]. Synchronous interictal discharges in this clinical tissue have substantial contribu-

tions from depolarizing GABAergic synaptic potentials, which appear to be associated with decreased expression of the KCC2 chloride transporter which maintains chloride equilibria hyperpolarized to rest.

Excitatory synapses also can be altered in epileptic foci. Receptors subunits can be modified, as is the case with AMPA receptors in more than one chronic model [55, 56]. Aberrant expression of different classes of receptor can affect synaptic function, as in the expression of kainic acid receptors in dentate gyrus, which prolong EPSPs and strengthen synaptic integration [2]. It has long been known that seizures induce changes in synaptic connectivity in the brain. The prototypical case is mossy fibre sprouting in the dentate gyrus [74], raising the prospect of increased recurrent excitation through this glutamatergic pathway. Recent studies suggest that aberrant postsynaptic receptors make the new synapses particularly effective [2], although it looks as though their effects may be controlled by inhibitory mechanisms, at least in vitro [51]. Despite the robust connection between chronic temporal lobe epilepsy and sprouting, it turns out that preventing sprouting with rapamycin after an episode of status epilepticus fails to prevent the development of chronic epilepsy [14, 26]. This is one example of the importance of testing the functional implications of cellular changes identified in epileptic foci: even plausible phenomena like sprouting of excitatory synaptic connections are not necessarily responsible for epileptogenesis.

Several neuropeptides change in chronic epilepsy, as reviewed in Casillas-Espinosa et al. [16], often in directions that suggest they may act as endogenous anticonvulsants. These effects have attracted attention for translational research and is providing leads for potential innovative treatments for epilepsies that are refractory to currently available drugs [72]. Finally, intrinsic properties due to voltage-gated ion channels change in both genetic and acquired epilepsies, with examples for sodium, potassium, calcium and HCN channels amongst others; this topic is beyond the scope of this chapter, but is reviewed in Poolos & Johnston [54].

This short overview covers a small part of the diverse cellular pathologies and pathophysiological changes that have been found in chronic focal epilepsies. Even the most reproducible of models reveals multiple distinct cellular changes. Major questions remain on the roles each plays. It may be that in isolation none are sufficient to induce epileptic foci, but that several need to be present. It also is clear that some changes are antiepileptic and could provide leads for new treatments, as in the example of the work on peptides mentioned above. Of course some changes may be epiphenomena, perhaps induced by repeated seizures, but with no material impact on seizure susceptibility or generation. It is likely that different kinds of epileptic foci, both clinical and experimental, may have their individual combinations of cellular abnormalities. Finally, epilepsy is more than the seizures, with a range of comorbidities that can be detected between seizures [13], many of which will have underlying cellular mechanisms that may be identified by the kinds of investigations outlined above.

### 15.3.2 The Epileptogenic Zone, Hypersynchrony and High-Frequency Oscillations

Around one in three persons with epilepsy fail to gain adequate seizure control with currently available drugs, maybe even more for medial temporal lobe epilepsy [84]. Surgical resection of the tissue responsible for seizures can be remarkably effective as long as the correct tissue is removed [48]. The epileptogenic zone is defined as the area that is necessary and sufficient for resection to result in seizure freedom [57]. If the seizures stop after surgery then the resection must have been sufficient, but it is harder to be certain that all the resection was necessary. Presurgical work-up can include non-invasive imaging, scalp EEG, subdural and depth recordings. Here I will focus on the discovery that high-frequency oscillations may help define the epileptogenic zone in clinical and experimental foci [11].

High-frequency oscillations have frequencies greater than used to be recorded by routine EEG, and typically are considered as 80 Hz and above. Paper-based EEGs meant that high-frequency oscillations were missed in clinical electrophysiological investigations, but increasing computerization and improved amplifiers led to their discovery in clinical recordings during the 1990s [1, 23]. The early studies of high-frequency oscillations divided them between physiological ripples and pathophysiological “fast ripples”, separated by a boundary at around 200–250 Hz [11, 12]. Fast ripples have been associated with neuronal loss and hippocampal sclerosis [24, 73]. However in an experimental model lacking status epilepticus and with minimal or no neuronal loss we found that fast ripples (>250 Hz) were reliably associated with the primary focus [37], which supports the idea that electrographic markers can extend surgery into more difficult cases.

It is well established that timing of gamma oscillations depends on inhibitory synaptic transmission [4, 83]. Inhibitory mechanisms also are important in physiological ripples [87]. However it is more difficult to see how fast ripples at >250 Hz, and reaching >500 Hz, can depend on synaptic mechanisms. Fast ripples appear to represent synchronous firing of excitatory pyramidal and granule neurons [8, 10, 35]. These neurons do not fire as fast as the fast ripple oscillation; rather they fire every few cycles. In vitro studies suggest that excitatory neurons are weakly but significantly synchronized in small fluctuating groups extending over distances of a few hundred microns [35]. The potential mechanisms of synchronization on a millisecond timescale are limited. Perhaps the most plausible is that groups of neurons which are close to threshold synchronize through electrical field (sometimes called ephaptic) interactions [30, 33]: this effect is relatively weak under physiological conditions but weak fields are sufficient to entrain neurons which are firing spontaneously [20].

As mentioned above, removal of the epileptogenic zone is “necessary and sufficient” for seizure freedom. Determining how much tissue needs to be removed is a difficult challenge. An interesting approach to this important problem

comes from a clinical study which found that successful surgical outcome is associated with the proportion of tissue generating high-frequency oscillations that was resected [28].

It is increasingly clear that, while the distinction between ripples and fast ripples have been quite successful, the frequencies of oscillations are not sufficient to define their functional significance [34], either in terms of markers for epileptogenic tissue or in terms of contributions to seizure generation (e.g. by strengthening synaptic summation) [6]. The distinction between interictal pathological and normal physiological activities may depend on many factors: some cortical areas may differ from the hippocampus [43], not all epileptic foci are necessarily alike, and, from a practical point of view, electrode size can have an impact on recorded frequencies of high-frequency oscillations [86]. What does appear useful is finding recording sites with anomalous features, which may include faster activity than found in other sites in the same person [25, 38].

A distinctive approach to identifying epileptogenic zone used multichannel microelectrodes, the Utah or NeuroPort arrays, in people undergoing invasive ECoG recordings to find the regions in which neuronal hyperactivity first occurs at seizure onset. These arrays contain ~100 microelectrodes extending 1 mm from their bases which can be inserted into the cortex to record from neurons, probably located in layer 4–5. The big surprise was when Truccolo et al. [79] found that neurons recorded by their microelectrode arrays mostly stopped firing at electrographic (ECoG) seizure onset. This deviates substantially from experimental models of focal epilepsies, but does confirm earlier clinical studies with single microelectrodes [3]. Subsequently Schevon et al. [59] did find neuronal firing accelerating at seizure onset in some of their arrays, but the advancing wave of neuronal hyperactivity and hypersynchrony was much more restricted than the epileptic ECoG. It is not clear whether they were luckier or more careful in their microelectrode positioning, but it is reassuring that seizures are associated with accelerating neuronal firing which is phase linked to the simultaneously recorded epileptic ECoG [49]. The discrepancy

in localization was attributed to “inhibitory restraint” outlined above [59]: to recap, the idea is that feedforward inhibition constrains the advancing front of hyperactivity of excitatory neurons participating in the chain reaction of the epileptic seizure. This has been demonstrated explicitly in rodent neocortical slices *in vitro* [59, 78]. On this model the focal slowly-propagating population of hyperactive hypersynchronous excitatory neurons projects to more widespread regions where excitation is held in check by feedforward inhibition so that most of epileptic ECoG occurs in the absence of accelerating neuronal firing. This challenges the original description of epileptic EEGs as “hypersynchronous” [52], but it can be argued that the original use is a reasonable label for the large-amplitude relatively rhythmic EEG or ECoG, as long as it is clear that it does not necessarily mean hypersynchronous neuronal firing in the same area. These clinical investigations are difficult, but (ethics permitting) need repeating, ideally with critical experimental tests to determine whether the cellular interpretation of the role of feedforward inhibition derived from reductionist experiments really do apply to epileptic cortex in humans.

Perhaps the biggest challenges for this line of research are (a) whether the regions initiating hypersynchronous firing really do mark the epileptogenic zone, and if so, (b) how to exploit it for presurgical evaluation in preparation for resection of medically intractable epileptic foci. Microelectrode arrays would be very difficult to implement in most clinical settings: inserting microelectrode arrays into human cortex can be difficult and can damage the recorded tissue, while analyzing the resulting data needs the tools of cellular electrophysiology normally found in basic neuroscience laboratories. However it may be that other, technically more straightforward, markers can be found. Perhaps the most promising are the high-frequency oscillations discussed above. They represent coincident firing of principal neurons, and may provide markers for hyperactivity and hypersynchrony of neuronal firing. If they do, they would prove much more straightforward for clinical investigation than unit recordings from penetrating arrays of microelectrodes.

Future work needs to refine presurgical identification of the epileptogenic zone, using high-frequency oscillations and other biomarkers. The relationship between these biomarkers and pathological high-frequency neuronal firing detected by penetrating arrays may play a role in solving that clinical challenge. This relationship may also provide insights into the cellular mechanisms responsible for high-frequency oscillations, providing an *in vivo* approach to complement *ex vivo* (or *in vitro*) investigations to provide insights into the organization of the pathophysiological networks of epileptic foci.

## 15.4 Concluding Remarks

The last few decades have seen spectacular advances in our understanding of the cellular pathophysiology of epileptic activity in acute and chronic models and in clinical foci. Multiple cellular pathologies and pathophysiological processes operate in chronic foci, whether clinical or experimental. One set of major challenges is to distinguish between those responsible for generating seizures, helping control them, responsible for comorbidities, and functionally neutral epiphenomena. Progressive advances in chronic experimental investigations *in vivo* have started to help us understand epileptic foci *in situ*, and to refine our concepts of the epileptogenic zone which should expand the application of surgery in cases of pharmacologically intractable epilepsies.

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# Does Epilepsy Cause a Reversion to Immature Function?

# 16

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## Abstract

Seizures have variable effects on brain. Numerous studies have examined the consequences of seizures, in light of the way that these may alter the susceptibility of the brain to seizures, promote epileptogenesis, or functionally alter brain leading to seizure-related comorbidities. In many –but not all– situations, seizures shift brain function towards a more immature state, promoting the birth of newborn neurons, altering the dendritic structure and neuronal connectivity, or changing neurotransmitter signaling towards more immature patterns. These effects depend upon many factors, including the seizure type, age of seizure occurrence, sex, and brain region studied. Here we discuss some of these findings proposing that these seizure-induced immature features do not simply represent rejuvenation of the brain but rather a de-synchronization of the homeostatic mechanisms that were in place to maintain normal physiology, which may contribute to epileptogenesis or the cognitive comorbidities.

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## Keywords

GABA receptor • Chloride cotransporter • Neurogenesis • mTOR • Dysplasia • Epileptogenesis

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## 16.1 Introduction

Epilepsies have multiple causes and phenotypes, leading to different seizure and epilepsy syndromes. A variety of genetic, toxic/metabolic, or structural abnormalities have been causally associated with epilepsies. Epilepsy may occur as a “system disorder”, attributed to dysfunction – but no overt structural pathology – of specific neuronal networks, as typically occurs in genetic generalized epilepsies, like absence epilepsy [4]. In other cases, specific pathologies, e.g., cortical malformations or hippocampal sclerosis, may lead to the generation of an epileptogenic focus.

Seizures and epilepsies may disrupt brain development. Often, these maldevelopmental consequences of seizures may manifest as age-inappropriate reversal to immature functions and developmental processes. For example, seizures may trigger the aberrant re-emergence of immature features of GABA<sub>A</sub> receptor (GABA<sub>A</sub>R) signaling in neurons from adult animals or may cause morphological changes reminiscent of immature neurons. Immature features include the generation of new neuronal progenitor cells, functional alteration of selected signaling pathways or morphological changes. Many of these immature features have been documented in surgically resected epileptic tissues from individuals with drug-resistant epilepsies, like temporal lobe epilepsy (TLE), hypothalamic hamartomas, cortical dysplasias, or peritumoral epileptic tissue. Comparisons with nonepileptic post-mortem or surgically resected tissues have indicated that some of these changes are specific for the epileptic tissue [46, 47]. Yet, the appearance of these changes after seizures in animal models often depends upon a variety of factors. Here we will discuss the animal studies that have supported these observations and have provided insights on the complex interactions between the immature features of the epileptic focus and epilepsies, their etiologies and treatments and how these can be modified by age, sex, region-specific or other factors.

## 16.2 Neurogenesis in TLE

Perhaps the most classic argument for a reversal of normal age-specific functions with a re-emergence of patterns observed during development is the observation that there is an increased number of newborn cells in the dentate gyrus, in response to seizures or during the epileptic state [101]. Increased neurogenesis in the dentate gyrus of adult rats has been shown using post-SE models of epilepsy [48, 86, 100] or kindling [84, 105] or hyperthermic seizures [[61] and reviewed in [85, 101]] (Table 16.1). Newborn cells manifest many of the electrophysiological and morphological features of the granule cells, but also some distinctive characteristics. For example, they may be more dispersed [48, 100], have bipolar rather than polarized dendrites and they do not stain for Neuropeptide Y (NPY) or glutamic acid decarboxylase (GAD) immunoreactivity [100]. Furthermore, newborn cells may integrate abnormally into the hippocampus after seizures. Newborn neurons that migrate towards the CA3 pyramidal region may synchronize with CA3 neurons into epileptiform bursts [100]. Doublecortin-positive newborn neurons in the hilar dentate of epileptic rats exhibit long and recurrent basal dendrites directed towards the granule cell layer and also receive excitatory synaptic input which is unusual in seizure-naïve rats [95]. These seizure-induced changes may contribute to the excitability of the hippocampus. It has also been proposed that newborn neurons may not be capable to integrate normally in processes controlling cognitive processes, contributing therefore to cognitive deficits [30, 88].

The effects of seizures on neurogenesis at the dentate is age, sex, region, model specific and may depend on the number and type of seizures that the animal experiences (reviewed in Table 16.1). In brief, neonatal rats may respond instead with reduced or unaltered neurogenesis. Furthermore, aged rats may not respond as robustly with neurogenesis following seizures as younger adults do. Longitudinal studies may

**Table 16.1** Effects of seizures on neurogenesis in the dentate gyrus

Animal characteristics	Model of seizures	Effect on neurogenesis in the dentate gyrus	Reference
PN0-4 Sprague–Dawley rats	PN0-4 flurothyl seizures (brief, repetitive)	1-5 brief flurothyl seizures had no effect on neurogenesis 25 flurothyl seizures over 4 days <b>reduced</b> neurogenesis in the dentate	[69]
PN1-7 Wistar rats	Recurrent pilocarpine SE (PN1, PN4, PN7) BrdU (PN7, PN13, PN20, PN48)	<b>Reduced</b> neurogenesis on PN8, PN14 <b>Increased</b> neurogenesis on PN49	[124]
PN9 rats	PN9: kainic acid SE (2–3 h) BrdU: 3 h after kainic acid	<b>Reduced</b> neurogenesis in the superior blade of the dentate	[59]
PN6-20 Sprague–Dawley rats	1–3 episodes of kainic acid SE between PN6-20 BrdU after each seizure and 4 h prior to sacrifice	<b>Reduced</b> number of BrdU-positive neurons in rats with 3 SEs, assessed on PN13, PN20, PN30, but not at earlier timepoints	[63]
PN10	PN10: Hyperthermic seizures (<30 min)	Normothermia-exposed males had more BrdU-positive cells than females	[61]
Sprague–Dawley male, female rats	PN11-16: BrdU injections	Hyperthermia had <b>no acute effect</b> on neurogenesis (assessed at PN17) Following hyperthermic seizures, newborn neurons in males <b>survived better</b> till PN66 than in females	
PN15 Sprague–Dawley rats	PN15: flurothyl SE PN17: BrdU injection	<b>Increased</b> neurogenesis after SE Further increased in malnourished animals	[78]
PN21, PN35 Sprague–Dawley rats, both sexes	Lithium-pilocarpine SE BrdU 3th–6th day after SE	<b>Increased</b> neurogenesis in both age groups No association with cell loss or subsequent probability for epilepsy	[98]
Adult Sprague–Dawley rats	Pilocarpine-SE (3–5 h) BrdU: 1–27 days post-pilocarpine	<b>Increased</b> neurogenesis at 3, 6, and 13 days post-pilocarpine SE	[86]
Adult Sprague–Dawley rats	Pilocarpine or kainic acid induced SE (1 h) BrdU: 4–11 or 26–30 days post SE	Newborn neurons born after SE migrate into the CA3 layer, maintain many granule cell characteristics (electrophysiological, morphological). However, they are NPY or GAD negative, have bipolar dendrites, and integrate abnormally, firing synchronously to CA3 pyramidal neurons	[100]
Adult female mice (nestin-GFP transgenic mice)? (8 week old)	Kainic acid SE (2–3 h) BrdU: 8 days post SE	<b>Increased</b> neurogenesis post-SE seen with the doublecortin positive neurons but not with the nestin or calretinin positive neurons Increased dispersion of newborn cells was seen with both doublecortin and calretinin positive neurons after SE	[48]
Adult male Sprague–Dawley rats	Amygdala kindling BrdU: 1 day after last kindled seizure or stimulation	<b>Increased</b> neurogenesis after ≥9 stage 4–5 seizures but not after 4–6 seizures Neurogenesis may not play a role in kindling development	[84]

(continued)

**Table 16.1** (continued)

Animal characteristics	Model of seizures	Effect on neurogenesis in the dentate gyrus	Reference
Adult male Wistar rats	Amygdala kindling	<b>Increased</b> neurogenesis seen only at the BrdU late group (after stage 5 seizures)	[105]
	BrdU early group: on the 2nd–4th stimulation days BrdU late group: on the days of their 2nd–4th stage 5 seizure		
Adult C57BL/6J mice	Flurothyl kindling	<b>Increased</b> neurogenesis after: 1–3 days following 1 seizure 0–7 days after 8 seizures	[28, 29]
	BrdU injections 0–28 days after 1 or 8 flurothyl seizures		
Adult F344 rats (4 months old)	Kainic acid i.c.v. or graded kainic acid SE (<6 h) i.p.	16 days post-SE: <b>Increased</b> number of doublecortin positive neurons in the dentate 5 months after SE: <b>decreased</b> numbers of doublecortin positive neurons in the dentate	[40]
Adult F344 rats (12 month old)	Kainic acid SE (i.p.) BrdU: day 0–12 after SE	<b>Increased</b> neurogenesis in the dentate, but to a less degree than in younger rats	[106]

Seizures have age and model-specific effects on neurogenesis in the dentate gyrus

*BrdU* bromodeoxyuridine, *GAD* glutamic acid decarboxylase, *GFP* Green fluorescent protein, *i.c.v.* intracerebroventricular, *i.p.* intraperitoneal, *NPY* neuropeptide Y, *PN* postnatal day, *SE* status epilepticus

reveal time-dependent changes in neurogenesis, which may be influenced also by the ability of these newborn cells to survive. For example, hyperthermic seizures caused the newborn neurons to survive longer in males than in females till adulthood, suggesting sex-specific factors controlling their function [61]. Few brief seizures may not be as sufficient to affect neurogenesis, as frequent or prolonged seizures do.

Investigations into whether aberrant neurogenesis may contribute to epileptogenesis have yielded variable results. Administration of anti-mitotics that prevent neurogenesis may decrease the frequency of spontaneous seizures in post-SE animals [51]. However, other treatments that reduce seizure-induced neurogenesis have resulted in either reduction [109] or no effect [88] on the frequency of spontaneous seizures. The developmental studies on the effects of SE in 2–3 week old rats which show increased SE-induced neurogenesis, even though neither cell loss nor epileptogenesis always ensue have also failed to associate the increase seizure-

induced neurogenesis with either of these consequences of SE [98]. Seizure-induced neurogenesis appears therefore to contribute to the excitability of the epileptic hippocampus and possibly to the associated cognitive dysfunction, but there is no definite evidence that it is required for or mediates the ensuing epileptogenesis. Future research into deciphering the mechanisms leading to seizure-induced neurogenesis and how these are modified by age or sex or seizure-specific factors would be needed.

### 16.3 Evidence for Immaturity of GABA<sub>A</sub> Receptor (GABA<sub>A</sub>R) Signaling in Epilepsies

GABA<sub>A</sub>R signaling is well known to undergo structural and functional changes through development. The subunit composition of the GABA<sub>A</sub>R complexes changes to include subunits that will provide electrophysiologic and pharmacological properties more akin to mature neurons. A typical

example is the developmental shift from alpha 2 or 3 (GABRA2 or GABRA3) to alpha 1 (GABRA1) subunits, which attribute faster kinetics of the inhibitory post-synaptic currents (IPSCs) and higher sensitivity to benzodiazepines [17, 47, 60]. In addition, GABA<sub>A</sub>R signaling changes from depolarizing early in development to hyperpolarizing in more mature neurons, rendering GABA<sub>A</sub>R-mediated inhibition more effective in older animals [70]. This is thought to be due to the developmental shift in the balance of the activity of cation/Cl<sup>-</sup> cotransporters (CCCs) that control the intracellular Cl<sup>-</sup> concentration to favor cotransporters that maintain high intracellular Cl<sup>-</sup> (i.e., NKCC1) in immature neurons and low intracellular Cl<sup>-</sup> in mature neurons (i.e., KCC2) [6, 26, 33, 90, 97]. The developmental increase in the expression and activity of KCC2, a Cl<sup>-</sup> exporting transporter, and the parallel decrease in NKCC1 eventually reduce intracellular Cl<sup>-</sup>, permitting the appearance of hyperpolarizing GABA<sub>A</sub>R signaling in more mature neurons.

The presence of depolarizing GABA<sub>A</sub>R signaling is critical for normal development, as it promotes neuronal growth, differentiation and synaptogenesis, by controlling calcium-sensitive signaling processes. In parallel, KCC2 may also modify the development of glutamatergic synapses in dendritic spines via interactions with cytoskeletal proteins, like 4.1 N, independently of any effects on GABA<sub>A</sub>R regulation [62]. The absence of depolarizing GABA<sub>A</sub>R signaling early in life can either be incompatible with life or disrupt neuronal differentiation and communication [6, 16, 26, 33, 43, 118, 119]. Considering the neurotrophic effects of depolarizing GABA<sub>A</sub>R signaling, it is not entirely surprising that depolarizing GABA<sub>A</sub>Rs are also found in pathologic conditions that favor neuritic growth and differentiation so as to promote aberrant synaptogenesis, connectivity and re-wiring, as occurs in various forms of acquired, focal-onset epilepsies (Table 16.2). Indeed, depolarizing GABA<sub>A</sub>R signaling can be facilitated by neurotrophins, like brain-derived neurotrophic growth factor (BDNF), which are released after seizures [96].

Abnormal shifts in the CCC activity towards an NKCC1-dominant state or depolarizing

GABA<sub>A</sub>R signaling have also been found in a number of pathological conditions predisposing to or leading to epilepsy, like trauma [11, 74], ischemia [45, 83], anoxia/glucose deprivation [36] as well as after kindling [80, 96] or during the latent or epileptic state in post-status epilepticus (SE) rodent models of epilepsy [7, 12, 13, 22, 87] (Table 16.2). Under such pathological conditions, the role of GABA<sub>A</sub>R signaling is not just to promote the healing and re-wiring of the brain but may acquire a pathogenic role, by promoting neuronal excitability, due to the impairment in inhibition. In further support, KCC2 deficient mice manifest early life epilepsy and histopathologic alterations reminiscent of hippocampal sclerosis [122]. Pharmacologic inhibition of depolarizing GABA<sub>A</sub>R signaling using the NKCC1 inhibitor bumetanide in combination with GABA<sub>A</sub>R agonists has shown antiseizure effects in certain seizure models [18, 25, 65, 68, 75, 94, 103], although model-, region-, age-, or time-dependent differences have been reported [65, 66, 68, 117, 127]. Administration of bumetanide with phenobarbital prior to seizure onset in the kainic acid induced SE model significantly enhanced the antiseizure effect of phenobarbital, in an age-dependent manner, that was attributed to the developmental decrease in NKCC1 expression [25]. Similarly, bumetanide inhibited rapid kindling of PN11 Wistar rats when it was administered prior to kindling stimuli [68] or hypoxic seizures when given prior to hypoxia in PN10 rats, even though the brain levels of bumetanide are significantly low [18]. On the other hand, *in vitro* studies demonstrated variable results of bumetanide when given after seizure onset that followed model, age, and region dependent patterns [54, 117]. In addition, NKCC1-knockout mice show greater susceptibility to 4-aminopyridine than wild type animals [127]. It is therefore possible that bumetanide administration prior to seizure onset and younger ages may facilitate its ability to enhance the antiseizure effects of GABA<sub>A</sub>R agonists. However it is also evident that model and region specific factors or other competing mechanisms may modify its effect.

Bumetanide has also been proposed to alleviate the febrile seizure-induced neurogenesis [56] and

**Table 16.2** Epilepsies associated with depolarizing GABA<sub>A</sub>R signaling

Epilepsy type/animal model	Stage in epilepsy	Findings	Reference
<b>Human epilepsies</b>			
Human TLE	Following surgery for intractable epilepsy	<i>Subiculum</i> Depolarizing GABA <sub>A</sub> R signaling Bicuculline inhibits interictal epileptic discharges in vitro Higher probability for depolarizing GABA <sub>A</sub> R in KCC2-negative neurons	[19, 42]
		Microinjections of hippocampal/temporal lobe extracts in <i>Xenopus</i> oocytes yield depolarizing GABA <sub>A</sub> R and high NKCC1 and low KCC2 mRNA expression	[81]
		Lower probability for NKCC1 to colocalize with KCC2 in epileptic subiculum / CA1	[72]
Human epilepsy due to hypothalamic hamartomas	Following surgery for intractable epilepsy	Depolarizing GABA <sub>A</sub> R signaling in hypothalamic hamartomas	[55]
Human epilepsy due to cortical dysplasias	Following surgery for intractable epilepsy	Reduced KCC2 expression in focal cortical dysplasias <i>TSC, FCD type IIB</i> Increased NKCC1, reduced KCC2 <i>TSC (single case)</i> Depolarizing GABA <sub>A</sub> R signaling <i>FCD type IIA</i> Increased NKCC1 and KCC2 <i>FCD type I or II</i>	[107] [110]
		Abnormal developmental changes in the expression of NKCC1, KCC2 Increase in NKCC1, altered subcellular expression of KCC2 in cortical malformations (FCD type IIB, hemimegalencephaly, gangliogliomas)	[47] [3]
Tumor-associated human epilepsy	Peritumoral cells	Increased NKCC1 expression	[20]
<b>Animal models of epilepsy</b>			
Post-SE epileptic rats, pilocarpine model, male Wister rats	Latent phase, 3 weeks post-SE	Depolarizing GABA <sub>A</sub> R signaling in layer 5 entorhinal cortex but not in entorhinal layer 3, subiculum, dentate gyrus, or perirhinal cortex.	[13]
Post-SE epileptic rats, pilocarpine model, adult male Sprague–Dawley rats	Established epilepsy, 2–5 months after SE	Depolarizing GABA <sub>A</sub> R signaling in granule cells of the dentate gyrus, insular, subicular neurons or the deep layers of the piriform cortex Reduction of KCC2 expression in the dentate gyrus, subiculum or the deep layers of the piriform cortex	[7, 12, 22, 87]

Abnormal shift to depolarizing GABA<sub>A</sub>R signaling and/or expression of cation chloride cotransporters KCC2 and NKCC1 have been described in both human tissue derived from epileptogenic areas of patients with epilepsies, as well as in animal models of epilepsy

*FCD* Focal cortical dysplasia, *SE* status epilepticus

the post-SE epilepsy-associated behavioral deficits [14], but has not been shown to have anti-epileptogenic effects in post-SE epilepsy or an in vitro model [14, 75]. Depolarizing GABA<sub>A</sub>R signaling also renders the injured neurons dependent upon neurotrophic factors, like BDNF, for survival, by augmenting the expression of the pan-neurotrophin receptor p75<sup>NTR</sup> [108]. Neurons with depolarizing GABA signaling are therefore

more amenable to dying in injured areas, which are deprived of BDNF.

Epilepsy and seizures have also been associated with disruption in the normal developmental patterns of expression of the subunits of GABA<sub>A</sub>Rs (see Table 16.3). In certain – but not all – cases these reflect a return to a more immature type of GABA<sub>A</sub>R subunit composition, as in studies demonstrating a reduction in the  $\alpha$ 1

**Table 16.3** Abnormalities in GABA<sub>A</sub>R subunit expression in human epilepsies and animal models of SE or epilepsies

Epilepsy type/animal model	Stage in epilepsy	Findings	Reference
<b>Human epilepsies</b>			
Human TLE	Following surgery for intractable epilepsy	Decreased GABRA3 protein in temporal neocortex (layers I-III), no change in GABRA1 or GABRA2	[64]
		Increased GABRA3, GABRA5, GABRB1, GABRB2, GABRB3 mRNA in subiculum compared to neocortex	[82]
		Decreased GABRG2 mRNA in subiculum compared to neocortex	
		Decreased GABRA1, GABRA3, GABRB3, GABRG2 protein expression in sclerotic but not in nonsclerotic hippocampus (CA1)	[89]
		Increased GABRB1, GABRB2, GABRB3 protein expression in both sclerotic and nonsclerotic hippocampus	
Human mesial TLE	Following surgery for intractable epilepsy	No change in GABRA1, GABRB1, GABRB2 mRNA expression in the amygdala	[23]
Human epilepsy due to hypothalamic hamartomas	Following surgery for intractable epilepsy	No change in GABA <sub>A</sub> R subunit mRNA	[123]
Human epilepsy due to cortical dysplasias	Following surgery for intractable epilepsy	<i>TSC, FCD type IIB</i>	[110]
		Decreased GABRA1 protein	
		<i>FCD type IIA</i>	
		Decreased GABRA4 protein	
		<i>FCD type I or II</i>	[47]
		Abnormal developmental changes in the expression of GABRA1, GABRA4, GABRG2 protein	
<b>Animal models of SE or epilepsy</b>			
Post-SE, pilocarpine model, PN10 rats	Nonepileptic (adult)	<i>In dentate gyrus granule cells</i>	[126]
		Increase in GABRA1 mRNA No change in GABRA4, GABRD mRNA	
Post-SE rats, pilocarpine model, adult male Sprague–Dawley rats	1–8 days post-SE	<i>In CA1 pyramidal neurons</i>	[38]
		Decrease in GABRA4, GABRB2/3, GABRG2, gephyrin protein	

(continued)

**Table 16.3** (continued)

Epilepsy type/animal model	Stage in epilepsy	Findings	Reference
Post-SE rats, pilocarpine model, adult rats	1–5 months post-SE	<i>In dentate gyrus granule cells (hippocampus)</i> Decrease in GABRA1, GABRB1 mRNA Increase in GABRA4, GABRB3, GABRD, GABRE mRNA	[15]
Post-SE rats, kainic acid, adult male Sprague–Dawley	1 month post-SE	<i>In dorsal hippocampus</i> Increase in GABRA1, GABRA2, GABRA4, GABRB2, GABRB3, GABRG2 protein Decrease in GABRD	[104]
Post-SE rats, kainic acid, adult male Sprague–Dawley rats	7–30 days post SE	<i>In dorsal hippocampus</i> Decrease in GABRA5 and GABRD mRNA	[113]
Post-SE, Electrically induced, adult male Sprague–Dawley rats	7–30 days post SE	<i>In dorsal hippocampus</i> Increase in GABRA1, GABRA4, GABRB1, GABRB2, GABRB3 mRNA Decrease in GABRD mRNA	[76]
Post-SE rats, pilocarpine, adult male Sprague–Dawley rats	3–4 months post-SE	<i>In CA1, CA3 pyramidal neurons</i> Decrease in GABRA5 protein	[41]
Post-SE rats, electrical stimulation of the amygdala, adult male Sprague–Dawley rats	Epileptic rats	<i>In hippocampus</i> Increase in GABRB3 mRNA (all regions) Decrease in GABRA2 mRNA (CA3c) and GABRA4 mRNA (CA1)	[58]
Post-SE, pilocarpine model, adult Wistar rats	Epileptic rats	<i>In cerebral cortex</i> Decrease in GABRA1, GABRG3, GABRD mRNA Increase in GABRA5 mRNA	[67]
Post-SE, electrical stimulation of amygdala, adult female Sprague–Dawley rats	Epileptic rats	<i>In hippocampus</i> Phenobarbital non-responders are more likely to have reduced GABRA1, GABRB2/3, GABRG2 protein expression in the hippocampus than responders	[8]

SE and epilepsies have different effects upon the expression of GABA<sub>A</sub>R subunits (GABR). Their effects depend upon the type and/or model of SE or epilepsy, age at seizure occurrence, the region and timepoint after seizures when the study is conducted, and the specific subunit examined

subunit, whereas in others they indicate disrupted development [47]. Changes in GABA<sub>A</sub>R subunits may contribute to either drug refractoriness [15] or epileptogenesis [93] or the development of comorbidities.

Most of the above studies have been done in either adult animals or are derived from individuals with drug-resistant epilepsy that underwent surgical resection of the epileptogenic focus at ages when the brain is relatively more mature. Age-specific patterns of regulation by seizures have been extensively shown for the seizure-induced changes in GABA<sub>A</sub>R subunits [126].

Similarly, the effects of neonatal seizures on GABA<sub>A</sub>R signaling and CCCs are not only age [32, 53] but sex-specific as well [32]. Kainic acid induced SE in PN4–6 rats accelerated the switch to hyperpolarizing GABA<sub>A</sub>R signaling in the CA1 pyramidal neurons of males, due to an increase in KCC2 expression and decrease in NKCC1 activity [32]. In contrast, kainic acid induced SE in PN4–6 female rats, in which GABA<sub>A</sub>R signaling is not depolarizing, causes a transient return to the depolarizing signaling mode due to an increase in NKCC1 activity [32]. In this study, the sexually dimorphic response to



neonatal seizures seemed to depend upon the earlier maturation of GABA<sub>A</sub>R signaling in the female hippocampus, attributed to a higher expression of KCC2 and lower NKCC1 activity in females [32]. Sex differences in the expression of KCC2 and NKCC1 or GABA<sub>A</sub>R signaling in the hippocampus have also been confirmed in other studies [73, 79]. In addition, brief kainic acid seizures augment the activity of KCC2 shortly after induction of seizures in neonatal male rats [53]. It should be noted however that these studies relate to the postictal – acute or sub-acute – stages of neonatal SE. During the acute ictal phase of the SE, there is plenty of evidence to support that GABA<sub>A</sub>R signaling becomes depolarizing [25, 52].

The consequences of these seizure effects on the direction of GABA<sub>A</sub>R signaling could impact upon the subsequent susceptibility of the animal to seizures, affect its ability to stop seizure propagation, or alter cognitive abilities. For example, activation of GABA<sub>A</sub>R signaling in the anterior substantia nigra pars reticulata (SNR) in rats has important age and sex specific role in controlling seizure propagation in the flurothyl model [114, 115]. Exposure of male and female PN4-6 rats to kainic acid induced SE, at the time when GABA<sub>A</sub>R signaling is depolarizing, causes a precocious appearance of hyperpolarizing GABA<sub>A</sub>R signaling due to increase in KCC2 expression [35] and disrupts the GABA<sub>A</sub>R-sensitive anticonvulsant function of the anterior SNR in the flurothyl seizure model (unpublished data). It is possible that the early deprivation of the SNR of the neurotrophic effects of the depolarizing GABA<sub>A</sub>R signaling effects may impair its development, leading to these long-lasting deficits.

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## 16.4 Other Immature or Dysmature Features Associated with Epilepsies

Seizures may cause long-lasting changes in other signaling pathways involved in neurodevelopmental plasticity. The mTOR pathway has attracted a lot of research interest recently because it is central to cellular differentiation

and growth. The mTOR pathway may become dysregulated in several seizure and epilepsy models [34, 92, 111, 121, 125] even if not necessarily caused by genetic disruption of components of the mTOR pathway. The ability of rapamycin, an mTOR inhibitor, to suppress epilepsy in these models as well as prevent or reverse certain of the histopathological or cognitive abnormalities has supported its role as a potential epileptostatic and potentially disease-modifying treatment. We use the term “epileptostatic” (i.e., epilepsy is on hold) to indicate that inhibition of the expression of epilepsy and associated histopathological abnormalities occur only in the presence of mTOR inhibition but re-appear after the mTOR inhibitor is withdrawn. Other neurodevelopmental processes may also be affected, such as excitatory signaling or myelination. A neonatal brief kainic acid seizure may reduce the surface expression of the NMDA receptor (NR) subunit that normally emerges through developmental maturation, NR2A [21]. Seizures during the period of myelination can halt or impair myelination in both animal and human studies [24, 50, 91].

Loss of dendritic spines and less frequently shortening of dendritic length or abnormal dendritic branching patterns may be seen in patients with TLE or focal epilepsies [5, 10, 31, 44, 71, 102, 116]. Whether dendritic pathologies cause epilepsy is a matter open for investigation. Certainly many known etiologies of epilepsies demonstrate similar dendritic pathologies, including Rett syndrome [2] and tuberous sclerosis (TSC) [112] implicating the affected pathways (MeCP2, mTOR) in their pathogenesis. However the evidence that dendritic pathology causes epilepsy is currently lacking. Animal studies of seizures or epilepsy, in models like kindling, iron-induced cortical epilepsy, tetanus toxin model, or post-SE models of epilepsy have demonstrated similar dendritic abnormalities suggesting that seizures may impair dendritic architecture and spine development [1, 39, 49, 57, 77, 120, 125]. The lack of selectivity of the dendritic abnormalities for the epileptogenic focus, rather poses this feature as contributory to the overall neuronal dysfunction and seizure-associated comorbidities and to a lesser degree as causative of epilepsy.

In addition, dysplastic lesions may be encountered in pathological specimens from patients with TLE [9]. These can be found as clusters of granular neurons in layer 2 of the neocortex, nodular heterotopias in the temporal lobe, or heterotopic isolated neurons in the gray-white matter junction or deep subcortical white matter. It is currently unclear whether these dysplastic lesions are causative of or secondary to TLE. However, the possibility that such lesions may predispose to the development of TLE is supported by studies that demonstrate epileptogenic potential of these dysplastic lesions [27] as well as the animal studies demonstrating the pro-epileptogenic potential of pre-existing dysplastic lesions in two-hit seizure models [37, 99].

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## 16.5 Conclusions

Seizures and several pathologies predisposing to focal-onset epilepsies may trigger the re-acquisition of immature features in mature neurons that are integrated in the epileptogenic focus. The appearance of these immature features is influenced by age and sex-specific factors, at least for certain of the events that precipitate epilepsies. We propose that this untimely re-acquisition of the immature features is not equivalent to rejuvenation of the brain but may rather represent a de-synchronization of the homeostatic mechanisms that were in place to maintain normal physiology. In other words the maladaptive interactions and integration of these immature components with otherwise appropriately functioning brain regions may contribute to the increased excitability and underlying pathological changes seen in the epileptic focus. Furthermore, such effects may disrupt normal brain development, leading to long-lasting impairments in networks that are critical for either seizure control, like the SNR, or for information processing leading to cognitive dysfunction.

A number of important unresolved questions arise. Under which conditions does the untimely presence of immature features and functions in the seizure-exposed brain promote epileptogenesis or cognitive decline? Conversely, what are

the factors that can compensate and prevent disease progression? Are these functional changes different in epileptogenic foci than in regions that are secondarily affected by propagated seizures and why? What are the mechanisms leading to seizure-induced neurogenesis and how are these modified by age or sex or seizure-specific factors? Under which conditions might aberrant neurogenesis or abnormal GABA<sub>A</sub>R signaling have a pathogenic role in epileptogenesis or cognitive processes? What is the key switch mechanism that shifts depolarizing GABA<sub>A</sub>R signaling from promoting neurotrophic and healing processes in seizure-exposed or injured brains to facilitating excitability, seizure maintenance, and potentially epileptogenesis? Does the altered expression of GABA<sub>A</sub>R subunits in post-seizure or epileptic brain impair inhibition or could it, in certain situations, protect from the potentially excitatory effects of depolarizing GABA? It is evident from the examples presented in Tables 16.1, 16.2 and 16.3, that there is significant variability across studies, animal models, disease states, and regions suggesting that the answer may not be ubiquitous. Therefore, even if certain answers may be obtained in specific experimental paradigms, it is critical to be able to translate them into the human situation and, most specifically, to a specific individual in need of specific prognosis or treatment after a specific insult. Identifying markers that will enable us to detect and follow longitudinally, in vivo, the evolution of these changes and their functional alterations would be critical in both validating their significance and implementing individualized targeted treatments to prevent disease progression.

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# Are Alterations in Transmitter Receptor and Ion Channel Expression Responsible for Epilepsies?

17

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## Abstract

Neuronal voltage-gated ion channels and ligand-gated synaptic receptors play a critical role in maintaining the delicate balance between neuronal excitation and inhibition within neuronal networks in the brain. Changes in expression of voltage-gated ion channels, in particular sodium, hyperpolarization-activated cyclic nucleotide-gated (HCN) and calcium channels, and ligand-gated synaptic receptors, in particular GABA and glutamate receptors, have been reported in many types of both genetic and acquired epilepsies, in animal models and in humans. In this chapter we review these and discuss the potential pathogenic role they may play in the epilepsies.

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## Keywords

Genetic generalized epilepsy • Acquired epilepsy • Voltage-gated ion channels • Ligand-gated ion channels • Animal models of epilepsy

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## 17.1 Introduction

Neuronal voltage-gated ion channels and ligand-gated synaptic receptors play a critical role in maintaining the delicate balance between excitation and inhibition within neuronal networks in the brain that enables normal brain electrical function [10, 80]. The consequence of disturbing the expression or function of these, even relatively subtly, can render neuronal networks liable to fire in an inappropriate, hyper-synchronous, oscillatory manner which can be self-sustained, engage other neuronal networks, and result in a clinical epileptic seizure. Changes in expression of voltage-gated ion channels and/or ligand-gated synaptic receptors have been reported in many types of both genetic and acquired epilepsies. The causative relationship between these changes likely varies between different epilepsy syndromes. In some cases these changes are clearly causative of the epilepsy; in some they may represent susceptibility factors that render the brain more liable to become epileptic following a second insult (acquired or genetic); while in others they may be compensatory or even epiphenomena – related to the precipitating insult but not directly impacting on the epilepsy.

In this chapter we will outline some of the changes in expression of voltage-gated ion channels and ligand-gated synaptic receptors that have been reported in association with the development of both genetic and acquired epilepsies, in animals and humans, and discuss their potential pathogenic role.

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## 17.2 Genetic Generalized Epilepsy (GGE)

### 17.2.1 Overview

The genetic generalised epilepsies (GGEs) represent approximately 20–30 % of epilepsy cases and have a particularly high prevalence among children and adolescents [29, 31]. Patients with GGE have seizures that arise synchronously in both hemispheres on the electroencephalogram

(EEG) without any identifiable structural brain abnormality [3]. Patients with GGE syndromes can experience a number of different seizure types, including generalised tonic-clonic seizures, myoclonic seizures and absence seizures. The GGEs are a complex group of disorders with the aetiology presumed to largely genetic [3]. The underlying pathophysiological basis of the GGEs is still incompletely understood, but it is generally believed that in most cases more than one genetic abnormality contributes to determine the epilepsy phenotype (i.e. polygenic).

There are many reports in the literature describing mutations in voltage-gated sodium [18], potassium [30], calcium [104], HCN channels [76] and GABA receptors [104] in patients with GGEs. However, because the epileptogenic networks that generate the seizures in GGE are bilaterally and diffusely distributed, patients do not undergo epilepsy surgery and therefore brain tissue to examine for protein expression are rarely available. As a result most studies investigating changes in expression of ion channels and receptors relevant to GGE come from animal models. Of particular importance has been the Genetic Absence Epilepsy Rat from Strasbourg (GAERS) and Wistar Albino Glaxo/Rij-rat (WAG/Rij), which are the two most validated polygenic rat models of GGE [12, 52]. GAERS and WAG/Rij rats were both independently derived from selective inbreeding of Wistar rat colonies that spontaneously expressed generalised spike-and-wave discharges (SWDs) on EEG recordings, to generate strains that express frequent and prominent spontaneous absence-like seizures accompanied by generalised SWDs that electrophysiologically resemble those seen in human GGE with absence seizures. In both strains, the rats usually begin to display seizures in the second and third month post-natal, becoming longer and more frequent as the animals mature, being fully manifest in most WAG/Rij and in all GAERS by 4 months of age [12, 53]. It is well established and accepted that GAERS and the WAG/Rij strains are excellent models of human GGE with absence seizures because they parallel many of the features seen in humans with GGE such as; seizure, behaviour, electrophysiology,

pathophysiology and pharmacology (reviewed by [12, 15, 68]).

Many different ion channels and receptors, including GABA, glutamate, sodium, chloride, calcium and HCN channels, have been implicated in the pathogenesis of epilepsy. There are numerous reviews detailing mutations in these genes, however this review will focus on alterations in expression that occur in ion channels and receptors in rat polygenic models of GGE – GAERS and WAG/Rij.

## 17.2.2 Voltage-Gated Ion Channels

### 17.2.2.1 Hyperpolarization-Activated Cyclic Nucleotide-Gated (HCN) Channels

HCN channels generate  $I_h$  current in the brain, which modulates pacemaker activity and cellular excitability [63, 79]. All four HCN isoforms, HCN1-4, are expressed with regional and developmental differences in the brain and are differentially modulated by cAMP [7, 101].

Within the thalamocortical circuit, which is critical in the generation of seizures in patients with GGE, HCN1 channels are abundantly expressed in the cortex and HCN2 and HCN4 channels are abundantly expressed in the thalamus [57]. Several studies have examined HCN channel expression in GAERS and WAG/Rij rats reporting similar results. Using *in situ* hybridization, HCN1 channel mRNA expression is increased in distinct thalamic nuclei in epileptic GAERS, namely the reticular nucleus and ventroposterior medial nucleus of the thalamus, with no changes in expression in the somatosensory cortex or of HCN2 and HCN4 channels [41]. In adult WAG/Rij rats a reduction in HCN1 protein expression has been reported in the neocortex, hippocampus and cerebellum [89] whereas HCN1 channel protein expression is increased in the dorsal lateral geniculate nucleus of the thalamus [9, 32]. In another study, HCN1 protein expression was reported to be down regulated by 33 % in the neocortex in 1 month old WAG/Rij rats that were not yet experiencing spontaneous absence seizures. The decrease in HCN1 expression became

more progressive by three (56 % reduction) and 6 months (68 % reduction) [40].

### 17.2.2.2 Voltage-Gated Calcium Channels

Low voltage-activated, “T-type”, calcium channels are recognized to play a key role in neuronal burst firing in neurons in the thalamus which are critical in generating the hypersynchronous thalamocortical oscillations that underlie generalized SWDs [68]. Therefore alterations in expression of T-type channels have significant potential to play a pathogenic role in GGE. The three T-type calcium channel subtypes,  $Ca_v3.1$ ,  $Ca_v3.2$  and  $Ca_v3.3$ , have unique biophysical, pharmacological and regulatory properties [67] with differential expression within the thalamocortical circuit [73]. Mutations in the human *CACNA1H*, which encodes  $Ca_v3.2$ , have been found in patients with different GGE syndromes [11, 31, 45]. Exogenous expression of mutant human  $Ca_v3.2$  channels reveal a variety of biophysical changes [37, 64, 100]. Neurons from the thalamic reticular nucleus in GAERS, which plays a key role in regulation of the oscillatory thalamocortical network activity that underlies absence seizure-associated generalized SWDs in these animals, have been found to have a significant increase in T-type calcium currents compared to non-epileptic control rats [95]. In a developmental expression study in GAERS,  $Ca_v3.2$  mRNA expression was found to be elevated in the reticular nucleus of the thalamus in young animals before the onset of spontaneous absence seizures and  $Ca_v3.1$  and  $Ca_v3.2$  mRNA expression was increased in the ventral posterior thalamic relay nuclei and reticular nucleus of the thalamus of adult epileptic animals respectively [93]. Complimentary findings have been documented in WAG/Rij rats. mRNA expression of all T-type calcium channels were found to be elevated in distinct thalamic nuclei in young WAG/Rij rats (P18-28) preceding seizure onset;  $Ca_v3.1$  was shown to be increased in the lateral geniculate nucleus and centrolateral nucleus,  $Ca_v3.2$  was increased only in the reticular nucleus and  $Ca_v3.3$  was increased in the centrolateral nucleus and reticular nucleus [8].

### 17.2.2.3 Voltage-Gated Sodium Channels

Voltage-gated sodium channels are responsible for the initiation and propagation of action potentials. Functional channels consists of one  $\alpha$  subunit (Nav1.1–Nav1.9) and a variable number of  $\beta$  ( $\beta$ 1– $\beta$ 4) subunits. Subunits are composed of four domains. Each domain contains six transmembrane domains, as well as voltage sensor and pore forming domains.  $\beta$  subunits are smaller and contain one anchoring transmembrane domain and large extracellular domain [51]. In the somatosensory cortex of epileptic WAG/Rij animals, a significant upregulation of Nav1.1 and Nav1.6 mRNA and protein levels was reported with the changes being localised to layer II-IV cortical neurons with immunohistochemistry [39]. Interestingly, long term treatment of WAG/Rij rats with ethosuximide commencing prior to the onset of spontaneous absence seizures not only suppressed seizures but it also completely abolished the abnormal expression of Nav1.1, Nav1.6 and HCN1 when examined in 5 month old WAG/Rij rats [5].

## 17.2.3 Altered Expression of Ligand Gated Ion Channels

### 17.2.3.1 GABA Receptors

Fast responses to GABA are mediated by ligand-gated GABA<sub>A</sub> receptors whereas slow responses are mediated by G-protein coupled GABA<sub>B</sub> receptors. Homeostatic balance of GABAergic and glutamatergic neurotransmission is critical for the maintenance of neuronal excitability. In absence epilepsy, the neuronal hyperexcitability which underlies absence seizure generation in the thalamocortical circuit is hypothesised to be due to an imbalance between excitatory and inhibitory neurotransmission [15].

### 17.2.3.2 GABA<sub>A</sub> Receptors

GABA<sub>A</sub> receptors are pentamers consisting of multiple subunit subtypes, including  $\alpha$  ( $\alpha$ 1– $\alpha$ 2),  $\beta$  ( $\beta$ 1– $\beta$ 3),  $\gamma$  ( $\gamma$ 1– $\gamma$ 3),  $\delta$ ,  $\epsilon$ ,  $\pi$ ,  $\theta$ , and  $\sigma$  ( $\sigma$ 1– $\sigma$ 3) subunits. Properties of GABA<sub>A</sub> receptor strongly depend on subunit composition [25]. The most common subunit composition contains two  $\alpha$  subunits, two  $\beta$  subunits and a  $\gamma$  subunit [50]. Expression of  $\alpha$

and  $\beta$  subunits is sufficient for the production of GABA-gated chloride channels, while the  $\gamma$  subunit is required for modulation by benzodiazepines. Alterations in GABAergic inhibitory neurotransmission can influence neuronal excitability and indeed alterations in expression of GABA receptors have been reported in GAERS and WAG/Rij. A study by Spreafico et al. [86] found decreased immunofluorescence for  $\beta$ 2– $\beta$ 3 subunits of GABA<sub>A</sub> receptors in the sensorimotor cortex and anterior thalamic areas of epileptic GAERS [86]. However, in WAG/Rij rats conflicting results have been reported. An increase in the expression of  $\alpha$ 4 and  $\delta$  subunits of the GABA<sub>A</sub> receptor was observed in the relay nuclei of adult epileptic WAG/Rij animals [70] whereas decreased immunoreactivity of  $\alpha$ 3 subunit of the GABA<sub>A</sub> receptor was reported at inhibitory synapses in the reticular nucleus of the thalamus [47].

### 17.2.3.3 GABA<sub>B</sub> Receptors

The GABA<sub>B</sub> receptors are metabotropic transmembrane receptors that are linked to potassium channels via G-proteins, thus GABA<sub>B</sub> receptors mediate GABAergic slow responses. They are composed of two subunits; GABAB1 and GABAB2 with two splice variants of GABAB1 [34]. The GABAB1a and GABAB1b subunits are thought to be the site of agonist binding, while the GABAB2 subunit activates the G-protein signalling pathway. Alterations in GABA<sub>B</sub> receptor subunit expression and distribution have been reported in WAG/Rij rats with a marked reduction in GABA<sub>B1b</sub>, GABA<sub>B1ac</sub>, GABA<sub>B1d</sub> and GABA<sub>B1bc</sub> mRNA levels in the cortex, whereas GABA<sub>B1a</sub> and GABA<sub>B2</sub> mRNA levels were unchanged [55]. Alterations in GABA<sub>B</sub> receptor expression in the thalamocortical circuit has been reported in one study on GAERS [74]. GABA<sub>B1</sub> mRNA expression was shown to be increased in the somatosensory cortex but decreased in the ventrobasal nucleus of the thalamus. However, protein expression showed a different pattern of expression. Both GABA<sub>B1</sub> and GABA<sub>B2</sub> receptors were shown to be increased in all regions of the thalamocortical circuit (somatosensory cortex, ventrobasal nucleus and reticular nucleus of the thalamus) [74]. Moreover, transgenic mice overexpressing either GABAB1 subunits show an epileptic

phenotype characterized by spontaneous, recurrent atypical absence seizures [87, 103].

### 17.2.3.4 Ionotropic Glutamate Receptors

The ionotropic glutamate receptor (iGluR) family of excitatory synaptic receptors, are divided into four distinct subgroups based on their pharmacology and structural homology, including the AMPA receptors (GluA1–GluA4), kainate receptors (GluK1–GluK5), NMDA receptors (GluN1, GluN2A–GluN2D, GluN3A, and GluN3B), and  $\delta$  receptors (GluD1 and GluD2) [16]. The iGluRs are tetramers with a binding site for glutamate on each subunit that assemble as dimers of dimers, and their composition can be homomeric or heteromeric [94].

AMPA receptors mediate fast glutamatergic neurotransmission, and GluA1 and GluA2 protein expression has been shown to be upregulated in adult epileptic GAERS in the cortical membrane fraction [35]. In conjunction with this increase, it was also shown that stargazin ( $\gamma$ 2), a transmembrane AMPA receptor regulatory protein (TARP), was also increased specifically in the membrane of the somatosensory cortex. Juvenile pre-epileptic GAERS did not show any alterations in AMPA receptor or TARP expression [35]. The epileptic and ataxic phenotype of the *stargazer* mouse was found to be genetically determined by a mutation in the stargazin gene (*Cacng2*) resulting in decreased expression of stargazin in the brain [43]. WAG/Rij at 3 and 6 months of age show a reduction in the NMDA receptor GluN1 subunit and AMPA receptor GluA4 subunit immunoreactivity in the somatosensory cortex compared to control rats, which was especially evident in layers IV, V and VI [98]. Similarly, GluN2B protein expression has been shown to be decreased in layers III and V of the somatosensory cortex of 2 month and 6 month old WAG/Rij rats [33].

### 17.2.4 Metabotropic Glutamate Receptors

Metabotropic glutamate receptors (mGluR) constitute a family of eight G-protein-coupled receptor subtypes that can indirectly modulate

ion channels via second messenger systems. The family of mGluRs is composed of eight receptor subtypes, grouped into three different families according to their amino acid homology, pharmacologic properties, and G-protein coupling [13]. Class I metabotropic glutamate receptors (mGluR1 and mGluR5) mediate an increase in neuronal excitability and their activation can induce seizures. Class II receptors (mGluR2 and mGluR3) and class III receptors (mGluR4, mGluR6-8) depress synaptic transmission.

Several mGluRs subtypes are localised at synapses of thalamocortical neurons and thus may play an important role in the generation of epileptic generalised SWD [61]. Indeed, mGluR1 $\alpha$  subtype has been shown to be down regulated in the thalamus of 8 month old epileptic WAG/Rij rats but in young pre-symptomatic WAG/Rij rats this reduction was not observed indicating that this change in mGluR1 $\alpha$  receptor is occurring as a consequence of the seizures [60]. Additionally, mGluR4 protein levels in the reticular nucleus and ventral posterolateral thalamic nuclei were significantly reduced in 2 month old pre-epileptic WAG/Rij, but in 8 month old epileptic WAG/Rij rats a significant increase in mGluR4 protein levels in the reticular nucleus of the thalamus was observed [59].

## 17.3 Acquired Epilepsies

### 17.3.1 Overview

Acquired epilepsies are caused by brain insult such as stroke, traumatic brain injury (TBI), brain inflammation, or status epilepticus. Consequently, the molecular and cellular pathology of acquired epilepsies is heterogeneous both in type, distribution, and temporal evolution [71, 72]. Previous studies in human tissue and animal models have shown that in addition to neurodegeneration, neurogenesis, vascular injury and angiogenesis, proliferation and activations of different types of glia, axonal/myelin injury and axonal sprouting, dendritic plasticity, and changes in the composition of extracellular matrix, also the composition of ligand and voltage-gated ion channels can change [72]. This is often referred as “acquired

channelopathy” which can contribute to both epileptogenesis, evolution of comorbidities, and drug-refractoriness [77].

Previous global analyses of gene expression in epileptic tissue indicate that changes in the expression of mRNA, encoding for receptors and ion channels is not prominent. Some changes have, however, been observed in the level of mRNA for subunits of calcium channels or GABA receptors (e.g. [26, 49, 72, 102]). Moreover, the changes in mRNA levels often do not correlate with the protein level [1, 44, 75]. The level of the expressed protein is the key for the function of receptors or ion channels. Therefore, we focus on changes in protein expression of channels and receptors. We compare the findings in human tissue to that in animals undergoing epileptogenesis or already having established epilepsy (changes detected >7 days post-injury). The data are summarized in Table 17.1.

## 17.4 Altered Expression of Voltage-Gated Ion Channels

### 17.4.1 Voltage-Gated Sodium Channels

Our literature search did not reveal any data on expression of  $\alpha$  subunits in human TLE. In a rodent model of TLE, in which epileptogenesis was induced by status epilepticus (SE), Hargus et al. [28] reported that  $\text{Na}_v1.6$  was present in axon initiation segment and  $\text{Nav}1.2$  in the soma of neurons located in layer II of the entorhinal cortex. In animals with epilepsy, the expression of both subunits was increased at 3 months post-SE. Instead, expression of  $\text{Nav}1.1$  and  $\text{Nav}1.3$  was low and did not differ between the control and epileptic animals. Authors concluded that changes in the expression of  $\text{Nav}1.2$  and  $\text{Nav}1.6$  participate in generation of hyperexcitability of layer II neurons [28].

There are few studies of expression of  $\beta$  subunits in human TLE.  $\text{Nav}\beta3$  was found to be expressed in principal neurons of the hippocampus proper. In TLE patients without hippocampal

sclerosis, the expression of  $\text{Nav}\beta3$  was reduced in the hippocampus as compared to that in TLE patients with hippocampal sclerosis [99]. In the normal hippocampus,  $\text{Nav}\beta1$  subunit was expressed in neurons and a weak immunoreactivity was also observed in astrocytes. The astrocytic expression of  $\text{Nav}\beta1$  showed a remarkable increase during epileptogenesis and epilepsy triggered by SE [27].

### 17.4.2 Voltage Gated Potassium Channels

Voltage gated potassium channels are six transmembrane proteins containing a pore consisting of two transmembrane fragments and a voltage sensor on N-terminal side. Usually channels are tetramers composed of identical subunits. Their function is to return the membrane potential to resting state after depolarization [14].

One of the most studied subunits in human TLE and acquired epilepsy models is  $\text{Kv}4.2$  ( $\text{KCND}2$ ) that is critical for mediating the A-currents crucial for regulation of neuronal excitability and control of threshold for action potential initiation. In TLE patients with hippocampal sclerosis, the level of  $\text{Kv}4.2$  was increased in the somata of pyramidal cells and in activated astrocytes [1]. Immunoreactivity for its phosphorylated form,  $\text{pKv}4.2$ , was increased in granule cell and in molecular layers of the dentate gyrus as well as in the hippocampal CA3 principal cells. In areas of neurodegeneration, however, the dendritic immunoreactivity of  $\text{Kv}4.2$  or  $\text{pKv}4.2$  was reduced. In some pyramidal neurons  $\text{pKv}4.2$  co-localized with postsynaptic markers.

Decrease in the expression of  $\text{Kv}4.2$  in CA1 has also been observed in animal models of epilepsy, including epileptogenesis triggered by SE or by TBI [4, 56, 82]. Increased expression of  $\text{pKv}4.2$ , similar to that observed in human TLE, has been found in the CA1 after SE in rats [4]. These observations suggested that a decrease in  $\text{Kv}4.2$  and an increase in  $\text{pKv}4.2$  by ERK kinases can contribute to increased dendritic excitability, resulting in reduced seizure threshold after epileptogenic brain insults [1, 4].

**Table 17.1** Changes in the expression in subunits of various ligand and voltage-gated ion channels after epileptogenic brain insults in animal models and in human temporal lobe epilepsy (TLE)

Subunit	Model	Brain area	Change	Reference
<b>GABA receptors</b>				
<b>GABA-A<math>\alpha</math>1</b>	Human TLE	Hippocampus	↑ in granule cell layer ↓ in subgranular region, CA2 and CA3	[48]
		Hippocampal formation	↓ in CA1	[69]
	Intraperitoneal KA in rat	Hippocampus	↑ in DG molecular layer at 1 month ↓ in CA1-CA3	[85]
	FPI in rat	Hippocampus	↓ at 7 days	[75, 23]
	Intrahippocampal KA in mice	Hippocampus	↑ in DG at 1 month ↓ in sr and slm of CA1 and hilus at 1 month	[6]
<b>GABA-A<math>\alpha</math>2</b>	Human TLE	Hippocampus	↑ in the DG granule cell layer and molecular layer ↓ subgranular region	[48]
	Intraperitoneal KA in rat	Hippocampus	↑ in the DG molecular layer and sr and so of CA3 at 1 month	[85]
	Intrahippocampal KA in mice	Hippocampus	↓ in the DG, ↓ in CA1 sr and slm and CA3 at 1 month	[6]
	Pilocarpine in rat	Hippocampus	↓ in CA3 at 6 weeks	[20]
	Human TLE	Hippocampus	↓ subgranular layer ↑ in DG and subiculum ↓ in CA1	[48] [69]
<b>GABA-A<math>\alpha</math>3</b>	Intraperitoneal KA in rat	Hippocampus	↑ in CA1-CA3 in pyramidal-shaped perikaryon at 1 month	[85]
	Intrahippocampal KA mice	Hippocampus	↓ in CA1 sr and slm and CA3 at 1 month	[6]
	Pilocarpine in rat	Hippocampus	↑ in DG	[20]
			↓ in CA3, hilus	
			↑ in DG molecular layer at 30 days	[85]
<b>GABA-A<math>\alpha</math>4</b>	HC stimulation induced SE in rat	Hippocampus	↑ at inhibitory synapse	[91]
	Pilocarpine in mice	Dentate gyrus	↑ immunoreactivity at 14–60 days	[65]
	FPI in rat	Hippocampus	↓ at 7 days	[75]
	Intraperitoneal KA in rat	Hippocampus	↑ in DG interneurons ↓ in CA1 at 1 month	[85]
			↑ in DG	[6]
<b>GABA-A<math>\alpha</math>5</b>	Intrahippocampal KA in mice	Hippocampus	↓ in CA1 sr and slm and CA3 at 1 month	[20]
	Pilocarpine in rat	Hippocampus	↓ in CA3 ↑ in granule cell layer	[20]
				(continued)

Table 17.1 (continued)

Subunit	Model	Brain area	Change	Reference
<b>GABA-A<math>\beta</math>1</b>	Human TLE	Hippocampal formation	↑ in DG and subiculum	[69]
	Intraperitoneal KA in rat	Hippocampus	↑ in CA1-CA3 sr and so at 1 month	[85]
	Human TLE	Hippocampus	↑ granule cell layer ↓ subgranular region	[48]
<b>GABA-A<math>\beta</math>2</b>		Hippocampal formation	↑ in DG and subiculum	[69]
	Intraperitoneal KA in rat	Hippocampus	↑ in CA1 so and sr, ↑DG molecular layer at 1 month	[85]
	Pilocarpine in rat	Hippocampus	↑ in granule cell layer at 6 weeks ↓ in CA3 and hilus	[20]
<b>GABA-A<math>\beta</math>3</b>	Human TLE	Hippocampus	↑ granule cell layer ↓ subgranular region	[48]
		Hippocampal formation	↑ in DG and subiculum ↓ in CA1	[69]
	Pilocarpine in rat	Hippocampus	↑ in granule cell layer at 6 weeks ↓ in CA3 and hilus	[20]
<b>GABA-A<math>\gamma</math>1</b>	Intraperitoneal KA in rat	Hippocampus	↑ in DG molecular layer at 1 month ↓ in whole hippocampus	[85]
<b>GABA-A<math>\gamma</math>2</b>	Human TLE	Hippocampus	↓ subgranular region	[48]
		Hippocampal formation	↑ in DG and subiculum ↓ in CA1	[69]
<b>GABA-A<math>\delta</math></b>	Intraperitoneal KA in rat	Hippocampus	↑ in DG molecular layer and hilar interneurons at 1 month	[85]
	Intrahippocampal KA in mice	Hippocampus	↑ in DG ↓ in CA1 slm, CA3 and at 1 month	[6]
<b>GABA-A<math>\delta</math></b>	Pilocarpine in rat	Hippocampus	↑ in molecular layer at 6 weeks	[20]
	Pilocarpine in mice	Hippocampus	↑ translocation from synaptic to perisynaptic localization at 1 month	[105]
		Dentate gyrus	↑ immunoreactivity at 7–60 days	[65]
	CCI in rat	Hippocampus	↓ at 90 days	[36]
	FPI in rat	Hippocampus	↓ at 7 days	[75]
	Pilocarpine in mice	Dentate gyrus	↓ immunoreactivity at 7–60 days	[65]
		Hippocampus	↓ at symmetrical synapses at 1 month	[105]
	CCI in rat	Hippocampus	↑ at 3 months	[36]
	FPI in rat	Hippocampus	↓ at 7 days	[75]

<b>GABA-B receptors</b>		
<b>GABA-BR1</b>	Human TLE Intrahippocampal KA in mice Intrahippocampal KA in mice	Hippocampal formation Hippocampus Hippocampus
		↓ in CA1 and granule cells [58] ↑ in DG at 1 and 3 months [88] ↑ in DG at 1 and 3 months [88] ↓ n CA1
<b>AMPA/Kainate receptors</b>		
<b>GluR1</b>	Intraperitoneal KA in rat	Hippocampus
<b>GluR5</b>	Human TLE Intraperitoneal KA in rat	Hippocampus and cortex Hippocampus
		↓ in the hippocampus at 1 month [83] ↑ in the hippocampus [44] ↑ at 3 and 6 months [97]
<b>NMDA receptors</b>		
<b>NR1</b>	HC stimulation induced SE in rat	Hippocampus
<b>NR2B</b>	Human TLE CCI in rat Intraperitoneal KA in rat HC stimulation induced SE in rat	Hippocampus Hippocampus Hippocampus Hippocampus
		↓ at 1 month [19] ↑ [54] ↑ at 3 months [36] ↓ at 4–6 weeks [92] ↓ at 1 month with transient ↑ following seizures [19] ↑ in astrocytes at 1 month
	HC stimulation induced SE in rat FPI at P19	Hippocampus Hippocampus, cortex
		↑ at 3 months [54] ↓ at 7 days [24]
<b>Metabotropic glutamate receptors</b>		
<b>mGluR2/3</b>	Pilocarpine in rat Angular bundle stimulation induced SE in rat	Dentate gyrus Hippocampus
		↑ increase in molecular layer at 2 months [78] ↑ in reactive astrocytes up to 3 months [2]
<b>mGluR4</b>	Human TLE	Hippocampus
<b>mGluR5</b>	Intra-amygdalar KA in rat Angular bundle stimulation induced SE in rat	Hippocampus Hippocampus
		↑ in the dentate gyrus [46] ↑ in reactive non-neuronal cells at 2 months [96] ↑ in reactive astrocytes up to 3 months [2]
	Pilocarpine in rat	CA1
		↓ in stratum radiatum 4–10 weeks [38]
<b>Voltage-gated Na<sup>+</sup> channels</b>		
<b>Nav1.2</b>	HC stimulation induced SE in rat	Medial entorhinal cortex layer II
		↑ cell bodies at 3 months [28]
<b>Nav1.6</b>	HC stimulation induced SE in rat	Medial entorhinal cortex layer II
		↑ axon initial segment at 3 months [28]

(continued)



Table 17.1 (continued)

Subunit	Model	Brain area	Change	Reference
<b><math>\beta 1</math></b>	HC stimulation induced SE in rat		↑ in astrocytes at 7 days–3 months	[27]
<b><math>\beta 3</math></b>	Human TLE	Hippocampus, cortex	↓ in the hippocampus but not cortex of non-HS patients when compared to HS patients	[99]
<b>Voltage-gated K<sup>+</sup> channels</b>				
<b>Kv1.4</b>	Pilocarpine in rats	Hippocampus	↑ in the inner molecular layer of DG and sl of CA3 at 1–12 weeks	[56]
<b>Kv4.2</b>	Human TLE with HC sclerosis	Hippocampus	↑ in neuronal somata and astrocytes ↓ in neuropil	[1]
	Intraperitoneal KA in rat	CA1	↓ at 1 month	[82]
	Pilocarpine in rat	CA1	↑ in phosphorylation ↓ protein levels	[4]
		Hippocampus	↓ in sr of CA1 translocation to outer molecular layer of the dentate gyrus at 1–12 weeks	[56]
	Pilocarpine in rat	Hippocampus	↓ CA1 and CA3 at 50 days	[90]
	CCI in mice	Hippocampus	↓ in ipsilateral CA1 and CA3 at 1 and 8 weeks	[42]
<b>Kv4.3</b>	Pilocarpine in rat	Hippocampus	translocation to outer molecular layer at 1–12 weeks	[56]
<b>Kv7.2 (KCNQ2)</b>	HC stimulation induced SE in rat	amygdala	↑ in basolateral amygdala at 15 days	[66]

**Abbreviation:** CCI controlled cortical impact, FPI fluid percussion injury, HC hippocampus, HS hippocampal sclerosis, KA kainic acid, SE status epilepticus, sl stratum lucidum, sm stratum lacunosum moleculare, sr stratum oriens, sr stratum radiatum, TLE temporal lobe epilepsy

The two other subunits, Kv4.3 and Kv1.4, are implicated in A-current. In SE model, Kv4.3 protein translocated within the molecular layer of the dentate gyrus, resulting in an increase in its concentration in the outer two thirds of the molecular layer [56]. Kv1.4 is localized in axons. Interestingly, in epileptic animals an increase in Kv1.4 was observed in stratum lucidum of the CA3 and in the inner molecular layer of the dentate gyrus, which are the areas of extensive axonal sprouting after epileptogenic brain insults [56].

Another voltage dependent potassium channel implicated in acquired epilepsy is Kv7.2 (KCNQ2). Kv7.2 contributes to M-current, controlling baseline excitability. Number of Kv7.2 immuno-positive neurons was increased in the basolateral amygdala in animals after SE. The increase was present only in animals with spontaneous seizures. An increase in Kv7.2 was proposed to decrease the baseline excitability of amygdaloid neurons [66].

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## 17.5 Altered Expression of Ligand Gated Ion-Channels

### 17.5.1 GABA<sub>A</sub> Receptors

In drug-resistant patients with TLE, studies using subunit-specific antibodies have revealed profound and complex alterations in the expression of GABAA receptor subunits in the hippocampus. In particular, decrease in immunoreactivity of  $\alpha 1$ ,  $\alpha 2$ ,  $\alpha 3$ , and  $\gamma 2$  was observed in the CA1 in TLE patients with hippocampal sclerosis [21, 48]. This was probably related to the CA1 neurodegeneration. In the granule cells of dentate gyrus, however, the expression of  $\alpha 1$  and  $\alpha 2$  subunits was increased.

An increased expression of  $\beta 2$  and  $\beta 3$  subunits was observed in the granule cell layer of dentate gyrus of patients with TLE, while the data available on the expression of  $\beta$  subunits in the CA1-CA3 subfields of the hippocampus proper are conflicting [21, 48]. Interestingly, several  $\beta$  subunits increase their expression in the apical dendrites and decrease the expression in the basal dendrites of granule cells [48].

The literature reporting the changes in the expression of GABAA receptor subunits in animal models of acquired epilepsy is extensive (Table 17.1). In the normal rat hippocampus, the distribution of GABAA receptor  $\alpha$  subunits is topographically organized in different hippocampal subfields and layers [81]. Fritschy et al. [20] showed that at 6 weeks after SE in rats, the expression of  $\alpha 1$  subunit was up-regulated in the granule cell and molecular layers of the dentate gyrus and down-regulated in the hippocampus proper. Interestingly, also the number of hilar  $\alpha 1$  positive interneurons was reduced [20]. Accordingly, a decrease in  $\alpha 1$  immunoreactivity in the CA1 subfield was reported at 1 month after SE. The decrease was accompanied by an increase in  $\alpha 1$  immunoreactivity in the granule cell and molecular layers of the dentate gyrus TBI caused a decrease in  $\alpha 1$  expression in hippocampal extracts at 1 week post-TBI. No such decrease in  $\alpha 1$  expression was found when assessed at 90 days post-TBI [23, 36, 75].

SE in rats resulted in an increase in  $\alpha 2$  subunit immunoreactivity in the molecular layer of the dentate gyrus which was accompanied with a decreased  $\alpha 2$  expression in the CA1 [20, 85]. When kainic acid was injected directly into the hippocampus, a decrease in  $\alpha 2$  immunoreactivity occurred in the ipsilateral CA1, CA3, and also in the dentate gyrus [6]. After TBI,  $\alpha 2$  expression did not differ from that in controls [23, 75].

Immunoreactivity of GABAA receptor  $\alpha 3$  subunit was decreased in the CA1 and CA3 subfields of the hippocampus after SE [6, 20]. In the dentate gyrus, however,  $\alpha 3$  immunoreactivity was increased in rats with epilepsy [20].

Decrease in  $\alpha 4$  subunits was observed in extracts from the whole rat hippocampus 1 week after TBI, but no changes was evident in rats 90 days after TBI nor in rats in which epilepsy was induced by SE [36, 75, 91].

Immunoreactivity of  $\alpha 4$  subunit was increased in the molecular layer of the dentate gyrus at 30 days after SE in mice [65]. Moreover, Sun et al. [91] demonstrated that in epileptic animals the  $\alpha 4$  subunits located on the somata and dendrites of the dentate granule cells were more commonly present within inhibitory synapses

than extra-synaptically. This coincided with a diminished action of neurosteroids on synaptic current, possibly contributing to facilitation of seizures in epileptic animals [91].

Expression of  $\alpha 5$  subunit of GABA<sub>A</sub> receptor decreased in the CA1 subfield of the hippocampus in rats that had experienced SE [20, 85]. Moreover, SE resulted in a slight increase in  $\alpha 5$  subunit immunoreactivity in the dentate gyrus [20]. An increased expression of  $\alpha 5$  in the dentate gyrus and a decrease in the CA1 were also observed at 1 month after SE [6]. However, no changes were observed in the expression of  $\alpha 5$  expression at 7 days after TBI induced by FPI [23, 75].

In the normal rat brain, GABA<sub>A</sub> receptor  $\beta 1$  and  $\beta 3$  subunits are expressed in the dendritic areas of the hippocampus, including the stratum oriens and stratum radiatum of the CA1-CA3, and the molecular layer of the dentate gyrus. Staining for  $\beta 2$  subunit is light in pyramidal cell dendrites or in granule cells, but is present in hippocampal interneurons [84]. At 6 weeks after SE, immunoreactivity for  $\beta 2$  and  $\beta 3$  subunits was increased in the granule cell layer and decreased in the CA1-CA3 subfields of hippocampus proper as well as in the hilus of the dentate gyrus [20]. However, at 7 days post-TBI the hippocampal expression of  $\beta 3$  subunit remained unaltered [23].

Immunoreactivity of GABA<sub>A</sub> receptor  $\gamma 1$  and  $\gamma 2$  subunits is light in the normal hippocampus.  $\gamma 1$  is expressed in astrocyte-like profiles.  $\gamma 2$  subunit is highly expressed in the dendrites of CA1-CA3 neurons and in the molecular layer of the dentate gyrus as well as in perikarya of a subpopulation of hilar neurons. Expression of  $\gamma 3$  subunit is most remarkable in fibers [81]. After SE there is an increase in the immunoreactivity for  $\gamma 2$  subunit in the molecular layer of the dentate gyrus [6, 20, 65]. Zhang et al. [105] showed that in epileptic rats  $\gamma 2$  subunits are translocated to the perisynaptic location in the dendrites of granule cells. This resulted in a decrease in the expression of  $\gamma 2$  subunits at the synaptic region, and coincided with a decrease in phasic inhibition in the dendrites of granule cells [105]. Data on expression of  $\gamma 2$  subunit expression in the hippocampus proper are less consistent. After SE, Sperk et al. [85] observed an increase in  $\gamma 2$

immunoreactivity in stratum lacunosum moleculare and stratum radiatum of the CA3. However,  $\gamma 2$  immunoreactivity was decreased in the ipsilateral CA1 and CA3 [85]. A decrease in the hippocampal expression of  $\gamma 2$  subunit was observed also after TBI using Western blot [36, 75].

In the normal hippocampus,  $\delta$  subunits are expressed in the molecular and granule cell layers and in interneurons of the dentate gyrus. Light immunoreactivity is also present in the CA1-CA3 subfields of hippocampus proper [81]. Chronically epileptic animals after SE showed a decrease in  $\delta$  subunit immunoreactivity in the molecular layer of the dentate gyrus and an increase in interneurons. This was accompanied by an increase in excitability in hippocampal slices sectioned from epileptic animal [65]. As shown by Zhang et al. [105] the expression of  $\delta$  subunit was decreased in the dendrites of dentate granule cells [105]. Unexpectedly, no impairment was observed in tonic inhibition, indicating that a reduction in the expression of  $\delta$  subunit is compensated by other GABA<sub>A</sub> subunits [105]. In addition to SE models, a decrease in  $\delta$  subunit was observed in the hippocampal extracts at 7 days following TBI [75].

In summary, the changes in the pattern of expression of different GABA<sub>A</sub> receptor subunits are complex and model specific. In several reports, the decrease in the expression of subunit protein correlated with the severity of neurodegeneration whereas the increases in the expression likely presented compensatory molecular plasticity in altered network. Undoubtedly, the reported alterations explain the impairment of GABAergic transmission tuning the network towards increased excitability.

### 17.5.2 GABA<sub>B</sub> Receptors

Contribution of the altered expression of metabotropic GABA<sub>B</sub> receptors to acquired epileptogenesis and ictogenesis are poorly understood as compared to that of GABA<sub>A</sub> receptors. In the normal human brain, neuronal expression of one of the GABA<sub>B</sub> receptors, GABABR1, has been reported in the hippocampus and entorhinal

cortex. In TLE, the expression of GABABR1 was reduced in the dentate granule cells as well as in the hippocampus, particularly in areas of neurodegeneration. Interestingly, no compensatory change in the expression of GABABR1 was found in surviving neurons [58].

Straessle et al. [88] investigated the distribution of the two variants of GABABR1 receptor, GABABR1a and GABABR1b as well as GABABR2 in a mouse model of TLE. At 4–6 weeks or 3 month post-SE, ipsilateral CA1-CA3 showed a remarkable reduction in GABABR1a and GABA-BR1b as well as in GABABR2 immunoreactivities, which was associated with extensive hippocampal neurodegeneration. On the contrary, expression of GABABR1a, GABABR1b, and GABAR2 subunits was enhanced in the dentate granule cells. Moreover, temporary loss and then reappearance of interneurons stained for GABABR1a,b or GABABR1b was observed in the hilus and CA3. In contrast to GABAA receptor subunits, no changes GABABR1a, GABABR1b, or GABABR2 immunoreactivities were observed in the molecular layer of the dentate gyrus [88].

### 17.5.3 AMPA/Kainate Receptors

Studies investigating the expression of subunit proteins forming AMPA (GluR1-4) and KA (GluR5-7) receptors in epileptic tissue are meager, despite the fact that some of the non-NMDA glutamate receptors are targeted by antiepileptic drugs.

The expression of GluR1 and GluR2/3 was increased in the molecular layer and GluR2 also in the stratum radiatum, in TLE patients either with or without hippocampal sclerosis. An increase in GluR1 was also found in the CA3 principal cells as well as in hilar mossy cells [17, 54]. In hippocampal stimulation model of TLE, rats with epilepsy showed increased expression of GluR1 in the molecular layer of the dentate gyrus [54]. In kainate model, however, hippocampal expression of GluR1 expression was decreased at 1 month post-SE [83]. A decrease in GluR1 protein expression was also observed in the hippocampus at 3 months after TBI [36].

Some information is available on kainate receptor subunit GluR5. Li et al. [44] reported an increase in GluR5 protein level in the hippocampus, but not in the temporal neocortex of TLE patients [44]. An increase in GluR5, but not in GluR6 protein expression was also observed in rats at 3 or 6 months after SE in rats [97].

### 17.5.4 NMDA Receptors

NMDA receptors are implicated in synaptic plasticity, including LTP and LTD. This has created an interest whether they could play a role also in the development of aberrant synaptic plasticity found in animal models and human TLE.

NMDA receptors are tetramers consisting of at least one NR1 subunit and NR2(A-D) or NR3 (A-B) subunits. Properties of NMDA receptor are determined by its subunit composition [22]. Changes in the expression of NMDA receptors have been studied mostly at mRNA level, and these studies have focused on early time points after epileptogenic insult [22]. Much less information is available on protein expression and on its localization in the epileptic tissue.

As NR1 subunit is an indispensable component of the NMDA receptor, its expression provides information on the presence and localization of all NMDA receptors. Frasca et al. [19] reported a decrease in the phosphorylated and non-phosphorylated forms of NR1 in animals with epilepsy after SE [19].

More information is available on NR2, a subunit that is critical for the localization of NMDA receptor. After SE, the expression of NR2B protein was decreased which was accompanied by a decrease in PSD-95 protein. Moreover, the decrease in NR2 correlated with behavioral deficits [92]. In another SE model, hippocampus showed a reduced expression of both NR2B as well as its phosphorylated form, p-NR2B. The decrease in p-NR2B in post-synaptic membranes was associated with its reduced interaction with postsynaptic density. Interestingly, spontaneous seizures in these animals caused a transient increase in p-NR2B. It was concluded that altered phosphorylation on NR2B leads to extra synaptic

localization of NMDA receptors in epileptic animals [19]. A decrease in hippocampal NR2B protein was also found in the CCI model of TBI in rats [36].

### 17.5.5 Metabotropic Glutamate Receptors

To our knowledge, only mGluR4 protein expression has been studied in the human epileptic brain. In the normal human brain, almost no mGluR4 immunoreactivity was present. The hippocampus resected from patients with TLE, however, showed a strong mGluR4 immunoreactivity, particularly in the dentate gyrus. Interestingly, mGluR4 was localized in periphery of pre- and postsynaptic membranes [46].

mGluR5, a member of class I receptors has been studied only in animal models of epilepsy. In the normal hippocampus, mGluR5 protein is expressed in the dendritic fields of pyramidal cells [96]. After SE, the expression of mGluR5 was reduced in the ipsilateral hippocampus, which correlated with the severity of neurodegeneration in CA1-CA3 pyramidal cells [96]. mGluR5 immunoreactivity was decreased also in CA1 following SE [38]. The decrease in mGluR5 immunoreactivity after SE occurred in neurons whereas astrocytes showed a strong immunolabeling [2, 96]. It was suggested that an increase in mGluR5 in astrocytes could associate with Ca<sup>2+</sup> oscillations in astrocytes [2, 96] whereas a reduction in mGluR5 in the CA1 principal cells could associate with an impairment in LTD which is one of the post-SE functional consequences [38].

Similarly to mGluR5, expression of mGluR2/3 receptor proteins has been studied in models of TLE triggered by SE. When mGluR2/3 immunoreactivity was analyzed at 1 week, 3 weeks, or 3 months after SE, its expression was increased in activated vimentin-positive astrocytes. It was proposed that this contributed to the propagation of calcium waves in astrocytic syncytium, resulting in generation of seizure focus [2]. Interestingly, at the 3 month time point the intensity of mGluR5 staining was reduced in the molecular layer and in stratum lacunosum moleculare, and these

changes coincided with neurodegeneration in the entorhinal cortex [2]. In another SE model, a decrease in mGluR2/3 immunoreactivity was detected in the stratum lacunosum moleculare of the CA1 and CA3 and in mossy fibers located in the CA3 and the hilus [62]. An increase in mGluR2/3 immunoreactivity was found in the molecular layer of the dentate gyrus [62, 78]. It remains to be studied whether the increase in mGluR2/3 also after SE occurs in astrocytes [78].

## 17.6 Conclusions

The spectrum of changes in expression of proteins forming ligand and voltage-gated ion channels in genetic and acquired epilepsies is wide and extends over different etiologies. Changes vary depending on the stage of epileptogenesis, brain area investigated, as well as the cell type and cellular compartment assessed. The overall picture of changes in receptors and ion channels is fragmentary and their functional analysis is limited. However, considering the multiplicity of molecular and cellular changes present in the epileptogenic regions, it remains a viable hypothesis that acquired channelopathies form a specific component of the molecular fingerprint for epileptogenesis, eventually leading to the development of epilepsy. To which extent they also contribute to the development of comorbidities and/or tissue recovery remains to be studied.

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## Part III

### Models and Methods

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# How Do We Make Models That Are Useful in Understanding Partial Epilepsies?

# 18

David A. Prince

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## Abstract

The goals of constructing epilepsy models are (1) to develop approaches to prophylaxis of epileptogenesis following cortical injury; (2) to devise selective treatments for established epilepsies based on underlying pathophysiological mechanisms; and (3) use of a disease (epilepsy) model to explore brain molecular, cellular and circuit properties. Modeling a particular epilepsy syndrome requires detailed knowledge of key clinical phenomenology and results of human experiments that can be addressed in critically designed laboratory protocols. Contributions to understanding mechanisms and treatment of neurological disorders has often come from research not focused on a specific disease-relevant issue. Much of the foundation for current research in epilepsy falls into this category. Too strict a definition of the relevance of an experimental model to progress in preventing or curing epilepsy may, in the long run, slow progress. Inadequate exploration of the experimental target and basic laboratory results in a given model can lead to a failed effort and false negative or positive results. Models should be chosen based on the specific issues to be addressed rather than on convenience of use. Multiple variables including maturational age, species and strain, lesion type, severity and location, latency from injury to experiment and genetic background will affect results. A number of key issues in clinical and basic research in partial epilepsies remain to be addressed including the mechanisms active during the latent period following injury, susceptibility factors that predispose to epileptogenesis, injury – induced adaptive versus maladaptive changes, mechanisms of pharmaco-resistance and strategies to deal with multiple pathophysiological processes occurring in parallel.

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**Keywords**

Posttraumatic • Prophylaxis • Mechanisms • Latent period • Pathophysiology  
• Translation • Maladaptive

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## 18.1 Introduction

I have chosen to limit discussion here to models of the partial or lesional epilepsies; however a number of the issues are generic to understanding the relevance of other models to prevention and treatment of clinical epilepsies. What follows is not meant to be a literature review, but rather a discussion of unsolved issues and my opinions relevant to the use of epilepsy models. For additional discussion and references, the reader is referred to *Epilepsia*, 54: Supplement 4, 1–74, 2013 and articles therein, generated by participants of a joint AES/ILAE translational workshop, and a number of recent reviews of epilepsy models and mechanisms [10, 11, 24, 25, 28–30, 32, 37, 44–48]. Issues and difficulties raised by these authors bear a remarkable resemblance to those highlighted in reviews more than 20 years ago (e.g. [9]), in spite of the introduction of a number of new models and antiepileptic drugs.

Making laboratory models “relevant” requires several considerations. We should recognize that important contributions to understanding the mechanisms and treatment of neurological disorders has often come from “non-targeted” research, not seemingly focused on a specific disease-relevant issue. Much of the foundation for current research in epilepsy falls into this category. For this reason, too strict a definition of whether an experimental model is relevant or non-relevant to progress in preventing or curing epilepsy may, in the long run, slow progress. How should one design a model relevant to our clinical understanding and treatment? The first step would be identification of specific key clinical issues that are roadblocks in preventing or treating epilepsy, and would be feasible to address in a critically- designed animal model. Such issues can only be identified through detailed observations of clinical phenomenology

and associated human research data, i.e. an important “bedside to bench” approach. Extensive basic research focused on one or more of these key issues should follow. The third step would use of data from the model to design a clinical experiment or trial. Here is where further definition of the too-often-used term “translational” becomes important. In literature, scholarly translation of a work requires intimate knowledge of the vocabularies and nuances of two languages. By analogy, application of data from a laboratory model to aspects of clinical disorder requires detailed clinical and basic experimental data. Inadequate definition of the experimental target and less than rigorous exploration of the laboratory results in a given model can lead to a failed effort, or the “Lost in Translation” phenomenon.

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## 18.2 Why Model at All? What Are the Long-Term Goals?

### 18.2.1 Prophylaxis of Seizure Development in Lesional Epilepsies

Reference to any classification of seizure disorders clearly reveals that the epilepsies are multifaceted and related to a large variety of etiologies. Why should one expect that the same model of epilepsy will be useful for research on prevention of seizures resulting from a stroke versus a focal tumor versus a traumatic brain injury? Although each of these etiologies likely has a different *combination* of underlying mechanisms that lead to seizure generation, they may all involve some *common abnormalities* that are sequellae of focal injury, such as aberrant rewiring of cortical circuits or vulnerability of specific inhibitory interneuronal subtypes. Preventative treatments that are selective for such specific subtypes of underlying pathophysiology might be effective in more

than one epilepsy syndrome and least likely to induce unwanted side effects. However, there are multiple mechanisms for epileptogenesis that occur in parallel for each subtype of lesional epilepsy. As a consequence, too focused an approach on one pathophysiology or pathway or gene may yield a false negative result, even though the target mechanism has been successfully affected. This may be particularly true in a model of epilepsy with a very high yield of seizures from diffuse brain lesions, such as some post-status epilepticus models. Therefore, progress may require a roadmap of pathophysiological mechanisms obtained from models of different types of epilepsies and even different models of a specific post-lesional epilepsy (e.g. [17]). Potential use of anti-epileptogenic cocktails containing more than one selective agent, treatments with single drugs that have multiple modes of action or treatments directed upstream to affect multiple pathways for epileptogenesis [11] would be a logical direction in studies of prophylaxis after injury.

### 18.2.2 Development of Selective Treatments for Established Epilepsies, Based on Underlying Pathophysiological Mechanisms

Unfortunately, as noted by many authors, in spite of the development of a number of new antiepileptic drugs, the proportion of individuals who have poorly controlled seizures remains the same, at about one third. It has been proposed that the reason for this is the use of the same models for initial drug screening over the years. There are a number of unknowns that should be considered in designing models for experiments to address this goal. The species and strain of the animal model selected for a given experiment will significantly affect the results [22, 43]. Susceptibility to seizures, and the efficacy and spectrum of toxicity of a given antiepileptic drug will also vary in individuals with different genetic backgrounds [15, 39]. Do such genetic differences extend also to the specific mechanisms underlying development of a particular epilepsy syndrome due to different etiologies, e.g., limbic

circuit epilepsy due to a head injury vs. following status epilepticus? Might these two etiologies for the same syndrome differ in their responses to a particular anti-seizure agent? Another important variable may be the temporal evolution of epileptogenesis after serious brain injury. This clearly varies markedly among individuals and may unfold over years [38, 41]. Is ongoing seizure activity responsible for the progressive loss of hippocampal volume seen in radiological studies? Do the mechanisms underlying seizures following an injury also vary over time, so that drugs might be selected on the basis of the duration of epilepsy in a particular model or patient? For example, early on after cortical injury, treatments that are directed against alterations in blood brain barrier, and immunological mechanisms or inflammation may be effective, however underlying mechanisms may shift over time so that later, formation of new synapses, recurrent excitatory circuits or disturbed inhibitory circuit function become important drug targets. A related question is whether emergence of drug resistance is in part due to shifts in underlying epileptogenic mechanisms over time? Are decreases in responsiveness related to progressive changes late after injury, such as increasing excitatory sprouting and/or death of neuronal subtypes, and what is the role the plastic changes in cortical circuits resulting from ongoing epileptiform activity in this process?

### 18.2.3 Disease as a Tool to Explore Brain Molecular, Cellular and Circuit Properties

*“Epilepsy represents one of the most exquisite experiments of nature and its study may provide basic insight into fundamental functions of the brain.”* [20]. Epilepsy has long been used as a research tool to explore brain mechanisms such as circuit properties and connections, and mechanisms of synchronization within normal brain. Clementi [3] described reflex epilepsies in which a selective afferent input would trigger local seizure activity and could be used to assess connectivity. Much of the early information about

localization of sensory and motor functions in cerebral cortex was derived from experiments in which the sites for seizure activity were mapped in human brain (e.g. [33]). Epilepsy research has revealed normal brain mechanisms such as cortical “surround” inhibition [35] and aspects of dendritic function [31, 53]. Plastic changes in brain structure and function are key to many normal processes during development, as well as after injury [19]. Epileptogenesis is a striking example of such brain plasticity [18]. Issues such as sprouting of new connections, changes in receptor subunit composition, and alterations in intrinsic membrane properties that are characteristic of neural development are also found during epileptogenesis and following prolonged recurrent seizure activity. Clinical studies done with multiple implanted electrodes in patients with epilepsy, together with functional MRI have revealed sites of pathophysiological interaction, pathways for spread of activity and modifications of epileptic brain to experience and treatment.

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### **18.3 Issues/Problems in Developing Models of Epilepsy**

#### **18.3.1 How Many Models Are Necessary for an Epilepsy with a Given Etiology?**

As there is no perfect model of human partial epilepsy, models must be chosen on the basis of the long-term goal to be addressed. If the goal is to determine whether a specific therapeutic agent decreases the incidence of seizures either prophylactically or after epilepsy is established, a “high throughput” model with a relatively short latency between injury and seizure activity and a high proportion of animals developing behavioral seizures would be necessary to adequately power the experiment without exhausting available manpower or other resources. Lesional models with long latencies from injury and lower rates of occurrence of seizures would be impractical. The choice of models of chronic focal neocortical epilepsy for use in development of prophylactic or therapeutic strategies is particularly vexing

due to long latencies, relatively low incidence of clinical seizures and variability between laboratories or even in the same laboratory. These requirements have led to the predominant use of models of chronic limbic system epilepsy following status epilepticus induced by pilocarpine, kainic acid or repetitive electrical stimulation. This approach begs the issue of whether other models of epileptogenesis such as those following traumatic injury in temporal lobes or neocortex have the same distribution of underlying mechanisms, and whether post-status epileptogenesis is a common pathophysiology in man. In other words, are we putting “all of our eggs into one basket?” Obviously, all pathologies cannot be represented in a given model. In other CNS disorders, such as autism, schizophrenia and Alzheimer’s disease, experiments have been done in a variety of models for a given condition, resulting in conclusions that multiple pathophysiologies may contribute to a given phenotype.

If, on the other hand, the experimental goal is to elucidate the basic cellular and synaptic mechanisms that may contribute to hyperexcitability and epileptogenesis, for example following focal cortical or hippocampal injury, a model in which hyperexcitability persists *in vitro* in a high proportion of cortical slices from a known focal area of injury would be preferred over the more diffuse or multifocal brain injuries that occur following status epilepticus or severe brain trauma. In this case, one might choose the partial cortical isolation model or epileptogenic focal areas in cortex due to infarction, controlled local cortical trauma, or experimentally induced focal inflammation/infection.

#### **18.3.2 Multiple Pathophysiological Processes**

Not only are there multiple abnormalities in any given model, but also these abnormalities do not occur in parallel over time. This has important implications for choice of therapy, be it prophylactic or after seizures have developed. For example, early on after injury, inflammation, alterations in the blood brain barrier, excessive release of glutamate from injured tissue and abnormalities

in membrane properties or receptors of acutely damaged neurons may be most important as targets for whatever agents are chosen. Further, it may be unclear which of these processes or combination of them is epileptogenic, and underlies the later development of seizures. Inflammation and blood brain barrier disturbances are present following any cortical trauma, yet only a minority of mild to moderate injuries result in partial epilepsy. Likewise, only a small proportion of gray matter infarctions result in focal epileptogenesis, even though similar acute processes occur following most injuries. Over time other more indolent processes may occur such as progressive loss of nerve cells following repetitive seizures or slowly activating mechanisms that induce either adaptive or maladaptive circuitry (e.g. [23, 27, 42, 49]). The choice of a therapeutic agent would depend on which of these processes was ongoing at a given point in time; it might not be effective to treat an area of injury with an anti-inflammatory agent after epileptogenesis is well established. The best experimental strategy would be to attempt to isolate or control one or another of these potential epileptogenic processes and assess the end result in a preparation that is sensitive enough to detect small changes in whatever is being measured.

### 18.3.3 Variables That May Affect the Development of Epilepsy After Cortical Injury and the Interpretation of Results of Modeling Experiments

- (i) Severity of injury and resulting epileptogenesis: It is clear that the severity of injury is a key prognostic factor in human posttraumatic epilepsy and one that may affect experimental results in a model [5]. Further, in models of severe traumatic injury or prolonged status epilepticus, multiple brain regions may be affected, making it difficult to determine site(s) of seizure origin. As discussed above, in experiments testing either prophylactic or therapeutic agents, it is desirable to use “high throughput” models in which there is frequent and intense seizure activity. Under these circumstances, it is possible that a therapeutic trial would appear to be negative because of the intense epileptiform activity, even though the agent employed was altering its target, as hypothesized. Other epileptogenic mechanisms might be powerful enough to hide favorable actions, leading to a false-negative trial.
- (ii) Site(s) and distribution of lesions (focal, multifocal, diffuse; hippocampus vs. neocortex) may influence results: Different cortical areas have varying susceptibilities to the development of epileptiform activity. Such differences in epileptogenic capacity from region to region with a given injury (e.g. [5]), or even within different laminae in the same cortical area [4, 36] may be due to variability in circuitry, receptors and intrinsic cell properties. Whatever the mechanisms, this variability makes it important to focus modeling experiments on specific neuronal types and structures comparable to those thought to be involved in clinical epileptogenesis. These intrinsic differences make it important to sample a given cell type or area, recognizing that there may be significant differences if experiments are carried out in another cortical region. There is marked variability in incidence, severity and frequency of seizures, even in the same posttraumatic model in the same laboratory [5]. This variability resembles that seen following human head injury, but also makes testing of antiepileptic or prophylactic strategies more difficult and raises questions about models in which almost all animals have frequent seizures.
- (iii) The etiology as well as severity of a human cortical lesion may be a factor that determines the likelihood of epileptogenesis and success of a planned intervention. Penetrating injuries and those that induce intracerebral bleeding have a higher incidence of seizures than those resulting from infarction or closed head injury. There are also sometimes striking differences between incidence of seizures in different models in the same laboratory [17], and between different laboratories using the same model. Some of this variability may be due to errors in experimental design [21, 34].



- (iv) Age and species: Assumptions regarding applicability of specific findings from models of epilepsy in one strain or species to another, or to human epilepsy, should be made with caution. These differences extend to transport of antiepileptic drugs [1]; induction of status epilepticus and its consequences [2, 22, 55]; seizure-induced cell injury or death [43]; kindling [12, 50], and effects of ischemia [54]. Susceptibility to epilepsy may be greater in the immature brain [16, 52], although some parameters that are thought to be important to epileptogenesis, such as the maturation of excitatory axonal arbors of cortical pyramidal cells, are slow to develop fully [40]. This makes results of experiments performed in models of epilepsy in immature in vitro slices difficult to generalize to mature cortical CNS structures.

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#### 18.4 What Are Some Key Issues for Clinical and Basic Research in Partial Epilepsies?

- (a) Latent period between injury and seizures provides evidence for ongoing epileptogenic processes following cortical injury and an opportunity for prophylaxis.

There may be a critical period within the first few days after injury when therapeutic intervention will be effective, even though the latency to seizures is significantly longer (e.g. [6, 13, 14, 26]). Further analyses of pathophysiological events that occur during the critical period and are interrupted during such experiments may lead to new effective antiepileptogenic approaches.

- (b) Non-epileptogenic vs. epileptogenic injury.

What are the genetic or other susceptibility factors that predispose an individual to epileptogenesis after injury (e.g. [8, 51])?

- (c) Which changes in epileptogenic brain are adaptive vs. maladaptive?

Injury-induced axonal sprouting has been considered a key *maladaptive* epileptogenic mechanism in a variety of models, and in human cortical structures. (reviewed in [25,

37]). However, establishment of new connectivity may also be an important *adaptive* mechanism that underlies recovery from stroke and other injury [7, 23, 27]. Recent results show that excitatory synaptic connectivity and epileptogenesis can be significantly reduced in cortically injured rats by treatment in vivo with gabapentin, a drug that interferes with synaptogenesis induced by astrocytic thrombospondins [26]. Will such drugs also limit behavioral recovery from brain injury? Additional experiments are required in models of injury-induced epileptogenesis to determine whether maladaptive connectivity can be limited without affecting adaptive mechanisms that foster behavioral recovery.

- (d) New (targeted) drug development; pharmaco-resistance.

Why have rates of seizure control not increased, in spite of introduction of multiple new anti-epileptic drugs?

- (e) Mechanisms of interictal-ictal transitions. Why a seizure today? What starts it? How does it propagate? What ends it? “*Why does the relatively restricted sporadic discharge of chronically epileptic neurons become periodically enhanced and propagated to produce overt seizures?*” [20].

- (f) What are the trigger mechanisms for sporadic seizures? Roles of “stress”, sleep, fever, hormones, etc.

- (g) Effects of epileptiform activity on neocortical and limbic structure/function.

- (h) What are the long term impacts of epilepsy on the mature and developing brain?

- (i) Co-morbidities as targets for research using animal models.

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#### 18.5 Conclusions

1. In assessing models of neocortical or temporal lobe epilepsy, it is important to first identify the specific issue to be addressed in the laboratory, derived from clinical observations and research, or results of previous experiments.

2. The long list of variables detailed above, that can influence results and conclusions, should be considered in advance and experimental and control groups and protocols planned accordingly.
3. Convenience is not the most important criterion for use of a given model. In the case of preclinical trials of agents for chronic partial epilepsy, the choices are quite limited, as seizure frequency sufficient to power the experiments is present predominantly in post-status models where widespread abnormalities are present and the analogy to the pathophysiology of spontaneously-occurring clinical partial epilepsy is unclear [29].
4. A major question for the model chosen will be whether expected results, based on known or expected variability of data, will yield an unambiguous answer to a specific issue, within practical limits of available resources.
5. A broader definition of the term “translation” is necessary, to include mechanisms by which defects at molecular, cellular and network levels are “translated” or evolve into the dysregulated cortical activities that generate epileptiform activity and behavioral events. Without knowledge of events at this level, the “Lost in Translation” phenomenon, i.e., failed clinical trials, false negative experiments, or collection of data irrelevant to the clinical issue, will be more likely.
6. “A really complete understanding of epilepsy might require almost total knowledge of the central nervous system” [20]. Much of our progress in epilepsy research derives from non-epilepsy related experiments in basic neuroscience. Too much emphasis on the “relevance” of a particular model or approach to epileptogenesis may limit discovery of major contributing mechanisms derived from less targeted experiments.

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# Aligning Animal Models with Clinical Epilepsy: Where to Begin?

# 19

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## Abstract

Treatment of the epilepsies have benefitted immensely from study of animal models, most notably in the development of diverse anti-seizure medications in current clinical use. However, available drugs provide only symptomatic relief from seizures and are often ineffective. As a result, a critical need remains for developing improved symptomatic or disease-modifying therapies – or ideally, preventive therapies. Animal models will undoubtedly play a central role in such efforts. To ensure success moving forward, a critical question arises, namely “How does one make laboratory models relevant to our clinical understanding and treatment?” Our answer to this question: It all begins with a detailed understanding of the clinical phenotype one seeks to model. To make our case, we point to two examples – Fragile X syndrome and status epilepticus-induced mesial temporal lobe epilepsy – and examine how development of animal models for these distinct syndromes is based upon observations by astute clinicians and systematic study of the disorder. We conclude that the continuous and effective interaction of skilled clinicians and bench scientists is critical to the optimal design and study of animal models to facilitate insight into the nature of human disorders and enhance likelihood of improved therapies.

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Animal models have played a critical role in epilepsy research dating back to 1937 when Putnam and Merritt [32] published the first animal model of seizures, the electroshock test in cats. Since that time, thousands of papers have been published that detail the development and utilization of animal models of seizures and epilepsy. Currently, there are over 100 different animal models employed in epilepsy research [33, 35]. These models utilize a wide array of species including drosophila, zebrafish, mice, rats, guinea pigs, cats, and even non-human primates. These models have provided insight into cellular and molecular mechanisms surrounding many aspects of the epilepsies. Moreover, some of these models have led to the development of novel therapies in the clinic.

That said, much work remains. Current pharmacologic treatments for epilepsy are “symptomatic” insofar as they suppress but do not prevent, modify, or cure the disorder. There is a critical need for new and improved treatment options that promote not only enhanced symptomatic therapy, but also (for the first time) provide disease-modifying or preventive therapy. Satisfying this need will require the use of appropriately designed and implemented animal models. To ensure success, we must address the following question: “How does one make laboratory models relevant to our clinical understanding and treatment?”

In our view, the answer to this question starts with a detailed understanding of the clinical phenotype one seeks to model. This understanding in turn guides design and analysis of the animal model. Here we choose two examples to illustrate our thinking. One consists of a monogenic disorder, Fragile X syndrome. The other is a subtype of the common, sporadic disorder temporal lobe epilepsy – namely the syndrome of mesial temporal lobe epilepsy emerging months to years after an episode of prolonged seizures (status epilepticus).

For each example, we will focus on a specific animal model that promises to inform clinical understanding and treatment.

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## 19.1 Evaluating Clinical Relevance

Evaluating the clinical relevance of an animal model is not a question unique to epilepsy research. In fact, this question is seminal to preclinical investigation of most human diseases. One approach to considering the relevance of a model for a particular human disease involves model evaluation using three criteria: its construct validity, its face validity, and its predictive validity [5].

“Construct validity” refers to how closely an animal model recapitulates the causal mechanisms underlying the disease in humans [5]. Construct validity is most readily addressed with monogenic disorders in which clinical and molecular genetic analyses have elucidated the molecular etiology of the syndrome in humans. For example, in Dravet Syndrome, the underlying cause for most human cases consists of *de novo* mutations of the *SCN1A* gene that result in loss of function [7]. Consequently, approaches to developing an animal model of Dravet Syndrome with high construct validity would include engineering an experimental animal (e.g. fly or zebrafish or mouse) with a null mutation of *SCN1A* or by substituting the wild type gene with an actual mutation identified in a human, a strategy referred to as “knock-in”.

“Face validity” refers to how closely an animal model recapitulates phenotypic characteristics of the human disease [5]. For example, patients with Rett Syndrome display several characteristic features, including cognitive impairment, breathing irregularities, and stereotypic hand movements [18, 29]. An animal model of Rett Syndrome with high face validity would reproduce most if not all of these phenotypic features.

Finally, “predictive validity” refers to how closely an animal model recapitulates treatment responsiveness observed in humans [5]. For example, absence seizures in humans are typically quite responsive to the pharmacologic agent ethosuximide [16]. An animal model for absence seizures that has high predictive validity would demonstrate a similar response to ethosuximide.

As we consider how to enhance the relevance of animal models to our clinical understanding and treatment, we will use these criteria as a framework. Ideally, animal models would have high validity for each of these three criteria. However, as we discuss below, this may not be possible, and – importantly – may not be necessary in order to inform clinical understanding and/or treatment of epilepsy. In fact, having high validity in only one criterion – construct, face, or predictive – may still provide a useful model for the appropriately selected question.

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## 19.2 Mendelian Disorders of Epilepsy

Mendelian disorders of epilepsy are those in which clinical and molecular genetic evidence establishes the mutation of a single gene as the cause of the disorder. To date, mutations in over 100 genes comprising a wide range of proteins have been linked to human diseases in which epilepsy is one of the phenotypic manifestations [28, 34]. Collectively, these Mendelian epilepsies account for only a small fraction of all epilepsies [28]. That said, study of these disorders will hopefully benefit individuals affected with these mutations, and insights derived from such studies may also inform mechanisms of non-Mendelian epilepsies.

For these Mendelian epilepsies, identification of the causal mutant gene by clinical and molecular genetic studies creates the opportunity to engineer an animal model by introducing the mutant gene into the genome of an experimental animal (e.g. fly, zebrafish, mouse, etc.) and examining its phenotypic manifestations. Such models typically have high construct validity because scientists can incorporate the precise genetic abnormality seen in humans, whether this

abnormality is a point mutation, chromosomal translocation, a frameshift mutation, etc. This high construct validity commonly equates to high face validity – the models recapitulate the key phenotypic features of the human disease. In these situations, the high construct and face validity strengthen the likelihood that such models will have high predictive validity as well.

However, high construct validity does not assure high face validity. For example, cystic fibrosis is a disease characterized by multi-organ failure with recurrent and persistent pulmonary infections being quite prominent. A common cause of cystic fibrosis is a mutation of F508 in the *CFTR* gene [17]. Mouse models with this exact mutation in their endogenous *Cftr* gene do not reproduce the severe pulmonary phenotype seen in humans. There are numerous possible explanations for this disparity including differences in the genetic background, immune response, etc. In spite of such a disparity, the low face validity of these models does not preclude their usefulness for addressing important questions. In fact, these cystic fibrosis models have been used extensively and with good success to probe questions surrounding other aspects of the human disease. The key issue is that the question addressed in the animal model must be carefully aligned with a specific and important question arising in the human disorder.

To illustrate these considerations in greater detail, we consider Fragile X syndrome – a disorder in which epilepsy is a prominent manifestation and for which engineering genetically modified mice have produced useful models that promise to inform our clinical understanding and treatment of this disease.

### 19.2.1 Characterizing Fragile X and Developing an Animal Model

Fragile X syndrome is a genetic disorder occurring in 1:5,000 males [8]. Phenotypic manifestations include seizures, autism, cognitive impairment, hypersensitivity to sensory stimuli, motor incoordination, growth abnormalities, and various physical characteristics such as an

elongated face, large protruding ears, and macroorchidism [6, 9, 30]. Clinical and molecular genetic investigations have led to identification of the molecular etiology of this syndrome, namely, a mutation of a gene termed “Fragile X”. Development of an animal model for Fragile X has been decades in the making. Its history began in 1943 when two clinicians (J. Purdon Martin and Julia Bell) described a family in England in which 11 males of two generations presented with mental retardation and social withdrawal [26]. Based on the pedigree, these clinicians hypothesized that this presentation of symptoms represented a novel, sex-linked recessive genetic disorder. Microscopic evaluation of chromosomal spreads isolated from these patients revealed the X-chromosome to be deformed or broken, thereby leading to the name “Fragile X” [24]. It took nearly 50 years, but the causative gene on the X-chromosome was finally identified to be *FMRI* [39]. The primary mutation within this gene leading to Fragile X was an expansion of the CGG trinucleotide repeat found within the 5′ untranslated region of *FMRI*. Investigators quickly demonstrated that this genomic expansion in turn leads to transcriptional silencing of the gene and thus a lack of the protein encoded by this gene – fragile X mental retardation protein (FMRP; [31]). With the gene, the mutation, and the effect on protein expression documented, scientists next set out to develop an animal model for Fragile X. By introducing a null mutation of the *Fmr1* gene into the genome of a mouse, an animal model was developed that recapitulated the loss of FMRP expression observed in humans [14]. By definition, this model does not exactly recapitulate the initial pathologic lesion underlying the human disorder, namely, the CGG trinucleotide expansion, and thus does not have *perfect* construct validity. However, it does recapitulate what is likely the primary consequence of the mutation, loss of FMRP expression.

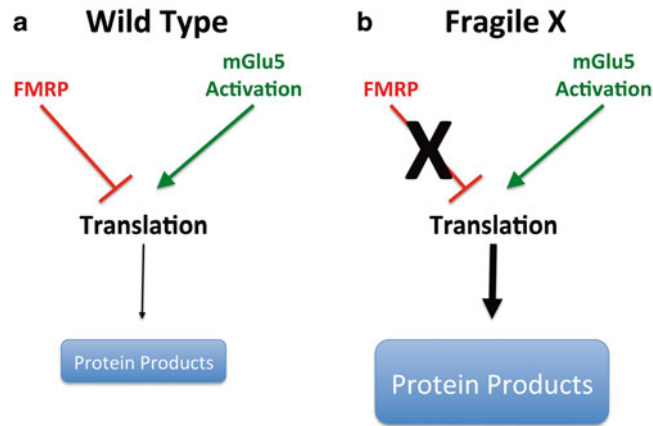
The fact that the *Fmr1* knockout mouse is not identical to the human genetic abnormality, yet recapitulates the human protein abnormality, raises an interesting point regarding animal model development. Specifically, a model lacking *perfect* construct validity may nonetheless shed light on clinical understanding and treatment of a

disorder. Indeed the *Fmr1* knockout mouse does recapitulate many aspects of the human phenotype, thereby giving it high face validity. Similar to humans with Fragile X, these mice exhibit seizures, cognitive problems, hyperactivity, and macroorchidism [4, 14, 30]. The high face validity of this mouse model has led to an intense search for how the genotype causes the phenotype. Briefly, FMRP is highly expressed within neurons, especially at synapses [4, 10, 30]. Here, it binds many messenger RNAs and represses translation of these mRNAs into protein [2, 22, 36, 42]. Upstream of these events is the G-protein-coupled glutamate receptor, mGluR, the activation of which leads to protein translation [21, 40]. The loss of the repressive effects of FMRP in Fragile X allows for unopposed mGlu5 signaling, which in turn results in excessive protein translation (Fig. 19.1). It is this unopposed mGlu5 signaling that likely contributes to the phenotypic manifestations of Fragile X syndrome, because crossing FMRP mutant mice to mice in which one allele of mGlu5 has been eliminated reduces seizures and other abnormalities of the FMRP mutant mouse [11]. These findings have led to development of potent and selective inhibitors of the mGlu5 receptor. Continuous treatment of FMRP mutant mice with mGlu5 inhibitors commencing early in life eliminates seizures and other phenotypic abnormalities. Moreover, initiating treatment with mGlu5 inhibitors in adult FMRP mutant mice *after* the development of seizures and other abnormalities reduces these seizures and corrects these other abnormalities. Collectively, these findings have provided the foundation of a clinical trial for patients with Fragile X syndrome with an mGlu5 inhibitor, results of which will inform the predictive validity of this model.

### 19.2.2 *Fmr1* Knockout Mouse – A Model Facilitating Clinical Understanding and Treatment of Fragile X Syndrome

In sum, clinical recognition of the distinctive phenotype and its familial aggregation provided the foundation for discovery of the mutant gene





**Fig. 19.1** FMRP and mGlu5 modulation of protein translation. (a) In wild type animals, FMRP inhibits protein translation while mGlu5 activation promotes translation. By balancing these opposing actions, the appropriate amount of

protein products is generated. (b) However, in mouse models of Fragile X (*Fmrp* knockout animals), the loss of the repressive effects of FMRP leads to unopposed mGlu5 signaling and ultimately excessive protein synthesis

decades later. Insight into the nature of the causative mutations, in turn, enabled engineering a genetically modified mouse model. This model illustrates how discovery of the molecular mechanisms by which the genotype leads to the phenotype can give rise to identification of a target for development of small molecules that could be used as drugs to treat the disorder. Careful alignment of the animal model with the clinical phenotype, an alignment simplified by knowledge of the molecular etiology afforded by molecular genetics, has led to a sequence of discoveries that in turn have enabled design of a clinical trial based upon disease mechanism.

### 19.3 Non-Mendelian Disorders of Epilepsy

The vast majority of the epilepsies do not exhibit a Mendelian pattern of inheritance; instead they typically arise sporadically as a consequence of various cortical lesions including developmental abnormalities, neoplasms, traumatic brain injury, and vascular insults. In contrast to a Mendelian disorder in which development of an animal model is based upon a known molecular etiology (i.e., a mutant gene), here the model must rely on recapitulating some feature(s) of the clinical syndrome. Once again, however, a detailed characterization

of the syndrome in humans is of critical importance to both appropriate design and evaluation of the animal model. One non-Mendelian epilepsy syndrome that has been extensively characterized by clinicians is a form of temporal lobe epilepsy (TLE) arising long after an episode of prolonged seizure activity (status epilepticus – SE).

#### 19.3.1 The Clinical Syndrome

TLE is the most common and also most devastating form of partial epilepsy in humans [33]. Broadly defined, TLE is an epilepsy in which seizures most commonly are initiated from the medial temporal lobe. A diversity of etiologies of TLE has been identified, implying that TLE comprises multiple disease subtypes. Despite such heterogeneity, some features are conserved – most notably the associated asymmetric pattern of hippocampal neuronal loss and gliosis, termed hippocampal or temporal lobe sclerosis [33].

One proposed subtype of TLE that presents with hippocampal sclerosis is that arising after an episode of SE [41]. Retrospective analysis of patients undergoing surgery for intractable TLE reveal that many of these patients experienced an episode of prolonged, focal, severe seizures (SE) many years prior to epilepsy development [15]. Most commonly, these severe seizures occurred

in the context of complicated febrile seizures during infancy or childhood but similar observations have been made following afebrile status epilepticus arising *de novo* in adults. Longitudinal studies have confirmed these observations in that up to half of individuals experiencing *de novo* status epilepticus of either febrile or afebrile origin in childhood or adulthood develop recurrent seizures (epilepsy) after a seizure-free latent period of variable duration [1, 37]. Importantly, inducing SE experimentally in an otherwise normal animal is sufficient to trigger the subsequent development of TLE. Based on these converging lines of evidence, it seems likely that the occurrence of *de novo* SE during infancy or adulthood contributes to TLE development in humans.

One prominent feature of this syndrome is a structural abnormality referred to as Ammon's Horn or hippocampal sclerosis [43]. It has long been recognized that many patients with TLE have atrophic and damaged hippocampi as visualized on MRI or histopathologic examination [25, 43]. Animal studies provide convincing evidence that severe seizure activity is sufficient to induce hippocampal damage similar to hippocampal sclerosis observed in humans, namely, neuron loss predominantly in the hilus and CA1 as well as mossy fiber sprouting in the dentate gyrus [12]. However, the specific relationship between hippocampal sclerosis and epileptogenesis has been highly debated. In our view, it seems plausible that hippocampal sclerosis is both a consequence of SE and can contribute to development of TLE. In the context of this controversy, there emerged an important clinical observation: MRI evidence of acute hippocampal injury within days following complicated febrile seizures, an event followed months later by hippocampal atrophy [38]. This MRI abnormality is evident in a subset of children following an episode of febrile status epilepticus. The question arises as to whether the subsequent emergence of TLE years later occurs in the subset with hippocampal damage and not in those with normal hippocampi (as detected by MRI) following status epilepticus. Addressing this question is the objective of a multicenter, longitudinal study (the FEBSTAT study) of children undergoing

complicated febrile seizures, a study that will permit correlating the occurrence of acute hippocampal injury and subsequent hippocampal sclerosis with the later emergence of TLE [19]. Importantly, the detailed analysis of this syndrome will provide the information needed to design and characterize animal models properly aligned with the human disease.

### 19.3.2 Animal Models of SE-Induced Epilepsy

SE-induced TLE is a heterogeneous disorder. Variability in presentation can be seen at almost every aspect of the disease – SE etiology, latent period duration, epileptic seizure severity, histopathology, etc. As such, it is unlikely that one single model can perfectly recapitulate all of its many facets. For this reason, it is no surprise that many different models for SE-induced TLE have emerged. That said, there are several key features that an animal model of this syndrome is expected to reflect. First, the model should begin with a brief episode of SE that is followed by emergence of spontaneous recurrent seizures after a latent period. Second, the model should correlate SE with *unilateral* hippocampal sclerosis, consistent with the pathologic findings noted in humans on both MRI and histopathology. Third and finally, the model should produce adult-onset of TLE as a result of either SE in adulthood [37] or in infants and children [1].

One model that fulfills these three criteria is TLE arising following SE induced by microinjection of the ionotropic glutamate receptor agonist, kainic acid (KA), into the amygdala. This model can be induced by infusion of KA into the amygdala of either young (P10) or adult rodents, the resulting SE leading to subsequent development of epilepsy. In adult mice and rats, microinjection of KA into the basolateral amygdala nucleus leads to almost immediate onset of status epilepticus [3, 27]. Typically, SE is allowed to continue for 40 min, at which point a benzodiazepine such as diazepam or lorazepam is administered to stop the seizure activity. Approximately 3 days after the initial

SE event, spontaneous recurrent seizures arise and appear to persist lifelong. In P10 rat pups, a similar approach has been utilized [13]. Kainic acid is microinjected into the basolateral amygdala nucleus leading to almost immediate onset of SE that lasts for several hours. Typically, SE is allowed to continue until its natural termination (as opposed to the pharmacologic intervention used in adult animals). When these animals are evaluated 4 months later, they exhibit both behavioral and electrographic seizure activity, demonstrating the emergence of TLE.

This model is one of several in which induction of SE in an otherwise normal rodent results in emergence of TLE. Other methods include systemic pilocarpine administration (a muscarinic agonist), systemic kainic acid administration, or focal electrical stimulation. These methods are effective and have been used extensively by many labs, with each of the models exhibiting advantages and disadvantages. Adapting the intra-amygdala KA model to mice [3] simplifies study of genetically modified animals, providing a powerful tool for elucidating molecular and cellular mechanisms of epilepsy. Additional advantages of the intra-amygdala KA model in the mouse include: 100 % of KA-injected mice develop SE; mortality is only 10–20 %; and 100 % of surviving animals become epileptic [3, 13, 27]. The efficiency together with low attrition provides important advantages, especially for studies of genetically modified animals with limited availability.

Importantly, a number of features of this model align with the clinical syndrome. To begin, this model mimics the initial pathologic insult observed in many patients, namely, SE. Furthermore, this model can induce epilepsy via SE in both young and adult animals, similarly to that observed in humans. However, the construct validity is not perfect in that in this model, SE is induced by a convulsant (KA) while in the majority of children, SE arises in the context of a febrile illness. That said, the fact that SE, whether induced by diverse chemical methods or electrical stimulation, causes TLE suggests that the key variable promoting epileptogenesis

in most instances is the occurrence of SE *per se*, not the cause of the SE.

In terms of face validity, this model does recapitulate many but not all components of the clinical syndrome. First, this model does mimic the temporal course of the disease in that SE leads to a latent period which in turn evolves into TLE. One area of debate with this model is the length of the latent period. In humans, the time between SE and TLE is on the order of months to years while in this model it is only a few days. Second, this model does yield unilateral hippocampal sclerosis following epilepsy onset that can be detected by both MRI and histopathologic analysis [13, 27]. One caveat is that the pattern of neuron loss within the hippocampus is different from that observed in humans. For most human specimens, neuron loss is most prominent in the hilus and CA1 regions [43]. In the intra-amygdala KA model, neuron loss is most prominent in CA3 and hilus, leaving CA1 relatively spared [13, 27]. Lastly, in humans with TLE, memory deficits and other comorbidities are common [25]. To date, there are no studies clearly documenting memory deficits following implementation of this animal model. However, recent work revealed the occurrence of anxiety-like behaviors in this model [23].

Regarding predictive validity, since there is currently no preventive therapy for TLE arising after SE in humans, it is not possible to assess the predictive validity of this model. However, the utility of this model has enabled discovery of two molecular targets that show promise for development of preventive therapy. First, David Henshall and colleagues reported that expression of the microRNA, miR-134, is increased following SE and that inhibiting miR-134 expression shortly after SE onset may be antiepileptogenic [20]. Second, work from our lab revealed that SE induced the enhanced activation of the BDNF receptor tyrosine kinase TrkB [23]. The utility of this model in the mouse enabled a powerful chemical-genetic approach using genetically modified mice. This approach led to the discovery that inhibition of the TrkB kinase activity, commencing following SE and continued for just 2 weeks, prevented development of TLE in

more than 90 % of animals when they were tested a month later. These discoveries, made possible by adapting this model to the mouse (40), provide novel targets for development of preventive therapy for this particular syndrome. Whether similar molecular mechanisms underlie development of TLE induced by different causes (e.g. trauma, developmental abnormalities, etc.) is uncertain.

### 19.3.2.1 Aligning Animal Models with Human Disease

The epilepsies represent a collection of heterogeneous disorders for which only symptomatic treatment is currently available. The lack of efficacy, together with undesirable consequences of symptomatic therapy for many patients, underscores the need to develop preventive therapies. Development of preventive therapies based upon disease mechanism requires properly aligning the animal model with the clinical syndrome. This is a challenging task, one that must begin with a detailed characterization of the clinical disorder. Such information provides a context critical to design and study of animal models that recapitulate key features of the clinical disorder. This descriptive first step underscores the importance of continuous and effective interactions of clinicians and bench scientists to assure the optimal alignment of animal models with human diseases, thereby enhancing the likelihood that study of the models will contribute to understanding the mechanisms of the disease and improving treatment. In short, the most effective and efficient way to develop animal models is to start at the bedside, move to the bench, and with a lot of hard work and luck, return to the bedside with a novel therapy in hand.

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# What Non-neuronal Mechanisms Should Be Studied to Understand Epileptic Seizures?

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Damir Janigro and Matthew C. Walker

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## Abstract

While seizures ultimately result from aberrant firing of neuronal networks, several laboratories have embraced a non-neurocentric view of epilepsy to show that other cells in the brain also bear an etiologic impact in epilepsy. Astrocytes and brain endothelial cells are examples of controllers of neuronal homeostasis; failure of proper function of either cell type has been shown to have profound consequences on neurophysiology. Recently, an even more holistic view of the cellular and molecular mechanisms of epilepsy has emerged to include white blood cells, immunological synapses, the extracellular matrix and the neurovascular unit. This review will briefly summarize these findings and propose mechanisms and targets for future research efforts on non-neuronal features of neurological disorders including epilepsy.

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## Keywords

Anti-epileptic drugs • Cerebrovasculature • Drug resistance • Brain endothelium • Glia-neuronal interactions • Extracellular matrix

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This book is devoted to one of the all-time leaders in epilepsy research, Philip Alan Schwartzkroin. Phil has not only changed the traditional understanding of mammalian neurophysiology but he also revolutionized the tools we employ to study the brain as one of the people to perfect the brain slice preparation [70, 71, 74–76]. Last but not least, Phil has edited many seminal books and papers, and incessantly contributed to the recruitment and scientific development of scores of young scientists. Under the shadow of this giant (and former mentor for one of us (DJ)) writing this review is a

humbling experience; one way to start the process is to refer the reader to Phil's recent introduction to the field of epilepsy research [72].

## 20.1 Why Study Non-neuronal Mechanisms in Neurology or Neuroscience?

We study the brain for many reasons, not least of which is for its intrinsic interest (see Schwartzkroin's recent introduction to the field [72]). The fascination with neurons and neuronal circuitries is not surprising since neurons are the collectors and effectors of our daily experiences and actions. In the specific case of epilepsy research (clinical or basic/translational), the quest for "epileptic neurons" or "epileptic circuits" has produced remarkable results, leading to the discovery of viable anti-epileptic drug targets and to the multimodal definition of the "epileptic focus", an invaluable clinical tool for the neurosurgeon. However, as in other neurological disorders, a neurocentric approach has left certain questions unanswered and experimental opportunities remain. The most striking example of why neuroscience should become more "holistic" is embolic stroke, a disease stemming from cerebrovascular disease that has devastating consequences on brain function. After the NIH convened a Stroke Progress Review Group in 2001, stroke research shifted from a purely neurocentric focus to a more integrated view wherein dynamic interactions between all cell types contribute to function and dysfunction in the brain. In the field of epilepsy research and treatment, there is no pressing need for such a sharp re-direction, since the field is already characterized by the study of many cell types, and non-neuronal processes. For example:

1. Many neuronal molecular, morphological defects or functional abnormalities described in human epileptic brain are present throughout the cycle of interictal-to-ictal states that characterize the epileptic brain. The persistence of these neuronal abnormalities does not fully explain why at a given time point an interictal cortex develops a seizure. Other

mechanisms, such as changes in cerebral blood flow or blood-brain barrier permeability have been proposed to mediate the interictal to ictal transition.

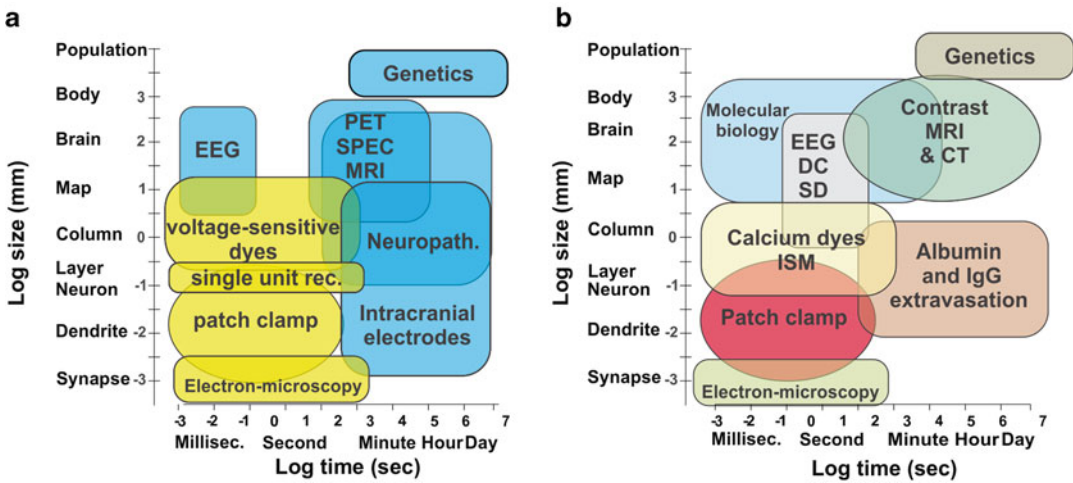
2. It has been proposed that the process of epileptogenesis is distinct from the process of ictogenesis. According to this hypothesis, what makes a brain epileptic (e.g., genetic mutations, acquired or inherited; malformations of brain development) does not directly cause seizures. In fact, seizures can occur in "non-epileptic" brain and people with epilepsy spend most of the time not having seizures, indeed many experience only a few seizures per year. Again, as in (1), non-neuronal mechanisms spanning from altered cerebral blood flow to glial dysfunction have been used to explain how an asymptomatic neurologic condition can suddenly develop into a seizure state or the fact that seizures can occur in non-epileptic brain (e.g., stroke).
3. Multiple drug resistance to anti-epileptic drugs affects over 20 % of patients with epilepsy. Multiple drug resistance cannot be fully explained in pharmacodynamic or neuronal terms, and great emphasis has been put on pharmacokinetic mechanisms that include the blood-brain barrier.
4. Analysis of resected or *post-mortem* epileptic brain reveals a number of pathophysiological changes in astrocytes and microglia. MRI studies show, in addition to persistent structural changes such as malformations of brain development, an array of transient changes that reflect post-ictal or interictal functional fluctuations in the extracellular space (increased FLAIR signal, perfusion changes *etc.*).
5. The analysis of molecular transcripts and changes in gene expression in patients with epilepsy reveal a surprising number of genes and proteins that are involved in astrocytic function, blood-brain barrier maintenance and transport, as well as immune signaling and extracellular matrix proteins.

The following paragraphs detail the rationale for new or corroborative experiments that will help understand the extent and nature of non-neuronal mechanisms of seizure disorders.

## 20.2 Identification of Important Problems

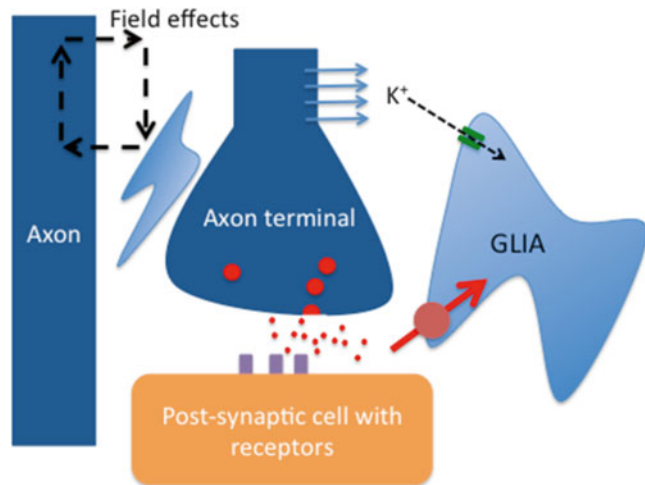
The translational nature of modern research affords the unique opportunity to use real life clinical problems and “translate” these into meaningful laboratory efforts. As beautifully illustrated by Phil in his summary of basic mechanisms [72], the tools used for research are not always the same used in clinical practice. In fact,

a substantial discrepancy in size and temporal resolution becomes evident when comparing clinical and laboratory-based approaches (Fig. 20.1). For this mini review, we will focus on three fundamental yet often neglected aspects of ictogenesis and epileptogenesis: the blood-brain barrier, glia (Fig. 20.2) and the extracellular matrix. The following paragraphs will summarize current understanding and knowledge gaps related to these cellular and molecular mechanisms of neuronal pathophysiology.



**Fig.20.1** Comparison of methods used in basic (a) or clinical (b) neuroscience. Note the partial overlap and significant differences

**Fig.20.2** Some mechanisms by which glia can affect seizure activity. Glia regulate the concentration of extracellular potassium, the size of extracellular space and so electrical field effects (ephaptic communication), and uptake of neurotransmitter





### 20.3 The Blood-Brain Barrier (BBB)

The BBB is the most important vascular barrier of the CNS. The BBB protects the brain from harmful substances of the blood stream, while supplying the brain with the nutrients required for proper function. The BBB strictly regulates the trafficking of cells of the immune system and pro-inflammatory cytokines from the blood into the brain. Recent findings indicate that neurovascular dysfunction is an integral part of many neurological disorders [35, 88]. In diseases with a compromised BBB, the microenvironment of neurons is altered; infiltration into the brain of cells, ions, or molecules may initiate a CNS response. Failure of the BBB is observed in association with a variety of pathological events, occurring as consequence of either systemic pathologies such as stroke, systemic inflammation and CNS disease such as multiple sclerosis (MS) and epilepsy. Increasing evidence has shown that BBB damage causes abnormal neuronal activity. For example, seizures are observed in MS patients, as consequence of stroke, or during systemic or local inflammation. As a proof-of-principle, we (DJ et al.) and others have demonstrated that failure of the BBB induced by “mechanical” means (such as osmotic shock) can play a key role in the onset of seizures [45].

*In vitro* and *in vivo* experiments on various models of neurological diseases have shown that blood-brain barrier damage accompanies the development of neurological symptoms; in contrast, managing BBB failure promotes recovery and affords neuroprotection. BBB disruption (BBBD) causes seizures in animal models and human subjects [19, 45, 46, 48, 50, 51, 85]. In particular, a model of temporal lobe epilepsy (pilocarpine, PILO) also depends on BBBD [19, 51, 84]. The currently accepted mechanism of BBBD-induced seizures predicts activation of adhesion molecules on endothelial cells and leukocytes [19]. According to this hypothesis, and in analogy to what is observed in multiple sclerosis, leukocyte adhesion to or interaction with BBB endothelial cells is an essential step leading to BBBD. Published results have shown that anti-

inflammatory therapy (e.g., glucocorticosteroids) effectively reduce BBBD and associated symptoms [48]. The specific cell types involved in inflammation-promoted blood-brain barrier dysfunction are poorly understood but many leukocyte families have been shown to be involved, including natural killer cells and cytotoxic lymphocytes [4, 46, 50]. Attempts to curb the immune response, such as the extreme case of splenectomy, have been shown to decrease experimental seizures [50]. While BBBD-induced seizures were independent from the means used to obtain disruption (osmotic, pilocarpine, albumin), a specific molecular effector of pilocarpine-induced seizures, perforin, was only recently identified [50]. Perforin released by T cells may explain how activation of T lymphocytes leads to increased BBB permeability; in fact, this molecule can effectively “perforate” the cell membrane causing a rapid loss of function and eventually cell death. In many ways, perforin actions mimic those of membrane-permeating antibiotics, nystatin or gramicidin.

Another reason to focus on the BBB when studying epilepsy is the failure to generate new brain therapeutics owing to insufficient knowledge of the mechanisms involved in brain drug distribution under pathological conditions. Drug resistance affects a significant number of people with epilepsy; it is estimated that approximately 20–30 % of people with epilepsy fail to respond to available anti-epileptic drugs (AEDs) [4, 26, 30, 37, 48, 61, 67]. In the past decade the over-expression of multidrug transporter proteins (e.g., MDR1) at the blood-brain barrier (BBB) has been proposed as a mechanism that contributes to the failure of AEDs to penetrate into epileptic brain [1, 9, 16, 41–43, 47, 49, 59, 77]. In addition to multidrug transporters, it was shown that transcripts of P450 enzymes are elevated in primary endothelial cells (EC) isolated from drug resistant epileptic (DRE) patients; these enzymes include AED-metabolizers such as CYP3A4, CYP2C19, *etc.* [21]. In addition, transcripts for PHASE II metabolic enzymes are present in DRE EC; these enzymes are responsible for the metabolism of 1st and 2nd generation AEDs; CYP3A4 and MDR1 co-localize at the BBB (and neurons) in human DRE brain [22] and overexpression

of CYP3A4 in DRE EC is associated with exaggerated carbamazepine (CBZ) metabolism. This new metabolic pathway produces the toxic CBZ metabolite quinolic acid (QA) leading to the paradoxical situation of an anti-epileptic drug being metabolized in the proximity of the epileptic focus to a seizure-promoting agent.

In summary, therapeutic considerations (use of anti-inflammatory therapy to treat seizures, BBB transporters in multiple drug resistance to anti-epileptic drugs) and etiologic factors (loss of BBB in seizures) suggest that the BBB is a viable and important target for studies aimed at the understanding and treatment of epilepsy. In addition to the role of the blood brain barrier, two other non-neuronal elements need to be considered – glia and brain extracellular matrix – both of which have been shown to have an increasing repertoire of roles in regulating network and brain excitability.

## 20.4 Neuroglia

“Glia” comes from the Greek meaning glue, and Virchow in his search for connective tissue in the brain, first coined the term neuroglia, considering them a sort of putty that supported the neurons [79]. Later, Golgi distinguished glia from neurons by the lack of an axon and ascribed to them a nutritive as well as supportive role. Ramon y Cajal determined that they were involved in the insulation of nerve cells and axons, a role later confirmed for oligodendroglia by a young Penfield who also established a role of glia in phagocytosis [24]. The repertoire of glia has, however, expanded in recent years from supportive tissue to playing an active role in determining network excitability, both modulating and responding to neuronal activity (Table 20.1).

### 20.4.1 Glia, Extracellular Space and Potassium Buffering

Glia play a critical part in the regulation of the size of the extracellular space, and extracellular ion homeostasis. In particular, they play a crucial role in the regulation of the concentration of

**Table 20.1** Role of glia in the central nervous system

Roles of Glia
A supportive and protective role for neurons
A role in inflammation
Regulation of the size of the extracellular space
Maintenance of ion homeostasis in the extracellular space
Neurotransmitter uptake and synthesis
Providing neurons with energy
Detecting glutamate release from neurons and other glia
Release of neurotransmitters, and regulatory proteins
Synapse formation and regulation
Communication between neuronal activity and cerebral blood flow

potassium [73]. Glia express both aquaporins and potassium channels (inward rectifying and delayed rectifying) that play a role in this glial function through maintaining potassium and water homeostasis [5, 11, 18]. In addition, the connection of glia through gap junctions results in a glial syncytium, which facilitates not only water and potassium buffering but also glial communication [23]. Abnormalities of glial buffering of potassium result in potassium accumulation during neuronal activity. Such an increase in extracellular potassium will result in the depolarization of neurons and may therefore play a role in seizure initiation and spread [44, 73]. Reductions in the size of the extracellular space can affect neuronal communication through enhancement of ephaptic transmission (electrical interactions occurring through juxtaposed neuronal elements, which are lessened by increasing the conductive space between these elements), alterations in neurotransmitter “spill-over” and clearance, and changes in the regulation of extracellular ion concentrations. It is noteworthy that decreasing the extracellular space can promote seizure activity, whilst strategies aimed at increasing the extracellular space and decreasing glial and neuronal swelling can terminate seizure activity [31].

### 20.4.2 Glia and Neurotransmitter Concentrations

Glia also regulate the extracellular concentration of glutamate and GABA. They express the

glutamate transporters GLAST (EAAT1) and GLT1 (EAAT2), which are responsible for most glutamate clearance [12]. These transporters determine the extracellular glutamate concentration, thus shaping the NMDA receptor response and the “spill-over” of glutamate following synaptic release onto other synapses (heterosynaptic activation) and extra-synaptic receptors [36]. Through this means, glial glutamate clearance plays a role in long-term synaptic plasticity. The expression of these transporters is regulated by an interaction between neurons and glia mediated by ephrins [55], which are extracellular proteins involved in neuronal development but which may be altered in injury and have been proposed to be involved in synaptic reorganisation following status epilepticus. Thus mechanisms that may play a part in synaptic reorganisation during epileptogenesis could also be involved in alterations in the expression of glutamate transporters. These possible roles of ephrins in epileptogenesis (see also below) have yet to be fully investigated.

The role of glia in the regulation of extracellular GABA is less clear since the glial GABA transporter (GAT3) seems to be mainly effective when the neuronal GABA transporters (predominantly GAT1) are blocked [34]. However, it is likely that GAT3 regulates a different pool of GABA that derives from non-vesicular sources. Further, GAT3 seems to play a greater part in regulating the extracellular GABA detected by interneurons than that detected by principal cells [80]. It has been proposed that GAT3 can reverse during periods of excessive activity, thus increasing extracellular GABA concentrations [28]. Finally, glia also are involved in the synthesis of neurotransmitters and in the glutamate-GABA shunt by which glutamate is converted to GABA [10]. Glutamate taken up by glia is converted to glutamine, which is then released into the extracellular space. Glutamine is taken up by neurons and converted to GABA. Inhibition of any of these processes results in a decrease in vesicular GABA content, GABA release and consequently GABAergic transmission [39]. Decreases in glutamate uptake that have been observed during

epileptogenesis could therefore not only increase extracellular glutamate but also decrease GABAergic transmission.

### 20.4.3 Glia and Metabolism

The uptake of glutamate by glia may have a further important role in neuronal energetics. Glutamate enters the Krebs cycle and therefore acts as an energy substrate. Glial glutamate uptake also activates the sodium-potassium ATPase, increasing glucose uptake and glycolysis [63]. Thus increases in extracellular glutamate during seizure activity can increase glial metabolism. Consequently, glia release lactate, which is taken up by neurons and used as an energy substrate, particularly during periods of excessive neuronal activity [6]. The role of glia in neuronal metabolism is probably even more extensive than this. Neurons lack pyruvate carboxylase [68], an enzyme that is crucial for replenishment of oxaloacetate in the Krebs cycle. As a result of this, the synthesis of GABA and glutamate can rapidly deplete Krebs cycle intermediaries in neurons. Replenishment of these intermediaries in neurons can, however, occur from direct transport of these intermediaries from glia to neurons. Glia are also a major producer of glutathione from glutamate, cysteine and glycine; glial glutathione production is necessary for protection of neurons from free radicals, which are produced during excessive neuronal activity [17]. Failure of glia to provide energy substrates for neurons could therefore promote neuronal death and disorders of neurotransmitter production and neuronal function. Indeed, glia play a crucial role in neurometabolism but how this is altered during and to what extent it plays a part in epileptogenesis are still unclear.

### 20.4.4 Glia, the Tripartite Synapse and Synaptic Plasticity

One of the main recent advances in our understanding of glia in modulating network activity has been the concept of the tripartite synapse in

which glia in close proximity to synapses play a part in synaptic transmission, along with the presynaptic terminal and postsynaptic cell [2]. Vesicles and vesicle-associated proteins have been detected in astrocytes, often in close association to nerve terminals. Glia can detect glutamate via metabotropic glutamate receptors, which mediate a focal rise in astrocyte calcium, which has been proposed to mediate vesicular transmitter release. Calcium rises in one astrocyte can trigger a calcium wave through the glial syncytium, suggesting a mechanism by which focal activity can spread. Glia can also release neurotransmitter through reverse transport and membrane channels. Most of the studies in this area support glial release of glutamate, d-serine and ATP (which is converted to adenosine by extracellular ectonucleotidases) [27]. Glutamate released from glia can act at post-synaptic NMDA receptors and has been proposed to contribute to paroxysmal depolarizing shifts underlying epileptiform activity [82]. D-serine is a co-agonist at NMDA receptors and d-serine release from glia seems to be necessary for NMDA receptor mediated long term potentiation [29]. Lastly, increased adenosine levels through glial ATP release modulates presynaptic release of glutamate in a bimodal fashion through A1 (decreasing release probability) and A2 (increasing release probability) receptors [81]. Thus glia can alter network excitability over short time periods, and could play a role in both seizure initiation (glutamate/D-serine release) and termination (adenosine).

Glia can also play a longer term role in modulating synaptic transmission through the interaction of ephrins, specifically ephrin-A3 on astrocytes with the EphA4 receptor on dendrites [55]. This is a bidirectional interaction, which regulates the expression of glutamate transporters in glia and modulates spine and synapse formation in neurons. Such interactions are important in synaptic plasticity. In addition, glial ephrin signaling is important for neurogenesis, indicating a role for glia in modulating neuronal development and connectivity [55]. The role that this plays in epileptogenesis has yet to be explored.

### 20.4.5 Glia and Neurovascular Coupling

When neuronal activity increases in an area of the brain, there is a concomitant increase in cerebral blood flow to that area – a phenomenon termed “neurovascular coupling.” There appear to be multiple mechanisms mediating this effect, but there is evidence that glutamate acting via metabotropic glutamate receptors and glutamate uptake by glia can affect the release of vasoactive compounds that directly affect cerebral vasculature [64]. One important consequence of this scenario is that neurovascular coupling may depend upon the release of glutamate rather than local neuronal firing. Indeed, there is accumulating evidence that, although neurovascular coupling correlates both with neuronal firing and local field potentials (i.e. post-synaptic receptor activation through glutamate release), the coupling with field potentials is stronger [40]. This increased blood flow is a critical component of seizure activity that can be detected with ictal SPECT or as an increase in the MRI blood oxygen level dependent (BOLD) signal.

### 20.4.6 Changes in Glia with Epileptogenesis

Brain injury and neuronal loss invariably leads to a reactive gliosis in which there is not only a proliferation of astrocytes but also changes in astrocytic morphology and gene expression [78]. Moreover, a reactive gliosis is observed in multiple pathologies associated with epileptogenesis, including traumatic brain injury, stroke, tumors, vascular lesions and hippocampal sclerosis. Abnormal glia are also found in tuberous sclerosis; specific knockout of the *Tsc1* gene in glia results in seizures [83].

Reactive gliosis may alter regulation of the extracellular space and promote ephaptic transmission. Aquaporin expression in astrocytes changes from astrocyte end feet (i.e., their perivascular location) to a more diffuse expression [5]. This has been proposed to lead to abnormal water

regulation, with perivascular water accumulation and increased water uptake by astrocytes resulting in astrocyte swelling and a decrease in the extracellular space. Breakdown of the blood brain barrier and accumulation of albumin within glia also leads to a reduction in glial inward rectifying potassium channel expression and so decreased buffering of potassium rises [13]. Moreover there is evidence in human epileptic tissue of a change in glial glutamate transporter expression and, from rodent studies of epileptogenesis, decreased efficacy of glutamate uptake [13, 65].

Glial metabolism also changes during epileptogenesis. There is an increase in the expression of adenosine kinase and along with astrocytosis, this leads to decreased adenosine levels with epileptogenesis [7]. There are decreased levels of glutamine synthetase, and a consequent decrease in the glutamate-GABA shunt, resulting in decreased inhibitory transmission [10]. Indeed, a specific reactive gliosis mediated by transfection with a viral vector had no effect on the intrinsic excitability of neighboring neurons, but selectively decreased inhibitory transmission, leading to an inhibitory deficit and increased propagation of excitatory transmission [60]. This is a clear demonstration that reactive gliosis alone is sufficient to promote hyperexcitability. Glial metabolism may also be affected by a reactive gliosis due to decreased glutamate uptake, although the role that changes in glial metabolism have on the development of epilepsy are unclear.

Although it is uncertain to what extent reactive gliosis affects the tripartite synapse, astrocyte calcium rises mediated by activation of metabotropic glutamate and purinergic receptors can promote the generation of seizure activity *in vitro* and *in vivo* [25]. Also glial metabotropic receptors are upregulated in epilepsy [3].

The critical role that glia play in the inflammatory process underlying epileptogenesis is discussed elsewhere in this book.

There has thus been growing evidence that glia can alter network excitability through multiple mechanisms. The possible roles of reactive gliosis and the part that it plays both in the development of epilepsy and the generation of seizures need to be further modeled and studied. The extensive

role that glia play in many critical functions will need to be carefully dissected in order to target specific glia mediated processes during epileptogenesis (Fig. 20.2).

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## 20.5 The Extracellular Matrix (ECM)

### 20.5.1 Physiological Role of the Extracellular Matrix

The extracellular matrix (ECM) consists of molecules that are secreted both by neurons and glia, and that aggregate in the extracellular space. About 20 % of the volume of the adult brain consists of extracellular matrix, and the extracellular matrix plays an essential role in determining the diffusion of small molecules [57]. In contrast to ECM elsewhere in the body, the brain ECM predominantly consists of proteoglycans, glycosaminoglycans (in particular hyaluronic acid), and glycoproteins of the tenascin family. There are also proteins that link the ECM to ECM and to molecules on neurons and glia [15].

The vast majority of the ECM is present in the extra-synaptic space. The ECM also makes up the basal lamina, which contributes to the blood-brain barrier. It has also been increasingly recognized that brain ECM consists of other well-defined components including peri-neuronal nets (mesh-like structures which surround cell bodies and proximal dendrites particularly of parvalbumin-expressing interneurons as a mesh-like structure), and specific components present at synapses which are linked to proteins at the post-synaptic and pre-synaptic membrane [15].

Peri-neuronal nets consist of proteoglycans of the lectican family which link with hyaluronic acid and tenascin-R [86]. Peri-neuronal nets are critical in development, closing critical periods and stabilizing synapses and neuronal plasticity. Digestion of proteoglycans associated with peri-neuronal nets or knockout of tenascin-R affect both synaptic plasticity and the excitability of interneurons. Peri-neuronal nets therefore play a crucial role in regulating network excitability and plasticity.

The extracellular matrix can undergo remodeling, which is dependent upon a series of serine proteases, such as plasminogen activators (in particular urokinase-type plasminogen activator), thrombin, metalloproteinase's, and reelin. All of these have been implicated in neuronal and network plasticity [14]. Alterations and remodeling of peri-neuronal nets permit neuronal reorganization following brain damage and seizures, and during development.

The interaction of the extracellular matrix with neurons can occur via specific receptors, integrins, which are transmembrane heterodimeric transmembrane glycoproteins composed of two of 26 subunits. Integrins bind to intracellular cytoskeleton and secondary messenger systems and extracellularly to other cells and the ECM [32]. They are closely associated with glutamate receptors and various ion channels. Integrins regulate multiple processes including synaptic plasticity, neuronal migration and development, axonal growth and synaptogenesis. They are also involved in angiogenesis.

### 20.5.2 Changes in the ECM in Epilepsy

There are persistent changes in multiple components of the ECM during the development of epilepsy. Peri-neuronal net components, including aggrecan, neurocan, hyaluronan, tenascin-R and some of the linking proteins, decrease during epileptogenesis; a progressive decrease in perinuronal nets is associated with a progressive decrease in inhibition and the occurrence of seizures (months after traumatic brain injury) [53, 62]. In addition, degradation of the ECM may permit aberrant neuronal and synaptic reorganisation. ECM remodelling and the increased secretion of proteases may also contribute to this process. There is robust evidence that expression of MMP-9 is increased during epileptogenesis, and that this increase may promote kindling [54]. Other serine proteases are also up-regulated in epilepsy including urokinase-type plasminogen activator (uPA) and its receptor (uPAR) [38]. Intriguingly, uPAR up-regulation may be

protective as uPAR knockouts develop a more severe epilepsy phenotype following status epilepticus [56]. This indicates that some of the changes of ECM during epileptogenesis may be adaptive rather than pathogenic.

In addition, mutations in the gene encoding SRPX2 (Sushi-repeat Protein, X-linked 2), one of the ligands of uPAR, results in bilateral perisylvian polymicrogyria and epilepsy in humans [66]. Integrin expression is also increased during epileptogenesis and in pathologies associated with the development of epilepsy [87].

Lastly, an extracellularly secreted molecule, leucine rich, glioma-inactivated 1 (LGI1) has been strongly associated with epilepsy [8, 20, 33, 58, 69]. LGI1 interconnects presynaptic disintegrin and metalloproteinase domain-containing protein 23 (ADAM23) to postsynaptic ADAM22 at the synaptic cleft. LGI1 is important for trafficking and kinetics of a presynaptic potassium channel, Kv1.1, and also for trafficking of post-synaptic AMPA receptors. In humans, mutations in LGI1 cause autosomal dominant lateral temporal epilepsy or autosomal dominant partial epilepsy with auditory features with onset in childhood/adolescence [58]. In addition, autoantibodies directed against LGI1 have been shown to underlie limbic encephalitis and temporal lobe seizures in humans [69].

Overall, there is growing evidence for the importance of the ECM in epileptogenesis, plasticity and determining network excitability. Further studies aimed at modeling disruption and reorganization of the ECM will be important for a greater understanding of the epileptogenic process. Moreover, the ECM provides an ideal target for therapies aimed at disrupting epileptogenesis and modifying established epilepsy, as it is extracellular and so easily accessible to drugs and has multiple downstream effects, regulating receptors, channels and synaptic transmission.

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## 20.6 Conclusions

There is burgeoning evidence to support a critical role for non-neuronal mechanisms in epileptogenesis and the generation of seizures.

Both animal experiments and experiments of nature (gene mutations) indicate that pathology of non-neuronal elements are sufficient for epileptogenesis. However, most of our present therapies are neurocentric, indicating that there may be enormous undiscovered therapeutic potential in targeting these non-neuronal elements. Moreover, it is a concern that many of the large scale mathematical models of brain function (e.g., the blue brain project [52]) have thus far ignored the role of these non-neuronal constituents.

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## What Epilepsy Comorbidities Are Important to Model in the Laboratory? Clinical Perspectives

Simon Shorvon

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### Abstract

In recent years, there has been a focus on studies of comorbidity in epilepsy. The concept of epilepsy comorbidity is complex. This is partly because epilepsy is essentially a symptom for which there are many underlying causes, with multiple genetic and environmental influences. These causal conditions themselves carry comorbidities which vary from condition to condition. The fact that some psychiatric comorbidities are ‘bidirectional’ complicates this further. These issues reduce the usefulness of any unitary study of ‘epilepsy comorbidity’. Epilepsy comorbidities can be divided into direct/indirect and somatic/psychiatric categories. Only some aspects are susceptible to experimental modeling. This chapter briefly reviews the clinical studies of cause, frequency, epidemiology and mortality of comorbidities, and their use as biomarkers for epilepsy.

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### Keywords

Epilepsy • Epileptic seizures • Somatic comorbidity • Psychiatric comorbidity

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### 21.1 Definitions and Divisions of Epilepsy Comorbidities

The term comorbidity has been said to have been first coined by Feinstein [7] to define the co-existence of different diseases or conditions. The original studies of epilepsy comorbidity emphasized migraine, psychiatric disorders and vascular disease, but since the early 2000s there

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has been a greater focus on this problem and more recent studies have demonstrated a wider range of comorbid disorders and have attempted to define their extent [10, 14, 15, 18] and underlying mechanisms [3, 11, 22, 31]. Some of the comorbid conditions are susceptible to experimental modeling and others are not. In broad terms, experimental or animal models are most appropriately employed to investigate the mechanisms and the causes of comorbidity.

The concept of ‘comorbidity’ of a condition such as epilepsy is complicated. As a rider to any discussion, it should of course be realized that epilepsy is essentially a symptom for which there are many underlying causes, with multiple genetic and environmental influences [25, 28]. These causal conditions themselves carry comorbidities, which vary from condition to condition, thus complicating any broad study of ‘epilepsy comorbidity’. Studies in epilepsy have divided and defined the range of comorbidities in a number of different ways.

- (i) Direct/indirect: The direct comorbidities are those that are due to epilepsy. The indirect comorbidities are those that are due to underlying causes of the epilepsies or to risk factors which are shared with epilepsy.
- (ii) Psychiatric/somatic: The psychiatric comorbidities refer to the primary psychiatric diseases and the somatic comorbidities to systemic and neurological disease.

It is probably not surprising to know that these divisions are artificial and there are in each system grey areas where conditions overlap or are not easy to pigeon-hole. Understanding comorbidity is important for various reasons:

- (i) The comorbidities may have an important influence on prognosis of epilepsy (including mortality) and indeed often have a greater influence than the epilepsy itself.
- (ii) The therapy of epilepsy may be influenced by their presence (as well as the fact that some comorbidity is due to therapy)
- (iii) The comorbidities may have diagnostic implications in some situations
- (iv) Doctors dealing with epilepsy should be alert to the risk of comorbidities as these too may require treatment

- (v) The comorbidities of epilepsy may in many instances cause more distress and dysfunction than the epilepsy itself.

There is also often a two-way relationship between comorbidity and epilepsy (often known as a ‘bidirectional’ relationship; discussed further below). Comorbidity can affect the course of the epilepsy directly (via organic effects on the brain) or indirectly (via chronic ill health, side-effects of treatment, secondary psychiatric effects). Comorbidities also affect health care utilization, and all the outcomes of epilepsy including mortality.

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## 21.2 Causes of Comorbidity in Epilepsy

‘Direct comorbidity’ is that due to the epilepsy or the effects of seizures themselves. Examples of seizure-related comorbidity are fractures due to falls in seizures, or memory disturbance due to cerebral damage. Laboratory studies offer the opportunity for prevention and especially neuroprotection and these are topics which can be modeled experimentally. Other direct morbidity is due to the secondary handicap of epilepsy which includes chronic ill health, psychiatric problems, social drift and other pressures. These are topics which cannot be studied in laboratory models.

The indirect comorbidities may be: (a) due to the underlying causes themselves, such as stroke or cerebral tumour, which cause epilepsy and also other effects; (b) due to shared risk factors which have been shown to predispose to epilepsy and also to other medical condition, examples include vascular disease which predisposes to stroke, or alcoholism which predisposes to head trauma. Sometimes the risk factors are genetic (discussed further below); (c) due to the treatment of epilepsy, examples include hepatic or bone disease, or interactions between medications; (d) conditions where the mechanisms underlying the association are quite unknown for instance associations with asthma, bowel disease or thyroid disease. Many of these aspects can be studied experimentally.

Psychiatric comorbidity (which can be both direct and indirect) is particularly complex with genetic, environmental, shared underlying causes and also treatment and direct cerebral damage all potentially contributing to the epilepsy and the comorbidities.

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## 21.3 Epidemiology and Frequency of Comorbidities of Epilepsy

There have been a number of large scale surveys of comorbidity, based on National Health Service statistics.

### 21.3.1 Somatic Comorbidity

The first database to be mined was the UK General Practice Research Database (GPRD) which covered a period between 1995 and 1998 [11]. Data were based on a population of 1.3 million, in which all ICD codes (codes defined by the International Classification of Disease (ICD) were recorded and of these 1,041,643 adults were studied. 5,834 persons with epilepsy were identified. The most common somatic conditions in adults with epilepsy were: fractures (10 %), with highest rates in women older than 64 years (17 %); asthma (9 % with the highest rates (11 %) in younger women); and migraine (8 %). Amongst the oldest patients, the most common somatic comorbidities were diabetes (9 %), transient ischaemic attacks (18 %), ischaemic heart disease (14 %), heart failure (12 %), neoplasia (7 %), and osteoarthritis (12 %). The most common neurologic disorders in this age group were brain degenerative diseases (14 %) and Parkinson's disease (4 %). The importance of environment is also shown by the study of Babu et al. [2] in India which showed increased rates of neurocysticercosis, sleep disorders, and tuberculosis compared with controls.

Télliez-Zenteno et al. [32] have used data obtained through two door-to-door Canadian health surveys, the National Population Health Survey (NPHS, N=49,000) and the Community Health Survey (CHS, N=130,882), covering

98 % of the Canadian population. They found that those with epilepsy had a statistically significant higher prevalence of many chronic conditions when compared to the general population; those conditions which occurred twice as often or more were (proportional risk): stomach/intestinal ulcers (CHS 2.5, NPHS 2.7), stroke (CHS 3.9, NPHS 4.7), urinary incontinence (CHS 3.2, NPHS 4.4), bowel disorders (CHS 2.0, NPHS 3.3), migraine (PR, CHS 2.0, NPHS 2.6), Alzheimer's disease (NPHS 4.3), and chronic fatigue (CHS 4.1). Of course several of these conditions are causal conditions of epilepsy (stroke, Alzheimers disease) and so it is not at all surprising that they cluster with epilepsy in population surveys, but the others were more surprising. It was postulated by the authors that gastro-intestinal diseases may be due drug therapy or autonomic ictal effects, although both explanations seem unconvincing.

### 21.3.2 Psychiatric Comorbidity

The commonest comorbidities of epilepsy are psychiatric. There are a number of epidemiological studies of comorbidities looking at this association. In the study mentioned above, Gaitatzis et al. [11] found the commonest psychiatric conditions in adults with epilepsy were: depression (18 %), anxiety (11 %) and psychosis (9 %). Overall, 41 % of patients with epilepsy received a psychiatric diagnosis at some point during the 3-year study period. Télliez-Zenteno et al. [32] used data from the Canadian Community Health Survey (CCHS) to compare the rates of psychiatric disease in those with and without epilepsy. The CCHS included 36,984 subjects. Those with a history of epilepsy reported higher lifetime anxiety disorders (odds ratio (OR) 2.4, 95 % confidence intervals (CI)= 1.5–3.8) or suicidal thoughts (OR 2.2 (1.4–3.3)). Surprisingly, the risk of major depressive disease or of panic disorder/agoraphobia were not greater in those with epilepsy (and may throw some doubt upon the methodology of this study).

There are also a number of case control studies, looking both at the frequency of epilepsy in

psychiatric populations and vice versa (the so-called bidirectional relationship). A variety of mental disorders, alcoholism and dementia are found more commonly in patients with epilepsy than in non-epileptic controls. The strongest associations of epilepsy are with major depression, bipolar disease, and schizophrenia. Major depressive episodes are more common in patients with epilepsy than in the general population, with prevalence ranging from 11 to 62 %, compared with 3.7–6.7 % for the general population [6, 8, 17, 23, 32]. There is an even stronger association with psychosis; the prevalence of the interictal psychosis of epilepsy ranges (in different studies) between 4.3 and 44 % and in a recent review, rates of 19.4 % and 15.2 % in generalized epilepsies and temporal lobe epilepsy groups are recorded.

The association of neurological and psychiatric disorders to epilepsy is complex. The fact that there is an association was fully recognized in the late nineteenth century and the concept of the ‘Neurological Trait’ was universally accepted [19, 26, 27]. According to this concept, epilepsy was an essentially inherited condition and inherited together with other neurological and psychiatric disorders. It was accepted that within a family the same inherited tendency might manifest in one person as epilepsy and in other family members as other conditions, but all reflected the same underlying inherited influence (of course, “genes” were not recognized, nor were Mendelian principles widely known at this time). Although different authorities included different conditions within the inherited tendency, at the core were mental disturbances such as insanity, mental retardation, behavioral aberrations, alcoholism – and epilepsy. Gowers, for instance, in 1881 wrote: “There are few diseases in the production of which inheritance has great influence.... It is well known that the neuropathic tendency does not always manifest itself in the same form.... The chief other morbid states (besides epilepsy), in which the neuropathic tendency is manifest are insanity, and, to a much smaller degree, chorea, hysteria, and some forms of disease of the spinal cord. Intemperance is probably also due, in many cases, to a neuropathic disposition” [13]. In Gowers’ personal series of 1,218 epilepsy cases, he found that 42 %

“presented evidence of neurotic inheritance.” In the nineteenth century, the concept was also linked to that of ‘degeneration’ and it was widely believed that the manifestations of the trait worsened as it was inherited from generation to generation.

Another topic of current interest is the “bidirectional nature” of the comorbidity epilepsy with various neuropsychiatric conditions. The association is often considered to be due to such factors as recurrent epileptic seizures, social stigma, adverse effects of drug treatment or the underlying structural or metabolic brain injury. However, recent studies have shown that the ‘bidirectionality’ may in fact predate the development of epilepsy [1, 4, 24] and be due to shared genetic propensities. Qin et al. [23] found a family history of epilepsy to be a risk factor for schizophrenia or schizophrenia-like psychosis, even after adjusting for personal history of epilepsy. Similarly, adults with new-onset epilepsy are seven times more likely to have a prior history of depression. Adults and children with newly diagnosed epilepsy have been noted to have a prior history of attempted suicide which is five times that of the general population. One development in the field was the finding that copy number variants (CNVs) underpin the pathogenesis of some neuro-developmental disease. Several studies have demonstrated that the same large CNVs underpin epilepsy, autism, schizophrenia, mental retardation and attention deficit hyperactivity disorder [1, 4, 5, 16, 20, 21, 24, 30, 33, 34].

If there are shared genetic influences, both the epilepsy and the neuropsychiatric conditions are frequently ‘neurodevelopmental’ in origin [19]. *Functional annotation analysis* is one attempt to understand shared pathogenic mechanisms, and the effect of the dimension of time is another factor which complicates analysis and renders simple ‘gene hunts’ unlikely to be very revealing. The reasons for this are the differing gene expression at different times, the effect of development of the activation of functional genetic pathways and the strong effect of environmental factors and chance in development (see 20 for further discussion of this point). The genetic mechanisms of these shared propensities (which has eerie

echoes of the concept of the “neurological trait”) are the subject of study and certainly can be modeled experimentally – this could be an area of promising future research.

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## 21.4 Mortality Associated with Comorbidity

The risk of death in epilepsy is elevated even when the epilepsy is in remission. In our own recently published 25-year follow-up study of people with newly diagnosed epilepsy the risk of premature death was twice that of the general population [29]. The underlying causes of epilepsy (stroke, brain tumour etc.) have an obviously increased rate of mortality. This not surprisingly increases the risk of premature mortality amongst those with epilepsy. Of more interest from the point of view of studies of comorbidities, are the ‘external’ causes of mortality in epilepsy (ie not underlying causes of epilepsy such as brain tumours or strokes) and the risk of premature mortality due to such causes.

An outstanding study in the field was recently published examining the relationship of psychiatric comorbidity to premature death [9]. This is the gold standard study in the area, and outshines all the others in terms of its comprehensive nature and intelligence. Data were obtained from all individuals born in Sweden between 1954 and 2009, via a variety of nationwide population registers in Sweden which were then linked: the Patient Register, the Censuses from 1970 to 1990, the Multi-Generation Register, and the Cause-of-Death Register. Epilepsy was identified through the National Patient Register, which includes individuals hospitalized or having outpatient appointments with specialist physicians in Sweden who had received a diagnosis of epilepsy (n=69,995). Data for causes of death were retrieved for all individuals who died between 1969 and 2009 from the Cause of Death register based on death certificates, which covers over 99 % of all deaths. Patients were compared with age-matched and sex-matched controls (n=660,869) from the general population as well as unaffected siblings (n=81,396). 6,155 (8.8 %)

people with epilepsy died during follow-up. The study had extensive sensitivity testing and the comparison with unaffected siblings was important for exploring interfamilial confounding.

The study found a very substantially elevated risk of premature death in epilepsy. The odds ratio for premature mortality was 11.1 [95 % CI=10.6–11.6] compared with general population controls, and 11.4 [10.4–12.5] compared with unaffected siblings. 15.8 % of the deaths were due to external causes. The external causes with the highest odd ratios were non-vehicle accidents (OR 5.5, 95 % CI 4.7–6.5) and suicide (3.7, 3.3–4.2). Of those who died from external causes, 75.2 % had comorbid psychiatric disorders, with the strongest associations being with depression (13.0, 10.3–16.6) and substance misuse (22.4, 18.3–27.3). This link between premature morbidity and psychiatric disease is of course of fundamental importance in clinical practice. Epilepsy was found in this study to be an independent risk factor for all-cause and external causes of death, a finding which was most clearly shown by the comparison of patients with their unaffected siblings, with the rate of mortality increased by 2.9× for suicide and 3.6× for accidents. Another important point recognized was that despite the high relative risks (odds ratios), the absolute rates of premature mortality from external causes was only 1.4 %. However, about a third of the epilepsy patients had at least one comorbid psychiatric diagnosis and about 10 % exhibited substance misuse [9].

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## 21.5 Comorbidity as Biomarker for Epilepsy

If comorbidities are common and easily measured, they can be used as biomarkers for epilepsy. Examples might include genetic markers or even physiological changes. The study of epilepsy biomarkers is not yet well developed, but biomarkers potentially have great importance for diagnostic purposes and also for prognosis and for studying the effects of therapy. Galanopoulou and Moshe [12] divided the search for biomarkers into four categories. Some are not likely to be

susceptible to experimental study, but others are. Two of the categories defined were (in their own words):

Biomarker of epileptogenicity: The desired features of these biomarkers include:

- Specificity in differentiating the epileptic state from reactive changes resulting from an initial precipitating event or the first seizure, and from developmental processes that have not yet reached maturity;
- Sensitivity in diagnosing epilepsy at the pre-clinical or early symptomatic stages, when clinical diagnosis has not yet been established;
- Ability to detect the reversal of epileptogenicity, to prevent unnecessary continuation of treatments.

Biomarkers of treatment implementation, tolerability or toxicity: Many antiepilepsy drugs have side effects, which result in comorbidity, the mechanisms for which offer the possibility of biomarker studies:

- Provide target identification for treatment selection, distinguishing it from age-specific relevant processes;
- Define the timing and therapeutic window of treatment administration, based on age- and sex-adapted criteria;
- Distinguish the treatment-responsive from the resistant patient populations early;
- Provide early risk identification and monitoring of treatment-related toxicities, based on age- and sex-adapted criteria, with sufficient specificity for the administered treatment;
- Have the ability to localize the epileptogenic focus accurately and facilitate more effective ablative treatments, if medical treatments are not curative.

## 21.6 Experimental and Animal Models of Comorbidity

As emphasized above, only some aspects of epilepsy comorbidity are susceptible to modeling in the experimental laboratory. Experimental studies which are most likely to be successful are those directed at the causal molecular, physiological and/or genetic mechanisms of the

relationship of epilepsy and its comorbid conditions. The relationship, especially for the psychiatric (and other brain-related) comorbidities are likely to be complex and have developmental and time-sensitive dimensions. Those comorbidities that have priority are, in the author's opinion, in the following areas:

- (a) Studies of the adverse effects of epilepsy on brain function, with experimental studies that focus on the mechanisms of brain damage and the role of neuroprotection
- (b) Studies of the adverse effects of epilepsy on somatic function, with experimental studies that focus on the molecular mechanisms of these effects and ways of blocking these (the role of osteoporosis for instance in fractures).
- (c) Studies of the underlying mechanisms of psychiatric comorbidities, with experimental studies focusing on the genetic and molecular basis, the bidirectionality of the relationship between epilepsy and comorbidities, and on shared pathways.
- (d) Studies of the role of comorbidity as biomarker of either epileptogenicity or of the adverse effects of treatment. These two may have a developmental or age-related effects, with different vulnerabilities at different ages.

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## Abstract

An epilepsy comorbidity is a condition or disorder that occurs at a frequency greater than chance in a person with epilepsy. Examples of common epilepsy comorbidities are depression, anxiety, and intellectual disability. Epilepsy comorbidities can be quite disabling, sometimes affecting a patient's quality of life to a greater extent than seizures. Animal models offer the opportunity to explore shared pathophysiological mechanisms, therapeutic options, and consequences of both the epilepsy syndrome and a given comorbidity. In this chapter, depression is used as an example of how animal models can inform translational questions about epilepsy comorbidities.

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## Keywords

Epilepsy • Comorbidity • Depression • Animal models

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## 22.1 What Are Epilepsy Comorbidities and Can Animal Models Help?

While epilepsy is primarily considered to be a condition of recurrent, unprovoked seizures, it is increasingly evident that epilepsy involves a lot more than seizures. Epilepsy comorbidities,

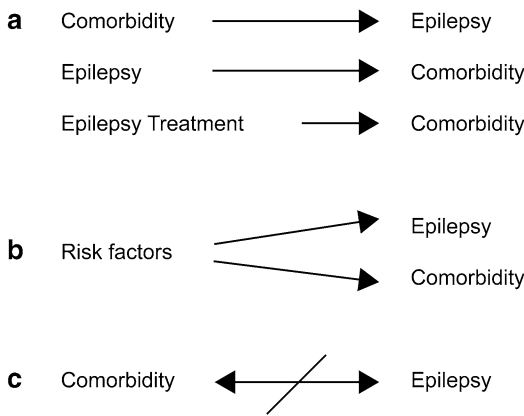
defined as medical or psychiatric disorders that occur at a frequency greater than chance in a patient with epilepsy, play a crucial role in the quality of life and treatment effectiveness in patients with epilepsy. Epilepsy comorbidities include disorders of cognition, mood, and behavior [3, 14, 26], as well as a variety of medical and neurological disorders [10]. Specific examples include depression, anxiety, intellectual impairment, autism, sleep disorders, migraine, and many others (Table 22.1). In some individuals, comorbidities can be more impairing than the seizures themselves [11]. Many patients have more than one comorbidity, underscoring the need to understand the roles played by single and multiple

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**Table 22.1** Examples of epilepsy comorbidities

Anxiety disorder
Autism spectrum disorder
Cardiovascular disease/stroke
Dementia/Alzheimer disease
Depression
Intellectual disability/cognitive impairment
Migraine
Sleep disorders
Suicidality



**Fig. 22.1** Possible relationships between epilepsy and comorbidity. (a) A causal relationship might exist whereby epilepsy or an epilepsy treatment might cause a comorbidity, or a comorbidity might lead to epilepsy. (b) Common risk factors might exist (e.g., environmental, genetic, structural) that lead to both epilepsy and a comorbidity. (c) Comorbidity and epilepsy might be independent, unrelated associations

comorbidities in epilepsy, epileptogenesis, and quality of life in persons with epilepsy.

The concept of epilepsy comorbidity has been under-recognized but is not new. Recent attention has been focused on epilepsy comorbidities with the addition of comorbidities as a separate NIH Epilepsy Benchmark [23]. In addition, recognition of a comorbidity as a major cause of impaired quality of life of persons with epilepsy is elaborated in the recent Institute of Medicine report [5].

There are several possible relationships between epilepsy and a comorbidity (Fig. 22.1) [3, 10]. First, the relationship can be causal, with one disorder causing the other or making the other disorder more likely (Fig. 22.1a). That is, a comorbid condition can lead to epilepsy, or conversely,

the comorbid condition occurs as a result of the epilepsy or its treatment. Several examples will clarify this concept. The etiologies of most symptomatic epilepsies (for example, traumatic brain injury, stroke) correspond to this cause-and-effect model [10]. Conversely, epilepsy itself can lead to a comorbidity such as anxiety disorder in predisposed individuals [34]. Finally, numerous examples of epilepsy treatments leading to behavioral comorbidities can be cited, such as the association of phenobarbital with hyperactivity in children and the association of levetiracetam with altered mood [35].

Second, shared risk factors, which can be genetic, metabolic, structural or environmental, can lead to the development of both epilepsy and a comorbidity (Fig. 22.1b). An example is the structural brain damage caused by perinatal hypoxia-ischemia that leads to both epilepsy and comorbid cerebral palsy [3]. This type of relationship also includes the comorbidities that are considered “bidirectional”, that is, common underlying mechanisms could facilitate the development of both epilepsy and the comorbidity. Depression is a common and critically important example of an epilepsy comorbidity and is discussed in detail below. Third, the relationship between epilepsy and a comorbidity could be incidental or even spurious (Fig. 22.1c).

Since so many people with epilepsy harbor one or more comorbidities, it is important to elucidate these relationships. For example, there could be shared pathophysiological mechanisms between epilepsy and a comorbidity, with the possibility that one or both conditions is amenable to a treatment or disease modification that exploits these common mechanisms. Of note, no specific therapy exists for a comorbidity in the context of epilepsy. That is, if a patient with epilepsy is diagnosed with a comorbidity such as anxiety or depression, treatment choice is limited to medications used to treat anxiety or depression, irrespective of the concurrent epilepsy. Novel treatments are needed that take into account the specific pathogenic mechanisms of both epilepsy and the comorbidity.

Given the prevalence of epilepsy comorbidities and the lack of understanding of their mechanisms,

**Table 22.2** Factors to consider in animal models of epilepsy comorbidities

Age of onset (of seizures and comorbid symptoms)
Brain region and neurotransmitter system underlying comorbidity
Environmental factors (e.g., cage size and density, light/dark cycle)
Food intake (e.g., may be decreased in depressed animals)
Gender of animal
Handling by laboratory personnel
Species/strain/genetic background
Symptoms versus syndrome (i.e., concurrent additional comorbidities)

**Table 22.3** What can be learned from studying epilepsy comorbidities in animal models?

Mechanisms of shared pathophysiology
Potential avenues for therapy and disease modification (e.g., relative roles of antidepressant and anticonvulsant medications on both epilepsy and depression)
Correlations between behavioral phenotype of the comorbidity and features of the epilepsy syndrome (seizure type, frequency, temporal relationship with comorbid symptoms, etc.)
Role of the comorbidity in epilepsy progression and epileptogenesis

the question arises as to whether animal models can provide useful information about pathogenesis or treatment [18]. The purpose of this chapter is to provide an overview of some of the theoretical issues in modeling epilepsy comorbidities in animals, followed by an example of how understanding one specific epilepsy comorbidity – depression – might enhance understanding of the pathophysiology of both disorders and could help to identify treatment targets. Comprehensive reviews of comorbidities in animal models of epilepsy already exist [3], as do detailed guidelines for testing specific cognitive functions in animal models of epilepsy [45].

The first question to consider is how closely an animal model resembles the human condition. This question applies to epilepsy as well as to the comorbidity, and when trying to model both conditions in one animal, obvious challenges arise (Table 22.2). Species differences are usually obvious, but not trivial. While at first glance, it

might seem implausible that a rodent could exhibit depression similar to that experienced by a patient. However, a burgeoning literature supports the idea that there are shared features and pathophysiological mechanisms between depression in animals and humans (discussed in greater detail below). Second, for any comorbidity under consideration, the experimenter must evaluate how the testing paradigm itself might contribute to the animal's performance; that is, does the test itself elicit stress or another set of behaviors that confound the original intention? Third, it is critical that longitudinal observations be employed – it is insufficient to test an animal only once in a behavioral paradigm since both epilepsy and most comorbidities are chronic (and often evolving) conditions (Table 22.3).

## 22.2 Depression as an Example of an Epilepsy Comorbidity

Depression is extremely common in the general population, but even more so among people with epilepsy [22]. In population-based studies, it has been estimated that approximately 25–35 % of individuals with epilepsy suffer from depression (even higher if the epilepsy is not well controlled) and that people with depression have a 3- to 7-times greater risk of developing epilepsy than the general population [15, 20, 47]. Depression also affects 8–26 % of children with epilepsy [9, 37]. These percentages far exceed those expected in the general population and may well underestimate the actual prevalence of depression in persons with epilepsy. A history of depression is a reliable predictor of worse epilepsy severity [20]. The bidirectional relationship of epilepsy and depression (epilepsy is more likely in people with depression, and depression is more likely among people with epilepsy) is validated by neurobiological data of several types, including neurotransmitter analyses, MRI and positron emission tomography studies of temporal or frontal lobe function, and investigations of hypothalamic-pituitary-adrenal (HPA) axis dysfunction [21]. The bidirectional relationship suggests that there may exist one or more common

neurobiological mechanisms and that these mechanisms might be exploited for therapeutic advantage.

Depression is a heterogeneous disorder with several distinct subtypes classifiable using the Diagnostic and Statistical Manual of Mental Disorders (5th edition, DSM-V [1]). It is important to recognize that the DSM is based on expert consensus not validated biomarkers. DSM-V criteria for the diagnosis of depression include despair, anhedonia (inability to experience pleasure), vegetative symptoms (weight loss, appetite decrease or increase, decreased energy, insomnia), feelings of worthlessness and guilt, decreased focus/attention span, and suicidal ideation. It is uncertain whether depression in persons with epilepsy is identical to depression in persons without epilepsy. Data suggests that many “atypical” features that do not adhere to the strict DSM criteria typify depression in individuals with epilepsy [20]. Atypical features include a greater degree of anxiety, irritability, and mood lability. Importantly, the timing of depressive episodes may relate to seizure occurrence; a bout of depression may precede a seizure (interictal episode) or occur around the same time as a seizure (peri-ictal episode) [22]. Despite their frequent co-occurrence, the severity of depression, at least in temporal lobe epilepsy, is not proportionate to the number of seizures [12]. The treatment goal is reduction of both seizures and depressive symptoms, although seizure control does not always correlate with improvement in depression [13]. Ideally, this goal would be achieved using monotherapy, with one drug improving both seizure control and depression. Specific data about the impact of antidepressants on depression in epilepsy are scarce but much needed.

The effects of antidepressants on epilepsy and antiepileptic agents on depression are complex. Some antiepileptic drugs are well known for their mood stabilizing properties (e.g., carbamazepine, valproate, lamotrigine). Likewise, antidepressants have been shown to exert anticonvulsant effects in both patients and animals – selective serotonin reuptake inhibitors (SSRIs), serotonin-norepinephrine reuptake inhibitors (SNRIs), and tricyclic antidepressants (TCAs) can increase brain

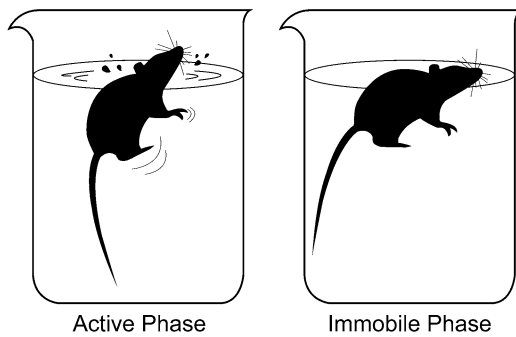
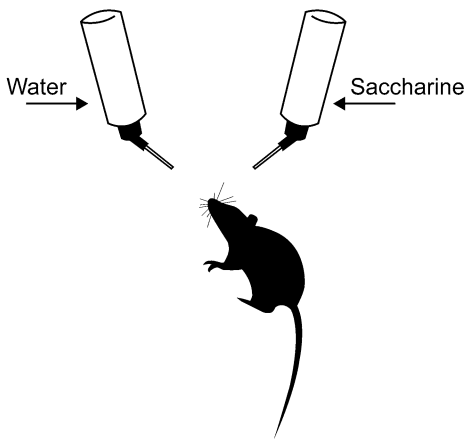
monoamines such as serotonin, norepinephrine, and dopamine, favoring an anticonvulsant action [16]. In addition, depression can be ameliorated by alterations of the primary neurotransmitter systems of the brain – glutamate receptor antagonists (e.g., dizocilpine, ketamine) or  $\gamma$ -amino-butyric acid (GABA)-receptor agonists [33, 40]. The multifaceted effects of these and other novel agents in epilepsy and depression are poised for study in animal models [28].

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### 22.3 Evaluating Depression in Animal Models of Epilepsy

Depression is an exemplary disorder in which to explore the opportunities and challenges between epilepsy and a comorbidity using animal models [8]. Obviously, many subjective symptoms of depression cannot easily be extrapolated to animals, but an approximation of some of the symptoms makes the study of this comorbidity in animals quite tenable. To that end, a set of modified criteria for depression in rodents has been proposed [3]. Of the depression criteria listed in DSM-V, despair and anhedonia are most readily testable in animals, with validated laboratory tests available for those symptoms. The forced swim test (FST) is a measure of despair in rodents, while anhedonia, the failure to experience pleasure, is assessed by the taste preference test (TPT). These tests have been widely used to screen potential antidepressant compounds. It is important to recognize that a single administration of those or any other experimental measure of depression in animals represents only a single point in time, whereas a comorbidity typically evolves over time, necessitating serial assessments.

The FST is performed by placing an animal in a water-filled chamber with smooth sides, from which it cannot escape. Initially, the animal typically swims around frantically, trying to escape by climbing the walls (active escape phase). Eventually, the animal seems to give up this futile effort and becomes immobile, simply floating in the water, striving to keep its head above water to prevent drowning (immobility phase). These two phases are easily quantified, with the immobility

**a** Forced swim test**b** Taste preference test

**Fig. 22.2** Laboratory tests of depression in rodents. (a) The forced swim test is a measure of despair, one of the core symptoms of depression. *Left*, active escape phase. *Right*, immobility phase, considered to represent despair. (b) The taste preference test is a measure of anhedonia, or loss of ability to experience pleasure. A normal animal prefers the sweetened liquid, while a depressed animal does not express this preference

phase comprising a validated measure of despair (Fig. 22.2a). In models of depression, animals that are depressed have shorter active escape phases and enter the immobile phase sooner. Importantly, the FST is itself a stressor for an animal. Antidepressant drugs increase the active escape phase duration, supporting the contention that the immobility phase represents despair. The FST has been used for antidepressant drug discovery in animal models of depression, but pharmacologic studies of antidepressants in epileptic animals have emerged only recently.

The TPT compares a rodent's preference for drinking a solution sweetened with saccharine (or sucrose) over plain water (Fig. 22.2b) [32]. Ordinarily, rodents prefer to drink the sweet solution. Sugar consumption stimulates dopaminergic fibers projecting from the ventral tegmental area to the nucleus accumbens, where the amount of dopamine released correlates with motivational aspects of reward [39]. In depressed animals, intake of sweet liquids such as sucrose or saccharine is decreased and there is no difference in rodents' consumption of the sweetened versus plain water, suggesting that they have less interest in the flavored fluid.

Other tests have also been employed for comorbidities in epilepsy research, some applicable to depression and others more reflective of anxiety, cognitive function, memory, or learning. Comprehensive lists of such tests (Table 1 in [3] and Table 1 in [8]) reveal that many are in need of validation in animals with epilepsy. As well, there is an urgent need for multi-dimensional behavioral tests to simultaneously assess concurrent comorbidities in the same subject – depression, anxiety, sleep dysfunction, etc.

## 22.4 Examples of Comorbid Epilepsy and Depression in Animal Models

To illustrate some of the insights that can be gained from animal models, examples are now provided that examine various aspects of the relationship between epilepsy and depression. These examples include both acquired and genetic etiologies. Space precludes detailed discussion of other relevant examples such as GAERs (genetic absence epilepsy rats from Strasbourg) [19] and genetically epilepsy prone rats (GEPRs) [17].

Chemoconvulsant models of temporal lobe epilepsy (TLE) in rats using either kainic acid (KA) or lithium/pilocarpine (LiP) allow detailed study of the relationship between seizures (number, frequency, duration and timing of spontaneous recurrent seizures) and the occurrence of behavioral and cognitive abnormalities. KA is a glutamate receptor agonist; pilocarpine is an agonist of

muscarinic acetylcholine receptors. Both forms of chemoconvulsant-induced epilepsy mimic limbic epilepsy, with initial status epilepticus followed weeks-to-months later by spontaneous recurrent seizures and behavioral and cognitive abnormalities. In both KA- and LiP-epilepsy, evidence of depression has been documented on the FST and TPT [24, 36]. Following KA-induced status epilepticus, rats had shorter latencies to the immobile phase on the FST and longer duration of immobility, suggesting that these rats were depressed (increased despair) [24]. Using microarray analysis, it was shown that depressed rats had a reduction in expression of the gene for serotonin receptor 5B. Most strikingly, environmental enrichment prevented both FST abnormalities and the underlying gene expression changes, suggesting that environmental factors play a crucial role in the development of depression as an epilepsy comorbidity. Investigation of structural brain injury and the roles of antidepressant and anticonvulsant drugs in this model would further clarify these relationships.

In the other chemoconvulsant model, intraperitoneal injection of LiP causes limbic status epilepticus, followed in subsequent weeks by behavioral deficits such as learning and memory impairment and a depression phenotype. Compared to naïve rats, LiP-treated rats demonstrated increased immobility time in the FST and loss of taste preference in the TPT [29], supporting the depression phenotype of despair and anhedonia. These behavioral deficits were rescued by treatment with a blocker of the serotonin 5HT1a receptor, but there was no effect of selective serotonin reuptake inhibitors (SSRIs) [29], suggesting that depression in this model does not respond to medications typically used to treat clinical depression. These observations support the conclusion that depression, at least in some epilepsy disorders, represents an atypical form of the condition. This model provides the opportunity to dissect contributions of the multiple serotonin receptors involved in various depression subtypes [27]. The effects of standard anticonvulsants on depression and antidepressants on epilepsy have not yet been reported in this model.

To investigate the mechanism linking depression to epilepsy in this model, the authors noted that dysregulation of the HPA axis is a marker of depression, with increased levels of plasma glucocorticoid (cortisol) due to loss of negative feedback of cortisol on corticotrophin releasing hormone and adrenocorticotrophic hormone release [25]. LiP-treated rats had elevated cortisol levels, supporting the depression phenotype [31]. After status epilepticus in these animals, there was reduced serotonergic innervation from brainstem raphe nuclei to the hippocampus due to upregulation of raphe 5-HT1A autoreceptors, as found in some human depression [4]. Furthermore, a blocker of 5-HT1A receptors, WAY-100635, improved performance on the FST, forming a link between abnormal serotonergic function, depression, and behavior [30].

Further studies showed that increased hippocampal interleukin 1- $\beta$  (IL1 $\beta$ ) signaling might mediate both depressive symptoms and heightened hippocampal excitability leading to spontaneous seizures in this model. The authors proposed a scheme whereby epilepsy leads to depression by increasing IL1 $\beta$  signaling, which upregulates raphe 5-HT1A autoreceptors, compromising raphe-to-hippocampus serotonergic neurotransmission. These findings raise the possibility of a link between mechanisms of epilepsy, depression, stress, and the inflammatory response [49]. Potential loci for intervention might include blockade of glucocorticoid action, downregulation of raphe 5HT1A autoreceptors, or anti-inflammatory agents. This model can also be utilized to further characterize the mechanisms of neuronal excitability underlying epilepsy and depression.

The next example is rats bred for susceptibility or resistance to depression-like behaviors during swimming in the FST (named SwLo and SwHi, respectively). SwLo rats display increased immobility in the FST and anhedonic tendencies. Importantly, SwLo rats also have increased predisposition to limbic seizures induced by kainic acid or pilocarpine, providing an excellent opportunity to examine the joint mechanisms of depression and epilepsy, with particular relevance to temporal lobe epilepsy [7, 46]. Chronic antidepressant treatment reverses the FST deficits in

SwLo rats [50], substantiating the validity of this model in depression. In addition, the existence of the converse model – SwHi rats that are resistant to depression – provides a unique opportunity to examine whether this strain is also relatively resistant to seizure development. To date, there are no data regarding the effects of anticonvulsants on either depression or seizure development in this model. Finally, this model provides further evidence for the interaction of environment and genetics in the expression of both depression and epilepsy, as aerobic exercise was found to improve both FST performance and seizure resistance in SwLo rats compared to SwHi rats [6].

Lastly, a genetic model of absence epilepsy has revealed a number of important relationships between epilepsy predisposition and psychiatric comorbidities. The inbred WAG/Rij (Wistar Albino Glaxo/Rijswijk) rat strain develops absence seizures at approximately 2–3 months of age, in parallel with the onset of depression and anxiety phenotypes [41]. Therefore, this model is ideal to investigate the age-related onset and causal relationship between depression and epilepsy with spike-wave discharges. WAG/Rij rats have deficiencies in the FST and TPT, as well as anxiety-related behaviors in the open field test [44]. The depressive symptoms in this model can be rescued by chronic treatment with the TCA, imipramine (but the effect of imipramine on seizures is unknown). Chronic treatment of WAG/Rij rats with the anti-absence drug ethosuximide from 3 weeks to 5 months of age led to persistent seizure suppression many months after discontinuation of treatment [2]. Chronic ethosuximide treatment also reduced immobility time on the FST, suggesting that this anticonvulsant exerted both antiepileptic and antidepressant effects [48]. The authors concluded that there is a causal relationship between the development of the epileptic phenotype and depressive symptoms in this model [43]. Prominent involvement of the dopaminergic system in these behavioral dysfunctions is supported by acute treatment with a dopamine receptor D2/3 antagonist, raclopride, which exacerbated FST deficiencies, and a D2/3 receptor agonist, pramipexole, which exerted antidepressant effects [42]. Recent work also implicates involvement of the mTOR pathway in both epileptogenesis and

depression in WAG/Rij rats [38]. Blockade of the mTOR pathway with rapamycin for either 7 days (“sub-chronic”) or 17 weeks (“chronic”) ameliorated absence seizures but had an opposite effect on depression using the FST and TPT – sub-chronic treatment with rapamycin had an antidepressant effect while chronic treatment produced a prodepressant effect. These results could form the basis of a novel treatment strategy for epilepsy and depression (mTOR inhibition), while raising the interesting caveat that the same agent (rapamycin) can exert different effects on depression, depending on the specific administration protocol. Taken together, data from more than three decades of study of the WAG/Rij rat absence epilepsy model strongly support a close interrelationship between seizures and psychiatric comorbidities, especially depression, and provide an excellent model in which to investigate correlations between seizure occurrence, cognitive dysfunction, and treatment parameters.

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## 22.5 Conclusion

Potential pathophysiological overlaps between epilepsy and epilepsy comorbidities are eminently amenable to study in the laboratory using animal models. While acknowledging species differences and other inherent limitations of animal models of epilepsy and psychiatric diseases, the shared pathophysiology between epilepsy and depression, anxiety, and other comorbidities are readily amenable to laboratory investigation and could yield insights into the pathophysiological mechanisms in one or both conditions, as well as potential therapeutic modalities. This rigorous approach to translational neurobiology has been typified by laboratory models championed by Dr. Philip Schwartzkroin and his colleagues.

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# What New Modeling Approaches Will Help Us Identify Promising Drug Treatments?

23

Scott C. Baraban and Wolfgang Löscher

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## Abstract

Despite the development of numerous novel antiepileptic drugs (AEDs) in recent years, several unmet clinical needs remain, including resistance to AEDs in about 30 % of patients with epilepsy, adverse effects of AEDs that can reduce quality of life, and the lack of treatments that can prevent development of epilepsy in patients at risk. Animal models of seizures and epilepsy have been instrumental in the discovery and preclinical development of novel AEDs, but obviously the previously used models have failed to identify drugs that address unmet medical needs. Thus, we urgently need fresh ideas for improving preclinical AED development. In this review, a number of promising models will be described, including the use of simple vertebrates such as zebrafish (*Danio rerio*), large animal models such as the dog and newly characterized rodent models of pharmacoresistant epilepsy. While these strategies, like any animal model approach also have their limitations, they offer hope that new more effective AEDs will be identified in the coming years.

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## Keywords

Zebrafish • Epileptic dogs • Epileptic rodents • Pharmacoresistant epilepsy • Antiepileptic drugs • Epilepsy syndromes

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## 23.1 Introduction

Rodent models of seizures and epilepsy have played a fundamental role in advancing our understanding of basic mechanisms underlying ictogenesis and epileptogenesis. They have also been instrumental in the discovery and preclinical development of novel antiepileptic drugs (AEDs) [12]. Indeed, animal models with a similarly high predictive value do not exist for other neurological disorders, such as bipolar disease or migraine [62]. Despite the availability of predictive rodent models, at least 30 % of epilepsy patients are not controlled by currently available AEDs. One reason is that, with few exceptions, most AED candidates were identified in simple evoked seizure models in otherwise healthy rodents such as the maximal electroshock seizure (MES) or acute pentylentetrazole (PTZ; metrazol) tests [48]. In these traditional models, in use since the 1940s, successful AED treatments suppress acute seizure events, but effects on drug-resistant seizure events or chronic spontaneous seizures are not routinely evaluated. Thus, we urgently need fresh ideas for improving preclinical AED development. Here, a number of promising models will be described, including the use of simple vertebrates such as zebrafish (*Danio rerio*), large animal models such as the dog and newly characterized rodent models of pharmacoresistant epilepsy. We will not discuss *in vitro* brain slice models or neurons derived from patients using induced pluripotent stem cell technology, because the network complexity of the brain and its alterations by seizure activity are difficult to recapitulate in the dish.

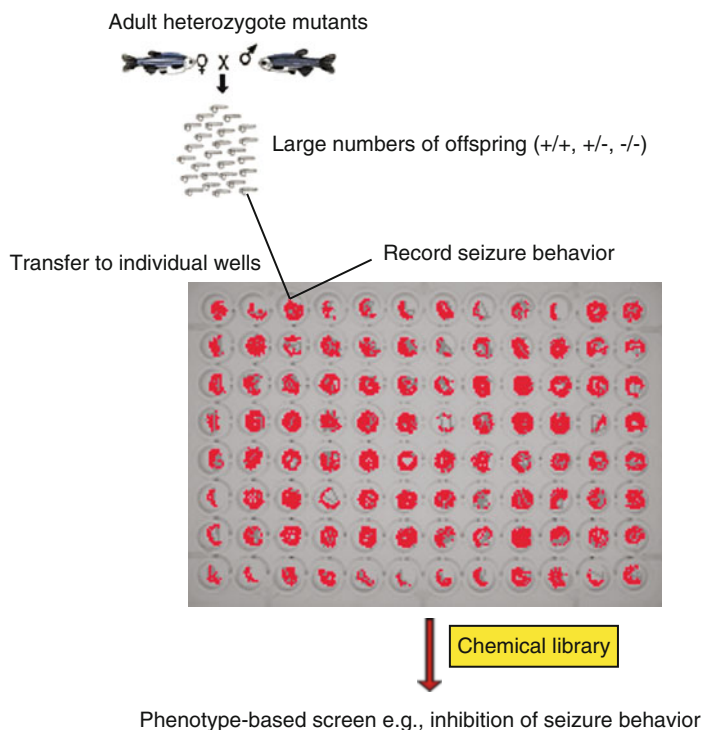
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## 23.2 Zebrafish-Based Approaches to Epilepsy and Drug Discovery

Traditionally used as a model organism to study vertebrate development and embryogenesis, zebrafish only recently emerged as an important model for epilepsy research [5, 17, 27, 29, 53, 65, 70]. The rapid *ex vivo* development, genetic trac-

tability and transparency of larval zebrafish make them ideally suited to these types of studies (Fig. 23.1). Because zebrafish are vertebrates with a fairly complex nervous system [2, 21, 61] recording electroencephalographic activity is also possible [7], and with exposure to standard convulsant manipulations (e.g., PTZ, pilocarpine, 4-aminopyridine, heat) abnormal electrical discharge with brief high-frequency small amplitude (interictal-like) and longer duration, complex multi-spike large amplitude (ictal-like) events can be readily observed. Sophisticated imaging approaches, taking advantage of the transparency of larval zebrafish and genetic modification to express calcium or bioluminescence indicators, provide additional evidence that central nervous system (CNS)-generated seizure-like activity is robust in response to PTZ. This is an important advantage of zebrafish as a model organism for epilepsy research as CNS-generated abnormal electrical events are often considered a hallmark feature of this disease. In the original description of the acute PTZ seizure model in wild-type zebrafish at 6 or 7 days post-fertilization (dpf), Baraban et al. [5] provided a framework for characterizing epilepsy in zebrafish: (i) evidence for seizure-induced gene (*c-Fos*) expression, (ii) a scoring system for seizure-like behaviours, (iii) electrophysiological examples of abnormal electrographic burst discharge and (iv) sensitivity to common AEDs (valproate, ethosuximide, carbamazepine, phenytoin, phenobarbital and diazepam). As expected from similar PTZ testing in rodents [71], valproate and diazepam were the most effective at inhibiting electrographic seizure events with approximate ED<sub>50</sub>s of 1 mM and 5 μM, respectively. Using this same model, Berghmans et al. [11] extended this dataset to include 14 standard AEDs. These follow-up experiments used an assay where wild-type larvae were “incubated” in a test compound for 24 h prior to acute PTZ administration and monitoring of seizure-like behaviour exclusively in a locomotion-based tracking assay. These studies confirmed the results of Baraban et al. [5] but also highlight the limitations of a behaviour-only assay as two drugs that failed to alter electrographic burst discharge amplitude (ethosuximide

**Fig. 23.1** Schematic illustration of the zebrafish assay



and carbamazepine) were identified as “anticonvulsant” as measured by a reduction in swim activity. A likely explanation is that overnight exposure to these AEDs was either toxic or sedative to developing zebrafish, as both possibilities would appear as suppressed locomotion in motion-based tracking assay. More recently, Afrikanova et al. [1] revisited this overnight exposure-PTZ challenge assay and evaluated a similar list of 13 AEDs using a combination of locomotion tracking followed by electrophysiology on agar-immobilized larvae. These latter studies aligned most closely with the original PTZ findings, identifying valproate and diazepam, while also showing that ethosuximide altered burst frequency but not amplitude. Maximum-tolerated drug concentrations were studied in both papers highlighting an additional advantage of the zebrafish platform for simultaneous *in vivo* evaluation of drug toxicities e.g., one of the primary reasons that most compounds identified in preclinical trials ultimately fail to reach the clinic. In a recent paper by Baxendale et al. [10] also using PTZ, a high-throughput screen of a ~2,000 bioactive

small molecule library was reported. These studies used a first-pass assay based on increased *c-Fos* mRNA expression (as measured by *in situ* hybridization) following PTZ exposure at two dpf and a secondary locomotion-based assay at four dpf for additional concentration-response studies. Unfortunately, it is unclear whether the 46 compounds identified using this approach are antiepileptic as previous studies indicate the earliest possible developmental stage where confirmed electrographic seizures could be observed in zebrafish larvae is three dpf [6, 27]. Before this age, larvae are still in chorion and do not swim freely. Furthermore, these non-physiological assays should be interpreted with caution as the Baxendale et al. [10] study identified several candidate compounds with known neurotoxicity profiles e.g., lindane, rotenonic acid, deguelin, endrin and propanil.

Although seizures can be easily induced, drug discovery using acute seizure models, even in zebrafish, are prone to the same limitations as in rodents. Namely, these approaches use healthy animals, the seizure-events are acute and evoked

using potentially non-physiological stimuli such as a stimulation electrode or convulsants, and most importantly they do not model spontaneously occurring seizure events. Zebrafish diverged from humans roughly 450 million years ago but recent genome sequencing revealed that the similarity between the zebrafish and human genome is ~70 % [28]. This fact, coupled with the fecundity of adult zebrafish (producing 100–200 offspring per week from a single adult breeding pair), the permeability of larvae to drugs placed in the bathing media, and ability to thrive in volumes as small as 100  $\mu$ l make zebrafish an attractive model for a drug discovery program targeted to genetic forms of epilepsy. In the Baraban laboratory, we have focused on zebrafish designed to mimic monogenic epilepsy disorders of childhood as they offer the advantages of spontaneous seizure activity and a genetic basis mimicking the human condition. In this approach, one can model specific forms of pediatric epilepsy – Type I Lissencephaly (*Lis1*), Angelman syndrome (*Ube3A*), Tuberous Sclerosis Complex (*Tsc*) or Dravet syndrome for example (*Scn1a*) – then design drug screening programs targeted to that patient population. In some cases these are stable mutations carried in the zebrafish germline, where other models involve acute antisense knockdown of gene expression in immature zebrafish. Thus, a form of “personalized medicine” aimed at identifying new therapeutic options for relatively rare, but catastrophic, forms of epilepsy. Our recent studies are based on a two-stage screening process. First, zebrafish mutants are placed in individual wells and behaviour (locomotion) is tracked using a 96-well format. Once a baseline level of spontaneous seizure activity is established a test compound is added, and then a second locomotion assay is performed to evaluate the effect on seizure behaviour (with distance travelled and mean velocity of swim movement used as surrogate markers) [5, 16]. As freely behaving larvae can simultaneously be observed for heart rate, edoema or touch-sensitivity, in vivo toxicity is also determined with this strategy. Using a 96-well format it is relatively easy to power this research for statistical analysis and multiple drug concentrations can

be assessed in a given plate. The same fish can subsequently be used for electrophysiological analysis, which allows a determination of “false positives” in the locomotion assay that are lethal, sedative or paralyzing. With even a modest zebrafish facility, this approach can easily be used to screen 20–50 drugs per week. The disadvantage of this strategy is that it is not well-suited to acquired forms of epilepsy that develop more slowly over time or in the adult nervous system, or compounds that are not easily dissolved in embryo media. It is also difficult to directly translate concentrations that are effective via bath application in larval zebrafish to those that may be useful clinically in humans.

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### 23.3 Rodent Models of Pharmacoresistant Seizures

The concept of developing rodent seizure or epilepsy models that do not respond to clinically approved AEDs and then using such models for the discovery of novel more effective AEDs is not new but, to our knowledge, was first proposed by Löscher in 1986 [38]. Since then, several models of pharmacoresistant seizures have been developed, including the phenytoin-resistant kindled rat [40], the lamotrigine-resistant kindled rat [68], and the phenobarbital-resistant epileptic rat [14]. In all these models, resistance to one AED extends to other AEDs (cf., [49]), thus fulfilling the criterion of pharmacoresistant epilepsy [32]. By using two of these models, Löscher and colleagues described several factors that differentiated AED-resistant from AED-responsive rats, including the extent of neurodegeneration in the hippocampus, genetic factors, AED target alterations, alterations in drug efflux transporters, and intrinsic severity of the epilepsy as a determinant of AED refractoriness [49]. Similar factors have been described for AED-resistant human epilepsy, so that the rat models obviously reflect clinically important mechanisms of refractoriness. The next logical step was to use such models for new treatment discovery. One example here is that inhibiting the drug efflux transporter P-glycoprotein (Pgp), which is increased at

**Table 23.1** A comparison of elimination half-lives of antiepileptic drugs in humans, dogs and rats

AED	Half-life (h)		
	Human	Dog	Rat
Carbamazepine	25–50 <sup>a,b</sup>	1–2 <sup>a,b</sup>	1.2–3.5 <sup>a</sup>
Clobazam	16–50	~1.5	1
Clonazepam	18–50	1–3	?
Diazepam	24–72 <sup>a</sup> (DMD=40–130)	1–5 <sup>a</sup> (DMD=4)	1.4 <sup>a</sup> (DMD=1.1)
Ethosuximide	40–60	11–25	10–16
Felbamate	14–22	4–8	2–17 <sup>c</sup>
Gabapentin	5–7	3–4	2–3
Lacosamide	13	2–2.5	3
Lamotrigine	21–50	2–5	12 to >30
Levetiracetam	6–11	4–5	2–3
Oxcarbazepine	1–2.5 <sup>a</sup> (MHD=8–14)	~4 <sup>a</sup> (MHD=3–4)	? <sup>a</sup> (MHD=0.7–4)
Perampanel	70	5	2
Phenobarbital	70–100 <sup>b</sup>	25–90 <sup>b</sup>	9–20 <sup>b</sup>
Phenytoin	15–20 <sup>b,c</sup>	2–6 <sup>b,c</sup>	~1–8 <sup>b,c</sup>
Potassium bromide	~300	~600	72–192
Pregabalin	6	6–7	2.5
Primidone	6–12 <sup>a</sup> (PB=70–100)	4–12 <sup>a,b</sup> (PB=25–90)	5 <sup>a</sup> (PB=9–20)
Tiagabin	5–8	1–2	1
Topiramate	20–30	3–4	2–5
Valproate	8–15 <sup>a</sup>	1–3 <sup>a</sup>	~1–5 <sup>a,c</sup>
Vigabatrin	5–7 <sup>d</sup>	? <sup>d</sup>	~1 <sup>d</sup>
Zonisamide	60–70	~15	8

Data are from previous reviews of Löscher [44, 46] and have been revised and updated for the present study. Note that rats and dogs eliminate most AEDs more rapidly than humans, which has to be considered when using such drugs for chronic studies in experimental animals

*DMD* desmethyl diazepam, *MHD* monohydroxy derivative, *PB* phenobarbital, ? indicates that no published data were found

<sup>a</sup>Active metabolites; <sup>b</sup>shortens on continuing exposure to the drug (because of enzyme induction); <sup>c</sup>non-linear kinetics (half-life increases with dose); <sup>d</sup>duration of action independent of half-life because of irreversible inhibition of GABA degradation

the blood–brain barrier of AED-resistant rats, counteracted resistance to phenobarbital in epileptic rats [15]. The increased Pgp functionality in epileptic rats can be visualized in vivo by positron emission tomography [4]. By using Pgp imaging, Feldmann et al. [19] demonstrated that about 40 % of AED-resistant patients exhibit increased brain functionality of Pgp and could potentially benefit from Pgp inhibition. This example illustrates that chronic rodent models of pharmacoresistant seizures are helpful to discover new strategies for treatment of medically intractable epilepsy.

The disadvantage of the described chronic epilepsy models is that they are not suited for large-scale testing of novel compounds but rather

for evaluation of selected treatment strategies as illustrated by the example of Pgp inhibition. Kindling models such as the phenytoin-resistant kindled rat [40] or the lamotrigine-resistant kindled rat [68] have the advantage that seizures can be induced at will, so that chronic drug administration is not needed, whereas models with spontaneous recurrent seizures (SRS) such as the phenobarbital-resistant epileptic rat [14] necessitate continuous (24/7) EEG/video recording for assessing drug efficacy. When testing drug effects on SRS in such rat models, the rapid elimination of most drugs, including AEDs, in rats (Table 23.1) necessitates the use of an adequate dosing regimen during prolonged drug administration to

avoid false negative results [46]. The same is true when administering potential antiepileptogenic drugs in the latent period following epileptogenic brain insults in rats [46]. Mice developing SRS after intrahippocampal injection of kainate have been proposed as a model of pharmacoresistant seizures; these mice have the advantage that the frequency of SRS is so high that drug efficacy can be determined after single dose administration [54, 66]. However, as yet this model has only rarely been used for investigating the antiepileptic efficacy of novel compounds [54].

Based on the logistical problems associated with drug testing in chronic models, models such as the zebrafish or acute rodent seizure models are indispensable when testing large numbers of investigational compounds before evaluating the most interesting compounds in chronic models. One of these acute seizure models, the 6-Hz model of partial seizures in mice, was initially proposed to provide a useful model of therapy-resistant limbic seizures [9], but more recent studies have not confirmed this idea [49]. Rather, the 6-Hz model is a valuable part of a preclinical test battery to further differentiate compounds. Also, a more recent genetic mouse model of Dravet syndrome, in which clinical symptoms of this syndrome occur after *Scn1a* heterozygous knockout, may be an interesting possibility for testing drugs or drug combinations for treatment of as yet pharmacoresistant types of seizures [59, 60]. Furthermore, a zebrafish *Scn1a* mutant, such as the one recently described by the Baraban laboratory [8] would be an efficient first pass high-throughput approach to identify potential candidate compounds that can be further investigated in chronic rodent models of pharmacoresistant seizures.

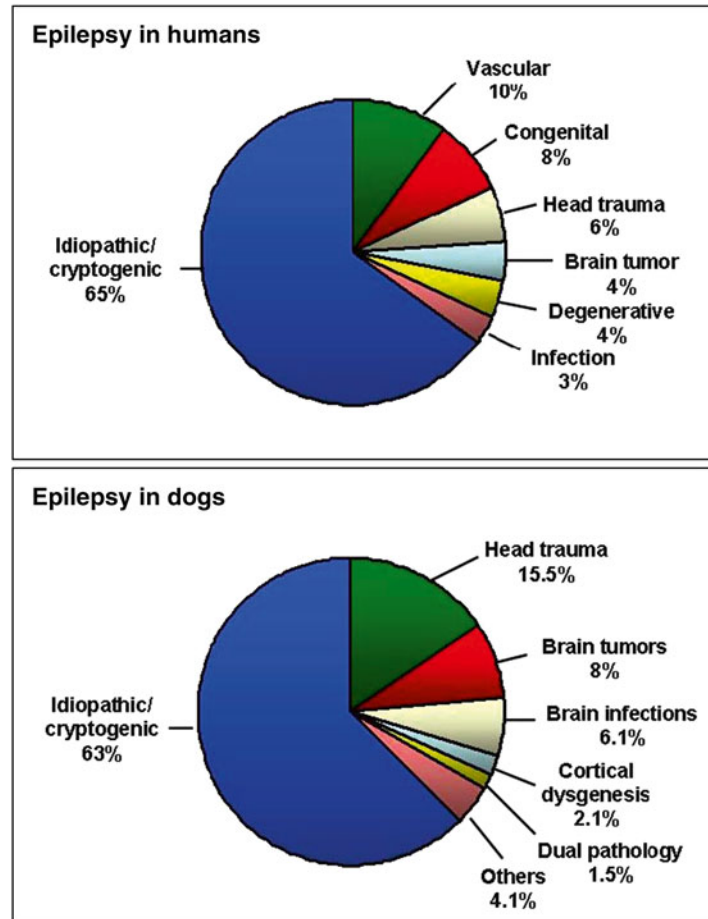
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### 23.4 Naturally Occurring Epilepsy in Dogs as a Translational Model

The dog is an important large animal model in various fields of biomedical research and fills a crucial step in the translation of basic research to new treatment regimens. For instance, because of

the relative large body size of dogs and many similarities in physiology and pharmacology between dogs and humans, scaling doses from dogs to humans is much easier than using rodents in selecting doses for clinical trials in humans. To our knowledge, Löscher et al. [37] were the first to propose naturally occurring canine epilepsy as a translational model of human epilepsy. The prevalence and phenomenology of epilepsy in dogs are very similar to human epilepsy. Indeed, epilepsy is the most common chronic neurological disease in dogs, affecting about 0.6–1 % of the dog population [64, 69]. Furthermore, causes of canine epilepsy are similar to those in humans (Fig. 23.2) except that cerebrovascular disease does not play any significant role, because it is rare in dogs [69]. About 50 % of dogs with partial and generalized convulsive seizures are not controlled by treatment with AEDs, so that epileptic dogs have been proposed as a valuable model of pharmacoresistant epilepsy that can be used to unravel mechanisms of resistance and evaluate new strategies for treatment [44, 64]. However, clinical trials on new AEDs in epileptic dogs are as laborious and time-consuming as clinical trials in human patients, necessitating randomized trial designs in which the new drug is compared with either placebo or a standard comparator [57, 58]. Recently, different treatments, including AEDs, vagal stimulation, and ketogenic diet were compared with placebo in epileptic dogs, and an unexpectedly high placebo rate was found, which was similar to that known from controlled clinical trials in humans with epilepsy [57, 58]. In contrast to humans, the placebo effect has been largely disregarded in veterinary medicine. In humans, a placebo response seems to require a recognition by the patient of the intent of treatment efforts. Because it is generally presumed that animals lack certain cognitive capacities, e.g. the ability to comprehend the intent of the veterinarian's manipulations, the power of suggestion, and expectations of recovery and healing, the existence of a placebo effect in animals seems counterintuitive [55]. However, in veterinary studies, the placebo response may be a result of expectations of the pet owner regarding treatment in studies as those conducted by Munana et al.

**Fig. 23.2** A comparison of the presumed causes of recurrent epileptic seizures in humans and dogs. The graph on humans illustrate the proportion of incidence cases of epilepsy by etiology in Rochester, Minnesota, U.S.A., 1935–1984 [24]; a similar graph was initially shown by Lowenstein [35]. The graph on dogs illustrates data from a recent epidemiologic study on canine epilepsy [69]



[57, 58] in epileptic dogs, where the owners are responsible for administration of treatment and outcome measures (i.e., seizure frequency) are derived solely from owner observations. Other factors that may be included in placebo responses in veterinary studies include regression to the mean, investigator bias, client bias, the potential for a higher level of care during the study, and improved adherence to treatment with active medication that is being administered in addition to the placebo during the study (for details see [57]). Furthermore, the placebo response can be because of effects of placebo administration on the animal, which is well documented in laboratory animals and may involve conditioned responses among others [55]. As a consequence, studies on new treatments in laboratory animals (or pets) should always include a “placebo” group receiv-

ing all manipulations (e.g., handling, injections, electrode implantation, seizure recording etc.) that are used for the new treatment.

In addition to chronic epilepsy, naturally occurring canine status epilepticus (SE) has been proposed as a translational platform for evaluating investigational compounds for eventual use in human trials [34] and a controlled study on i.v. levetiracetam for treatment of SE in dogs has been published recently [23].

One important caveat that has to be considered when using dogs for long-term studies on AEDs is that dogs, similar to rodents, eliminate many drugs, including most AEDs, much more rapidly than humans (Table 23.1). Thus, when using AEDs such as phenytoin, carbamazepine or valproate with too low half-lives for maintenance treatment in epileptic dogs, no sufficient drug



levels and, hence, no antiepileptic effects are obtained in this species [20, 36, 37]. The few AEDs with sufficiently long half-lives for maintenance treatment include phenobarbital, primidone (because of its metabolism to phenobarbital), and potassium bromide, which is the reason why until recently only these old drugs were approved for treatment of canine epilepsy in the US or Europe. This situation has changed by the recent approval of imepitoin for treatment of dogs with newly diagnosed epilepsy (see below). Furthermore, several newer AEDs, including levetiracetam, felbamate, zonisamide, topiramate, gabapentin, and pregabalin are used as add-on treatment in dogs with pharmacoresistant seizures [64]. It has been tried to overcome the problem of too rapid elimination of most AEDs by dogs by using sustained-release formulations; however, sustained-release preparations developed for use in humans are not suited for dogs because of the much higher gastrointestinal passage rate in dogs (~24 h) vs. humans (~65–100 h) [36, 44]. Thus, AED formulations that exhibit retarded release of the drug in the gastrointestinal tract have to be adapted to the dog to overcome problems associated with too rapid drug elimination in this species. For phenytoin, a slow-release preparation has been developed for dogs, by which therapeutic plasma levels could be maintained despite the rapid elimination of this drug in dogs [18], but, to our knowledge, no clinical experience with this preparation has been published. Vigabatrin has been evaluated for control of epilepsy in dogs, because its mechanism of action (irreversible inhibition of GABA degradation) allows an effective treatment which should be independent of species differences in drug elimination. Vigabatrin proved to be effective in epileptic dogs with phenobarbital-resistant seizures, but at least in part vigabatrin had to be withdrawn because of development of severe adverse effects, such as haemolytic anaemia [67].

Löscher's group has used dogs as a translational model over the recent 25 years in the development of a new category of AEDs, i.e., drugs that act as partial agonists at the benzodiazepine (BZD) site of the GABA<sub>A</sub> receptor. Such drugs have the wide spectrum of antiepi-

leptic activity against diverse types of seizures as the traditional full BZD agonists such as diazepam, clonazepam or clobazam, but are much better tolerated and lack the tolerance and abuse liability of the full agonists [22, 41]. In our studies, we either used a canine seizure model, in which seizures are induced by i.v. infusion of pentylenetetrazole, or epileptic dogs. The first partial BZD agonist that was characterized in dogs (and compared with full BZD agonists) was the  $\beta$ -carboline abecarnil, providing proof-of-concept that partial BZD agonists are advantageous for treatment of seizures compared to traditional, full-agonist BZDs [39, 41]. More recently, the low-affinity partial BZD agonist imepitoin, an imidazolin derivative, was evaluated in the dog seizure model and epileptic dogs and reported to provide efficacious antiepileptic activity without the known disadvantages of full BZD agonists [45, 51]. Based on several randomized controlled clinical trials in epileptic dogs, imepitoin was recently approved in Europe for treatment of canine epilepsy [13, 51]. That imepitoin is an effective and safe AED in epileptic dogs indicates that low-affinity partial BZD agonists may offer a new mechanistic category of useful AEDs.

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### 23.5 Network Approaches for Development of Novel Treatments

Several of the models described in this review may be particularly interesting for evaluating a novel strategy of AED development, the network approach [3, 26, 50]. One of the dominant strategies in drug discovery is designing maximally selective ligands to act on individual drug targets [26]. However, many effective drugs act via modulation of multiple targets rather than single proteins. Furthermore, most epilepsies develop not from alterations of a single target but rather from complex alterations resulting in an epileptic network in the brain. The only existing cure of epilepsy is resective surgery in which the regional epileptic network or part of this network is removed. Thus, treatments focusing exclusively

on a single protein or individual biochemical pathway may be less effective than treatments targeting different proteins or pathways involved in the network. The latter approach has been recently termed “network pharmacology” and relates to principles of systems biology [3, 26]. The principle of network pharmacology is to develop combinations of existing drugs, which regulate activity via different targets within a biological network, for diseases that do not sufficiently respond to single drug treatment or for which no treatment exists. Integrating network biology and polypharmacology holds the promise of expanding the current opportunity space for druggable targets [26]. However, the rational design of polypharmacology faces considerable challenges in the need for new methods to validate target combinations and optimize multiple structure-activity relationships while maintaining drug-like properties. The advances in zebrafish chemical screening technologies may allow rapid identification of the most interesting drug combinations resulting from network approaches, followed by evaluating these combinations in chronic models of epilepsy.

Some examples for interesting network approaches include combinations of glutamate receptor antagonists that target different glutamate receptor subtypes. We reported that extremely low doses of the NMDA (N-methyl-D-aspartate) receptor antagonist MK-801 (dizocilpine) markedly potentiated the anticonvulsant effect the AMPA (alpha-amino-3-hydroxy-5-methyl-4-isoxazolepropionic acid) receptor antagonist NBQX (2,3-dihydroxy-6-nitro-7-sulfamoylbenzo(F)quinoxaline) without increasing its adverse effects [42]. Similar over-additive effects were seen when NBQX was combined with the competitive NMDA antagonist CGP39551 (the carboxyethyl ester of DL-(E)-2-amino-4-methyl-5-phosphono-3-pentenoic acid) or the low-affinity, rapidly channel blocking NMDA receptor antagonist memantine [42, 43]. We are currently evaluating combinations of clinically approved NMDA antagonists (ketamine, memantine) and the novel AMPA antagonist perampanel in models of difficult-to-treat seizures. Another interesting example is the combination

of phenobarbital with the diuretic bumetanide, which is currently evaluated clinically following promising preclinical data [31, 52]. The biologically plausible idea behind this combination is that a shift from inhibitory to excitatory GABA may be involved in difficult-to-treat neonatal and adult seizures [30, 56]. GABA-mediated excitation has been observed when expression of the chloride importer NKCC1 is higher than expression of the chloride exporter KCC2; e.g., early during development and in the hippocampus of adults with temporal lobe epilepsy [30, 56]. Bumetanide inhibits the neuronal chloride cotransporter NKCC1, thereby reverts the GABA shift and enables GABAergic drugs such as phenobarbital to potentiate inhibitory GABAergic transmission [52]. This recent work builds on an earlier demonstration from the Schwartzkroin laboratory that furosemide, another chloride cotransporter inhibitor, exhibits powerful anti-convulsant activity across a range of *in vitro* and *in vivo* seizure models [25]. Further examples for interesting network approaches include combined targeting of different inflammatory pathways, which are involved in seizure generation [33]. These examples strongly indicate that combinatorial treatment strategies offer new options for epilepsy therapy.

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## 23.6 Conclusions

Models for the discovery of drugs with antiepileptic activity have traditionally relied on a relatively small number of acute seizure models employed in otherwise healthy rodents. While useful in the discovery of most drugs currently available in the clinic, more resistant types of epilepsies including temporal lobe epilepsy patients who are unresponsive to available AEDs and catastrophic, often genetically-based, types of epilepsies seen in children necessitate alternative drug discovery strategies. Zebrafish, canine and novel rodent approaches are described here and offer several unique advantages over these traditional models. While these strategies, like any animal model approach also have their limitations, they offer hope that new classes of

AEDs will be identified in the coming years. Furthermore, animal models in which epilepsy develops after brain insults or gene mutations are essential in the search for novel antiepileptogenic treatments that prevent or modify the development of epilepsy in patients at risk [47, 63]. Previously, this field was dominated by studies in SE models in rats, although SE is only rarely a cause of symptomatic epilepsy [47]. Thus, models of more common causes of acquired epilepsy, such as traumatic brain injury, and models in which epilepsy develops after gene mutations should be used more extensively in research on antiepileptogenesis. We have started to use the zebrafish and canine approaches to identify molecular pathways that may be involved in the epileptogenic process and may offer new targets for antiepileptogenic treatments.

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# What Are the Arguments For and Against Rational Therapy for Epilepsy?

# 24

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and H. Steve White

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## Abstract

Although more than a dozen new anti-seizure drugs (ASDs) have entered the market since 1993, a substantial proportion of patients (~30 %) remain refractory to current treatments. Thus, a concerted effort to identify and develop new therapies that will help these patients continues. Until this effort succeeds, it is reasonable to re-assess the use of currently available therapies and to consider how these therapies might be utilized in a more efficacious manner. This applies to the selection of monotherapies in newly-diagnosed epilepsy, but perhaps, more importantly, to the choice of combination treatments in otherwise drug-refractory epilepsy. Rational polytherapy is a concept that is predicated on the combination of drugs with complementary mechanisms of action (MoAs) that work synergistically to maximize efficacy and minimize the potential for adverse events. Furthermore, rational polytherapy requires a detailed understanding of the MoA subclasses amongst available ASDs and an appreciation of the empirical evidence that supports the use of specific combinations. The majority of ASDs can be loosely categorized into those that target neurotransmission and network hyperexcitability, modulate intrinsic neuronal properties through ion channels, or possess broad-spectrum efficacy as a result of multiple mechanisms. Within each of these categories, there are discrete pharmacological profiles that differentiate individual ASDs. This chapter will consider how knowledge of MoA can help guide therapy in a

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rational manner, both in the selection of monotherapies for specific seizure types and syndromes, but also in the choice of drug combinations for patients whose epilepsy is not optimally controlled with a single ASD.

### Keywords

Mechanism of action • Anti-seizure drugs • Monotherapy • Polytherapy • Drug-refractory epilepsy

## 24.1 Introduction

Approximately 50 million people worldwide suffer from epilepsy. While more than 20 anti-seizure drugs (ASDs<sup>1</sup>) are currently available and many patients can be successfully treated with just one drug, there remains a substantial population (up to 40 % of all newly diagnosed patients) whose seizures are unresponsive to monotherapy [48, 57]. Most, if not all, of these patients will receive combination therapy at some point in the clinical management of their epilepsy [86]. Many will be exposed to newer ASDs, which are invariably brought to market as “adjunctive treatments” to those ASDs that are currently approved.

Emerging evidence suggests that combining ASDs with different mechanisms of action (MoAs) may be the most effective means to successfully manage difficult-to-control epilepsy [11]. Since many patients with refractory epilepsy may take three or more ASDs concurrently, it is essential that clinicians select drugs with the greatest potential for synergism and the lowest risk for adverse effects [11]. This can be considered “rational” polytherapy. Ultimately, this approach may offer the greatest potential to effectively manage seizures in patients with pharmacoresistant epilepsy: maximizing benefit and minimizing harm. Such an approach may also highlight novel

pathways or targets that might be exploited in future drug development efforts.

## 24.2 Does Mechanism of Action Really Matter?

It is logical to suggest that MoA should be considered at a number of steps in the treatment spectrum: when choosing an initial monotherapy for some primary generalized epilepsies; when considering a switch to a new ASD after a previous monotherapy has failed; or when adding a second or even third ASD in the therapy-resistant patient. Unfortunately, the absence of important clinical data from appropriate double-blind randomized clinical trials, which attempt to compare mechanistically distinct ASDs in discrete patient populations, prohibits such a logical therapeutic approach. Designing and delivering such a trial would be an enormous undertaking and one that is unlikely ever to be fully realized, on both logistical and financial grounds. Moreover, much of the evidence that is available to the patient with epilepsy and his or her clinician has been derived from clinical observation and often as a result of the desire to avoid poor outcomes, rather than to optimize the likelihood of good ones. This is most evident in the case of seizure worsening, where knowledge of MoA, the syndromic diagnosis and, in some cases, the underlying etiology can be beneficial. For example, clinical experience has demonstrated that GABAergic agonists and sodium channel blockers can worsen generalized spike-wave seizures in absence epilepsy. Similarly, sodium channel blockers, but

<sup>1</sup> Anti-seizure drugs (ASDs) is a new descriptive term considered by some to better reflect the effects of current therapies for epilepsy, in that they prevent only one of many sequelae of the disorder, i.e. the seizures, but not other comorbidities associated with epilepsy [14, 45].

not GABAergic agonists, can worsen seizures in patients with Dravet's syndrome or severe myoclonic epilepsy of infancy (SMEI). Time will tell whether ongoing improvements in our understanding of the underlying molecular etiology of the epilepsies will direct the choice of treatment in other seizure types.

Robust empirical evidence to support mechanism-driven therapy may be lacking, but proof-of-principle can be derived through post-hoc analysis of clinical trial data. This is unfortunately not done with sufficient regularity. Results from head-to-head monotherapy studies (where available) can be scrutinized for any evidence of preferential efficacy of a specific MoA within a specific seizure type [10, 37]. Likewise, add-on clinical trial studies of new ASDs can be interrogated for evidence of preferred combinations of ASDs, as was done with post-hoc analysis of lacosamide trial data [76]. In this analysis, lacosamide appeared to possess less efficacy and to be associated with more adverse effects when added to existing treatment regimens that contained at least one "traditional" voltage-gated sodium channel (VGSC) blocking ASD (*i.e.* phenytoin, carbamazepine, lamotrigine, oxcarbazepine) than when added to regimens that were devoid of sodium channel blockers [76]. Although the power of such post-hoc analyses is questionable and the original studies on which they are based are both heterogeneous and not necessarily reflective of real life, the results are important to direct future rational therapy decisions.

Such insight allows for some generalizations, not least of which is that, for newly-diagnosed focal epilepsies, MoA is mostly irrelevant. The majority of these patients will respond to a modest dose of whichever drug is chosen [9], with choice more often dictated not by MoA, but by clinical and demographic characteristics. In this population, MoA becomes more relevant when patients start to fail ASDs due to a lack of adequate seizure control at a therapeutic dose. Under those circumstances, for example, it would not make sense to replace one VGSC blocker with another. Failure due to adverse effects is different and it would be reasonable to replace carbamazepine with lamotrigine if carbamazepine

was effective, but not well tolerated. Arguably, MoA becomes most important in this population when the decision is made that monotherapy is not sufficient and that polytherapy is required. Under those circumstances, the best outcomes are often seen with drugs that work in different ways. For the drug refractory patient, the question then becomes: what is meant by "different"? Are lamotrigine and lacosamide different? Are benzodiazepines and barbiturates different? Is it enough to consider the class into which the drug might be arbitrarily placed, or is discrete consideration of the pharmacological minutiae more important? That remains unclear. A related issue is the supposed promiscuity of the majority of ASDs in terms of their cellular effects, resulting in negative perceptions of the efficacy of the drug in that particular circumstance. This unfortunate attitude often undermines efforts to explore and to implement rational treatment strategies for therapy-resistant epilepsy on the basis of MoA.

The understanding of how ASDs exert their effects at the cellular level has improved immeasurably in the past 25 years [52]. This advance will only further optimize treatment outcomes in epilepsy. Admittedly for some ASDs, the precise MoA remains frustratingly elusive, but for most, the primary cellular effects are now well described [93]. In the remainder of this chapter, we describe current understanding of ASD MoAs, categorized by target type (Table 24.1), and thereafter discuss the clinical implications of those actions and how therapeutic management may develop in future years from such observations.

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### 24.3 Compounds That Target Neurotransmission and Network Synchronization

Epilepsy, in its broadest sense, is generally considered to arise due to an imbalance in, or abnormal synchronization of, inhibitory and excitatory signaling within neuronal networks [2, 80]. As such, it is not surprising that most currently available ASDs target ion channels or receptors involved in excitatory and/or inhibitory neurotransmission (Table 24.1) [49, 93]. Similarly, it



**Table 24.1** Mechanism of action of approved anti-seizure drugs

Mechanism of action	Anti-seizure drug(s)
<b>Neurotransmission and network synchronization</b>	
Inhibitory neurotransmission: GABA system modulation	Barbiturates, Benzodiazepines, Felbamate, Stiripentol, Tiagabine, Topiramate, Valproate, Vigabatrin
Excitatory neurotransmission: Glutamate receptor modulation	Felbamate, Perampanel, Topiramate
Synaptic vesicle modulation: SV2A protein binding	Levetiracetam
<b>Neuronal voltage-dependent ion channels</b>	
Sodium (Na <sup>+</sup> ) channels	
Fast-inactivated	Carbamazepine, Eslicarbazepine, Lamotrigine, Oxcarbazepine, Phenytoin, Rufinamide, Topiramate, Zonisamide
Slow-inactivated	Lacosamide
Calcium (Ca <sup>2+</sup> ) channels <sup>a</sup>	Ethosuximide, Gabapentin, Lamotrigine, Pregabalin, Topiramate, Valproate, Zonisamide
Potassium (K <sup>+</sup> ) channels: Kv7.2/7.3 selective	Ezogabine

<sup>a</sup>As noted in the text, ASD effects on voltage-gated calcium channels have to be differentiated on the basis of whether they modify low or high voltage-gated calcium channels; e.g., ethosuximide, valproate and zonisamide have all been reported to modify the low voltage-gated T-type calcium current

makes sense that ASDs which display broad mechanistic profiles, *i.e.*, those that target multiple processes and pathways that are known to contribute to abnormal network synchronization, such as valproate, felbamate, topiramate, or zonisamide, often display broad spectrum clinical utility [51, 62]. Thus, understanding the specific MoAs of various ASDs may improve treatment outcomes when drug combinations are selected that display the most promising synergistic interactions at both inhibitory and excitatory synapses while conferring the least risk for adverse events.

Curbing excitatory neuronal activity can be achieved through GABAergic neuromodulation [5]. Some of the earliest marketed ASDs, including barbiturates and benzodiazepines, directly target

the GABAergic system (Table 24.1) and although the MoA of valproate remains to be definitively identified, one of its many pharmacological effects is to increase synaptic GABA turnover [49, 51]. Of the newer ASDs, two were specifically designed to enhance synaptic GABAergic inhibitory neurotransmission. Tiagabine blocks synaptic GABA reuptake [66, 87] thereby prolonging the inhibitory action of GABA at GABAergic synapses, whereas vigabatrin selectively inhibits GABA transaminase [49, 77], an action that prevents the catabolism of GABA and increases readily releasable GABA within presynaptic terminals [49]. Topiramate enhances GABAergic neurotransmission through non-benzodiazepine site effects on the GABA<sub>A</sub> receptor [82, 83]. More recently, stiripentol has been approved for the treatment of Dravet's syndrome. Stiripentol is a positive allosteric modulator of  $\alpha 3$ - $\beta 3$ - $\gamma 2$ -containing GABA<sub>A</sub> receptors, increasing GABAergic neurotransmission in neuronal circuits where this receptor subtype is expressed. Preference for stiripentol over non-selective GABA<sub>A</sub> receptor drugs in Dravet's syndrome suggests that pursuing subunit selective agents in drug development may provide improved seizure control or tolerability in other epilepsies [18]. With multiple ASDs that target the GABAergic system (Table 24.1), pharmacological enhancement of inhibitory neurotransmission can be considered a well-proven strategy for seizure control.

Until recently, efforts to target glutamate-mediated excitatory neurotransmission have met with disappointment. Within the brain, excitatory synaptic transmission is mediated predominately by AMPA- and NMDA-type glutamate receptors [20, 28]. Early preclinical evidence suggested that modulating glutamatergic signaling could effectively control or suppress seizures [78]. However, efforts to develop NMDA-receptor selective antagonists for the clinical management of epilepsy met with difficulty due to significant adverse behavioral effects [90]. To date, only felbamate possesses any substantial effects on NMDA-type glutamate receptors [41, 46, 73, 92]. Conversely, the modulation of AMPA-type glutamate receptors holds more clinical promise [74]. Modulation of AMPA, but not NMDA [42], receptor signaling exerts fewer effects on synaptic plasticity [38]

and has greater potential to modulate network hyperexcitability [74], an effect largely attributable to activity-dependent AMPA receptor localization dynamics that underlie fast synaptic excitatory neurotransmission [1]. To this point, perampanel is the first glutamatergic system-selective ASD that acts as a noncompetitive AMPA receptor antagonist [72], decreasing neuronal excitability and synchronization [74]. Amongst many other proposed MoAs, topiramate has also been shown to suppress excitatory neurotransmission by blocking non-NMDA type-glutamate receptors [36] and can reduce high basal concentrations of extracellular glutamate in the hippocampi of spontaneously epileptic rats [44]. Taken together, the effect of felbamate on NMDA receptors, the AMPA-selective effects of perampanel, and the non-NMDA effects of topiramate further demonstrate that glutamatergic modulation can efficiently suppress epileptic activity.

Excitatory neurotransmission may also be influenced by the binding of the ASD levetiracetam to the SV2A protein (Table 24.1), a membrane glycoprotein of synaptic vesicles [3]. The specific role of SV2A protein is still under active investigation. It is currently hypothesized that SV2A contributes to excitatory neurotransmission by participating in synaptic vesicle exo- and endocytotic processes in response to calcium-triggered vesicle fusion [24, 97]. Interestingly, SV2A knockout mice develop severe seizures [22] and resected brain tissues from patients with temporal lobe epilepsy show decreased immunoreactivity for SV2A protein [89], suggesting a possible role of reduced levels of SV2A in epileptogenicity. Levetiracetam is effective in the 6 Hz model of psychomotor seizures, but ineffective in other “traditional” animal models of epilepsy, further highlighting its unique pharmacological profile [47]. The availability of a drug like levetiracetam might be considered advantageous from a rational therapy perspective. The unique and novel MoA may be effectively combined with other ASDs with diverse MoAs to mitigate seizure frequency and susceptibility. More importantly, where it is possible to use selective combinations of such diverse MoAs to enhance efficacy, it may be possible to minimize the likelihood of adverse events by decreasing the total exposure burden of

the ASDs. Indeed, the fact that levetiracetam possesses a unique mode of action could explain why it has been so successful clinically, as both monotherapy and adjunctive treatment.

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#### 24.4 Compounds That Modulate Intrinsic Neuronal Properties

The above-described drug mechanisms modulate seizure susceptibility by selectively regulating excitatory and inhibitory neurotransmission and thereby suppressing aberrant neuronal network synchronization. However, many of the currently available treatments for epilepsy can also modulate the intrinsic excitability of individual neurons by targeting ion channels (Table 24.1). Multiple ASDs, such as carbamazepine, phenytoin, lamotrigine and oxcarbazepine modulate the fast-inactivated state of VGSC, whereas lacosamide is thought to have a preferential effect on the slow-inactivated state of the channel. In contrast, the gabapentinoids (gabapentin and pregabalin) and ezogabine decrease neuronal firing by selectively targeting calcium and potassium channels, respectively. It is likely that ion channel-selective mechanisms have naturally emerged amongst ASDs because ion channel dysfunction is so heavily implicated in the pathophysiology of many idiopathic epilepsies [55]. Recent evidence also demonstrates a link between genetic channelopathies and acquired epilepsies and supports the further development of ion channel modulators for the management of seizure disorders in general [68].

Ion channel-targeting drugs modulate depolarization and action potential generation and propagation. Many ASDs, including carbamazepine, phenytoin, lamotrigine and oxcarbazepine, bind VGSCs, with preferential affinity for the channel protein when in the fast-inactivated state. This leads to a prolongation of recovery following transient depolarizations [54], thereby limiting repetitive action potential firing. This effect is also both use- and frequency-dependent, meaning that it is enhanced during periods of high-frequency neuronal firing, as during epileptic discharges. Topiramate may also exert some anti-seizure effects through blockade of use-dependent VGSCs, although this effect appears to be different

from traditional VGSC-blocking ASDs [56]. Conversely, lacosamide also targets VGSCs but with preferential effects on the slow-inactivated state of the channel which predominates during sustained depolarization [23]. A selective action on the slow-inactivated rather than fast-inactivated state of VGSCs promotes the stabilization of hyperexcitable neuronal membranes, and suggests that lacosamide is pharmacologically distinct from traditional ASDs that target the fast-inactivated state (Table 24.1) [23]. The characteristic use- and frequency-dependence of sodium channel block is the only example of selectivity for a disease-related mechanism amongst current ASDs and explains why these drugs can interfere with a fundamental neurophysiological mechanism without significantly affecting normal neuronal activity.

In addition to blocking sodium channels, several ASDs act via blockade of voltage-gated calcium channels (Table 24.1), an action that effectively decreases intracellular calcium ion concentration. In the dendrites and cell soma, elevated intracellular calcium can promote destabilization of VGSCs and increase cellular excitability and the likelihood of action potential firing [69]. In pre-synaptic nerve terminals, elevated intracellular calcium is the trigger for neurotransmitter release. ASDs that selectively target high voltage-activated calcium channels have found success in the management of epilepsy and also neuropathic pain [85]. The gabapentinoids (gabapentin and pregabalin) selectively bind to the accessory subunit  $\alpha_2\delta$ -1 of voltage-gated calcium channels [32] to block P/Q-type calcium currents at nerve terminals, reducing the calcium-dependent release of glutamate [31]. Lamotrigine has also been shown to target P/Q-type, N-type and R-type channels [30, 91], all of which are expressed on pre-synaptic nerve terminals. Topiramate and felbamate are reported to have similar effects, although the channel subtypes are less well defined. However, a different action is seen with ethosuximide. This ASD interacts with the low voltage-activated T-type calcium channel that is predominantly expressed on thalamocortical relay neurons [21], which have in turn been implicated in the generation of the hypersyn-

chronous discharges that underlie generalized absence epilepsy. Blockade of T-type channels by ethosuximide almost certainly explains its efficacy in this regard, and may also explain the anti-absence effects of both valproate and zonisamide [13, 88]. Thus, several currently available ASDs modulate high and low voltage-activated calcium channels; an effect that indirectly reduces excitatory neurotransmission at glutamatergic synapses and limits neuronal synchronization.

Rather than targeting cellular excitability by limiting depolarization, it is also possible to promote hyperpolarization via a facilitatory effect on potassium currents. This is the primary MoA of ezogabine, which is a positive allosteric modulator of  $K_v7.2/7.3$  voltage-gated potassium channels that carry the so-called M-current [96]. The M-current is a non-inactivating potassium conductance, which exerts a hyperpolarizing influence on the resting membrane potential [53]. It serves as a natural brake on excitability in regions prone to synchronous network activity, with enhancement of the M-current by ezogabine suppressing epileptiform activity by prevention of spike bursting [75, 98, 99]. The role of potassium channels in modulating neuronal excitability is further underscored by the finding that mutations in  $K_v7.2/7.3$  potassium channel genes provide the basis for seizures in benign familial neonatal convulsions (BFNC), a rare form of epilepsy that arises due to mutations in these channels [7, 8, 16]. Thus, ezogabine provides another example of how targeting intrinsic neuronal properties can attenuate epileptic activity.

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## 24.5 Compounds That Reduce Seizure Susceptibility Through Multiple Mechanisms

As outlined above, currently available ASDs exert their effects through multiple mechanisms, including suppression of neuronal excitability, reduction in the propensity to fire an action potential, disruption of neurotransmission through interactions with synaptic vesicles, or an increased

inhibitory tone. Some ASDs, namely valproate, felbamate, topiramate and zonisamide, are reported to possess two or more of these effects (Table 24.1). These ASDs are effective across multiple epilepsy indications [51], suggesting that one way to treat a heterogeneous disorder like epilepsy is to broadly target ion channels and neurotransmitter receptors. Indeed, several clinical and preclinical studies provide strong proof-of-concept for such a treatment strategy [29, 76], with some authors suggesting that combining a drug with a single MoA with a drug with multiple MoAs may improve seizure control [11]. However, this approach might also result in mechanistic redundancy (where a specific mechanism is unnecessary or unhelpful) or reinforcement (where a specific mechanism is duplicated) – either scenario could potentially elevate the risk of adverse effects without necessarily enhancing seizure control. This can be considered an inherent limitation of drugs with multiple MoAs, that in some patients not all of those mechanisms will be beneficial and some may indeed be detrimental. This would explain why these compounds are often considered to be powerful drugs with proven broad-spectrum efficacy but which are occasionally not well-tolerated.

Broad-spectrum ASDs pose an interesting pharmacological conundrum: whether a single drug with multiple MoAs (*i.e.* polypharmacology) is equivalent, superior, or inferior to multiple drugs, each with single MoAs (*i.e.* polypharmacy)? Would it be better to use a combination of phenytoin, ethosuximide and acetazolamide, which target sodium channels, T-type calcium channels, and carbonic anhydrase respectively, or zonisamide, which targets all of these mechanisms simultaneously? Rational polytherapy would suggest that the single drug should behave in exactly the same way, in terms of both efficacy and tolerability, as the three-drug combination, assuming dose equivalents can be found and drug interactions can be compensated for. This is a puzzle that will probably never be solved because there are likely to be few prescribers who would choose the combination therapy under these circumstances. Most would opt instead for zonisamide on the grounds of ease of use, but

also to limit adverse effects that are perceived to hinder polypharmacy approaches.

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## 24.6 Preclinical Evidence of Potential for Polytherapy

Preclinical data suggest that the greatest potential for synergistic effects with polytherapy arises when ASDs with multiple MoAs are combined with ASDs with single, distinct MoAs [34, 43]. Preclinical isobolographic studies in rodent models of epilepsy have played an important role in teasing out either favorable or synergistic interactions from those that may be negative or antagonistic [25, 64, 79]. This preclinical evidence supports the concept of rational polypharmacy to control seizures using ASDs with diverse MoAs, although critics would argue that data from studies involving experimental animals are far more frequently positive than is observed clinically. With the possible exception of valproate combined with lamotrigine, synergistic combinations identified in animal models do not appear to reliably extend to the clinic.

Such discrepancies between preclinical and clinical observations can be explained, at least in part, by the inherent limitations in the preclinical studies. First, they are almost always conducted using high-throughput models of acute seizures in non-epileptic rodents. Given that the epileptic brain is undoubtedly remodeled relative to the normal brain [50], there are likely to be changes in the pharmacological responsiveness. Second, these studies are invariably conducted following acute drug dosing and with efficacy determined at or around the time of peak effect. This clearly does not reflect the clinical situation, and ignores any pharmacokinetic interaction that may exist between the compounds being tested or any tolerance that might develop from repeated administration. Such effects would also bypass the role of hepatic induction and/or inhibition seen when two or more ASDs are chronically combined in the patient population. This does not imply that these types of preclinical studies of ASD combinations lack value, but simply that the results should not be automatically assumed to translate

to the clinic. The apparent discrepancy between the results of preclinical and clinical combination studies could just as easily be explained by the fact that clinical studies have never been systematically explored. It is possible that the same combinations are synergistic in both animal models and human patients; it is just that we do not yet possess the clinical evidence to prove it.

If nothing else, preclinical evaluation can provide important insights into potentially synergistic and antagonistic combinations, which can be taken forward for more detailed clinical investigation. In this regard, such an approach can help to triage the myriad of possible combinations and allow clinical researchers to focus on those likely to be most beneficial (or least detrimental). With the introduction of more etiologically relevant animal models, future studies can be designed to examine combinations using chronic dosing in animals with therapy-resistant epilepsy. Such studies should more clearly define the true clinical potential for synergism. However, as the models become more elaborate and the treatment schedules more demanding, the likelihood of undertaking in-depth isobolographic studies of every possible ASD combination diminishes, not least because of the time and cost involved. A compromise may be required such that combinations are initially identified using acute seizure models, later confirmed using chronic treatment in models of drug-resistant epilepsy, and only then advanced to clinical validation studies.

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## 24.7 Clinical Evaluation of Polytherapy

Validating the findings of preclinical studies and thus establishing a basis for rational polytherapy requires the formal clinical evaluation of combinations of ASDs. This is a complex and challenging task that is unlikely to ever be systematically completed. It is more likely that clinical validation will be reserved for only the most robust combinations and in circumstances where evidence is considered absolutely essential to clinical implementation.

Clinical combination studies are notoriously difficult to undertake. They require the investigation of efficacy and tolerability of both single drugs and combinations in relatively homogeneous populations of patients using a design that is sufficiently powered to separate synergism from additivity alone, as well as adjusting combinations to balance overall drug load. Not surprisingly, such a study has never been attempted in epilepsy and it is debatable whether one ever will be. In the meantime, we are largely dependent on small proof-of-principle studies and anecdotal observations for evidence of effective polytherapy regimens. These include the classic and unexpected observations suggesting synergism with valproate and lamotrigine [12], which were later proven to hold true [67]. These studies provided validation of an ASD combination that was already in widespread use and probably considered useful by many investigators but for which there was no specific evidence of benefit. However, it is debatable whether there will ever be sufficient imperative or resources to pursue such a validation in the future.

The more applicable alternative strategy is the utilization of existing resources to search for at least indirect evidence of synergism. In this regard, post-hoc analysis of Phase III regulatory trial data provides a potential opportunity. The study by Sake and colleagues used the reanalysis of Phase III clinical trial data to demonstrate efficacy and tolerability with add-on lacosamide stratified by background therapy (*i.e.* whether it contained sodium channel blockers or not) [76]. Although the trials were never designed for this purpose and the analysis was arguably under-powered, some interesting findings were reported. Not least of these was the observation that lacosamide, which targets the slow-inactivation state of the VGSC, showed reduced efficacy and enhanced adverse effects when combined with traditional sodium blockers (which target the fast-inactivated state) than when combined with non-sodium channel blocking drugs [76]. If a similar approach were undertaken with all newly licensed ASDs, we could rapidly develop a picture of which mechanisms work best together and over time, with sufficient numbers of studies, it may

be possible to start to investigate individual drug combinations. Making this a mandatory requirement in the regulatory approval process would expedite the generation of such data and insisting on its release for independent meta-analysis would add further validity.

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## 24.8 Other Considerations for Rational Therapy

Consideration of multiple factors including MoA, route of metabolism and excretion must be made when determining the therapeutic treatment strategy. While it is generally considered best to combine therapies with different MoAs to maximize clinical effect, the risk for drug-drug interactions due to convergent induction or inhibition of cytochrome P450 enzymes poses a risk for extraneous adverse events in the context of polytherapy if such considerations are not accounted for in advance. Such risk/benefit assessments should be based on metabolic pathway and pharmacokinetics. The newer generation ASDs, at least those that do not undergo hepatic metabolism, are thought to possess the least potential for pharmacokinetic interactions as they are not primarily metabolized in the liver [35]. In contrast, phenobarbital, phenytoin, carbamazepine, primidone, valproate, lamotrigine, felbamate, rufinamide, and to some extent topiramate and oxcarbazepine are all associated with a risk of drug-drug interactions due to their route of metabolism [61, 63, 95]. When these drugs are combined, pharmacokinetic interactions may confound apparent pharmacodynamic effects and mask any possible synergism (or antagonism). That said, pharmacokinetic interactions can also be beneficial in a therapeutic sense, allowing a more rapid attainment of steady state with long half-life drugs in enzyme-induced patients (*i.e.* with zonisamide; [81]), or combining a beneficial pharmacokinetic interaction with a pharmacodynamic one (*i.e.*, valproate with lamotrigine). Such understanding of pharmacokinetics and drug interactions is essential when evaluating combination therapy and when reporting potential synergism between

ASDs, which will invariably be interpreted as being pharmacodynamic in nature.

Of additional consideration is the influence of genetics on therapeutic response. Siblings with epilepsy show similar responses to monotherapy or polytherapy [84]. There is, thus, strong evidence for a genetic contribution to the pharmacological management of epilepsy although the individual variants that predispose to treatment success or failure in general remain to be identified. However, there is now substantial evidence for specific genetic mutations in certain epilepsies; e.g., greater than 25 “epilepsy genes” have been identified [71], many of which encode the voltage- and ligand-gated ion channels that are also the predominant targets of ASDs [33, 70]. This is most clearly illustrated in Dravet’s syndrome, which is associated with loss-of-function mutations in one allele of the Na<sub>v</sub>1.1 channel [17]. Na<sub>v</sub>1.1 channels are the predominant sodium channels in inhibitory interneurons [58]. Importantly, this information provides a better understanding of why the sodium channel blocker, lamotrigine, can exacerbate seizures in patients with Na<sub>v</sub>1.1 mutations by possibly inhibiting the remaining functional sodium channels in inhibitory interneurons. Of course, this very elegant scientific explanation came long after clinical experience had already taught us to avoid sodium blockers in patients with a Dravet’s phenotype [39]. In the future, however, it may be possible to predict likely treatment response on the basis of drug MoA and the underlying molecular etiology of the epilepsy. In this regard, Dravet’s syndrome provides us with a clear example of the importance of translation (both forward and back) in directing a rational therapeutic approach. In the context of this chapter, it is also interesting that combination therapy with stiripentol, valproate and clobazam appears to be the most effective treatment in patients with Dravet’s syndrome [19]. Finding an effective therapeutic approach with ASDs that do not specifically target the mutations in VGSCs in Dravet’s patients thus illustrates one of the best examples to date of rational polytherapy for genetic epilepsy.

Indeed, several other genetic disorders present with seizures, which, unlike Dravet's syndrome, may be effectively managed with therapies that specifically target the mutated protein or pathway [26, 94, 100]. For example, tuberous sclerosis (TSC) arises as a result of a mutation in one of two proteins in the mammalian target of rapamycin (mTOR) proliferation pathway (TSC1 or TSC2) and which has recurrent seizures as a characteristic phenotype [65]. The mTOR inhibitor rapamycin effectively suppresses aberrant cellular proliferation in TSC [27] and has been proposed to be disease-modifying, although whether adjunct rapamycin will effectively reverse the epileptogenic process and protect against seizures remains to be determined [65]. A similar situation applies to Fragile X Syndrome (FXS), an autism spectrum disorder in which approximately 14 % of patients present with mild seizures [6]. FXS arises due to a triplet repeat in the *FMR1* gene that leads to loss of the RNA-binding protein, Fragile X Mental Retardation Protein (FMRP) [4]. This protein interacts with machinery essential to synaptic plasticity processes mediated by group I metabotropic glutamate receptors (mGluRs), including mGluR5 [4]. Clinical trials are currently ongoing to examine the use of mGluR5 antagonists in the targeted treatment of FXS [40]. Additionally, FXS patients may benefit from a rational polytherapy approach as preclinical studies in *FMR1* knockout mice, which display audiogenic seizures [59], suggest that acute, combined treatment with an mGluR5 inverse agonist and a GABA<sub>B</sub> receptor agonist can synergistically suppress seizures better than either treatment alone [60]. Furthermore, studies are underway to determine whether treatment for FXS could translate into effective means to suppress network hypersynchronization and changes in synaptic plasticity that arise in epilepsy in general.

Our limited experience from Dravet's syndrome, TSC, and FXS suggests that understanding the molecular etiology of epilepsy can promote rational therapeutic approaches by identifying pathways for targeted intervention or those that should be avoided. With current large-scale efforts to unravel the genetic contribution to epilepsy, including those coordinated by Epi4K, EpiPGX,

CENet, and the ILAE Genetics Consortium, it is probable that opportunities for rational therapy guided by the underlying etiology of the disorder will expand considerably. For example, emergent evidence suggests that *de novo* mutations of ion channel-encoding genes are prevalent in severe childhood epileptic encephalopathies [15, 33]; this information will then likely be informative to direct personalized treatment strategies for patients with similar mutations. Obviously, the hope of such collaborative research endeavors is that the emerging data will eventually inform clinical practice and could play an important role in individualized therapy. Such observations will further illustrate the need for critical evaluation of the disease characteristics and genetic associations *a priori* before deciding on a rational mono- or polytherapeutic approach. We may not be able to hit every target in every patient, but a better understanding of the disorder from a molecular perspective can only be an improvement over current practice in which most patients are treated from a position of blissful ignorance regarding the cause of their epilepsy.

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## 24.9 Summary and Conclusions

MoA is an important criterion in the selection of ASDs for individuals with epilepsy, particularly in the avoidance of seizure worsening in generalized epilepsies, in the replacement of ineffective monotherapies, and when instituting or adjusting polypharmacy regimens. In all of these cases, therapy (whether mono- or poly-) can be said to be rational when MoA is considered. As detailed above, currently available ASDs often share similar features in their MoA, allowing for selective application of ASDs in certain epilepsy patients. These MoAs dictate how individual drugs behave clinically, in the control of specific seizure types and in their propensity to elicit specific adverse effects, and also how ASDs perform within polytherapy regimens. For patients in whom monotherapy has proved inadequate, current evidence supports the combination of a drug with a single, selective MoA with one that possesses multiple cellular effects. Future effort to understand how

drug combinations work in certain patient populations is clearly of critical importance. In most cases, however, clinical validation of combinations identified in experimental models is lacking and greater efforts should be made to conduct post-hoc analysis of clinical trial data, which may provide essential information to direct basic research efforts, and vice-versa. At present, rational therapy for epilepsy describes the use of existing medications to treat seizure types and syndromes of mostly unknown cause using knowledge of how those medications work and interact at the cellular level. In some ways, it is not surprising that this approach is sub-optimal. Future advances in our understanding of the underlying molecular etiologies of the epilepsies, driven at least in part by current global genomics efforts, are likely to improve rational therapy of epilepsy immeasurably. These rationally applied strategies to mono- and polytherapeutic management will thus be critical to future efforts to better treat the refractory epilepsy patient, as well as the newly diagnosed patient.

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# How Can Advances in Epilepsy Genetics Lead to Better Treatments and Cures?

# 25

Renzo Guerrini and Jeffrey Noebels

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## Abstract

Advances in genetic analysis are fundamentally changing our understanding of the causes of epilepsy, and promise to add more precision to diagnosis and management of the clinical disorder. Single gene mutations that appear among more complex patterns of genomic variation can now be readily defined. As each mutation is identified, its predicted effects can now be validated in neurons derived from the patient's own stem cells, allowing a more precise understanding of the cellular defect. Parallel breakthroughs in genetic engineering now allow the creation of developmental experimental models bearing mutations identical to the human disorder. These models enable investigators to carry out detailed exploration of the downstream effects of the defective gene on the developing nervous system, and a framework for pursuing new therapeutic target discovery. Once these genetic strategies are combined with interdisciplinary technological advances in bioinformatics, imaging, and drug development, the promise of delivering clinical cures for some genetic epilepsies will be within our reach.

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## Keywords

Mutation • Phenotype • Gene testing • Comorbidity • Modifier • Complexity

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## 25.1 Introduction

The last decade has witnessed several revolutions in our ability to understand the genetic basis of the epilepsies and its role in diagnosis and treatment. Ten years ago, only a few genes, mostly for ion channels, had been linked to the appearance of epilepsy in large, multigenerational pedigrees. The general belief was that such families were rare, that the numbers of causative genes for epilepsy were few, and that each of the clinically

defined Mendelian syndromes was the exclusive product of one gene alone. Furthermore, the absence of a positive family history in a patient with epilepsy suggested that a purely genetic etiology was unlikely. Following this logic, it was anticipated that detecting a mutation in one of these genes would be highly predictive of a specific seizure syndrome, and that only a rare, inherited mutation causes disease. Understandably, most investigators concluded that knowledge of the gene defect could lead shortly to dramatically improved treatments, if not a cure, for the disorder.

In fact, none of these pioneering assumptions has proven to be entirely correct. However, as in the field of DNA sequencing, we have entered the 'next generation' of epilepsy genetics, and what began as a search for a few inherited gene errors that could explain why some epilepsies are familial has expanded into a set of powerful research tools and discoveries that have immeasurably accelerated our ability to correctly diagnose and, in some cases, treat the disease. Major strides in clinical phenotyping and classification of epilepsy syndromes have been driven by, and contribute to the identification of, new monogenic epilepsies, both inherited and *de novo* in origin. Advances in neuroimaging have proven critical to the discovery of genes leading to malformations of cortical development. New methods in molecular genetics and gene sequencing have allowed rapid identification of candidate genes for an increasing number of epilepsy syndromes and potential comorbidities, including sudden unexpected death. Advances in genetic epidemiology, genome-wide association studies and whole exome candidate gene profiling have stimulated the analysis of complex genetic traits. The mathematical aspects of these analyses, as well as the emergence of mutation and polymorphism databases and genotype-phenotype correlations, are now included in the growing new field of epilepsy bioinformatics.

In the neurobiology laboratory, identified genes arising from both human and experimental genetic studies now offer an unparalleled opportunity to examine basic mechanisms of the disease. Genetically engineered models enable the electrophysiological validation of a candidate

epilepsy gene using *in vivo* and *in vitro* approaches, and are essential to pinpoint the specific brain networks involved. Stem cells derived from patient's fibroblasts can now be reliably transformed into neurons to evaluate the effects of the mutation on cell biology and signaling within the affected nervous system. Contemporary experimental mouse models not only give investigators the ability to selectively express a predefined human gene mutation in the brain at different stages of brain development, but also to reverse its effects with drugs and other genes. High resolution, chronic imaging techniques using fluorescent reporters of gene expression permit the study of the pathophysiology of a genetic lesion over time, tracking the 'downstream' molecular biology of the seizure pathways. Seizures typically arise after prolonged periods of abnormal neural development, and in these cases where the damage is already done, correcting the actual gene defect may come too late to reverse the epileptic condition. However careful analysis of these secondary changes in the physiology and anatomy of the affected neural circuits may offer a second opportunity to discover a novel target for therapy, fulfilling the promise of a cure.

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## 25.2 The Emerging Picture of Epilepsy Genetics

### 25.2.1 Gene and Mutation Diversity

It is now estimated that genetic factors contribute to at least 40 % of all epilepsies. While there has been considerable progress in identifying genes for Mendelian epilepsies, the extent of genetic susceptibility to more common sporadic epilepsy syndromes remains unknown. Although limited evidence, in both animal models and human disease, has been gathered that susceptibility to epilepsy conferred by specific mutations might be influenced by non-pathogenic alleles at other genetic loci [1, 15] the characterization and validation of susceptibility variants appears particularly complex and requires large-scale collaborative efforts. Moreover, our understanding

genetic susceptibility to a major heterogeneous disorder such as epilepsy would likely be incomplete without reference to a specific syndrome. Over 600 entries for pedigrees showing Mendelian phenotypic inheritance patterns can be found in the Online Mendelian Inheritance in Man (OMIM) database, and genetic loci have been identified in over 160 of these cases. These genes arise not only among the >400 members of the ion channel gene family, but across an extraordinarily diverse group of molecular pathways that also regulate membrane excitability, synaptic plasticity, and rhythmic network firing behavior. Causative genes also include those for presynaptic neurotransmitter release, postsynaptic receptors, transporters, cell metabolism, and importantly, many formative steps in early brain development, such as the proliferation and migration of neuronal precursors, dendritogenesis, synaptogenesis, and glial biology. However, inherited mutations in these known epilepsy genes currently only account for a small fraction of patients. Thus, many additional genes causing seizures are likely to be identified. Within each of these genes, the molecular rearrangements themselves are typically novel, or occur with a very low allele frequency within the epilepsy population. Thus monogenic epilepsies are disorders of many, individually rare, errors in an increasingly broad spectrum of biological pathways.

Most of the idiopathic epilepsies arise sporadically among unaffected family members, or do not appear to follow single gene inheritance patterns. From a purely genetic perspective, this finding may be explained by an inadequate family size, or an underlying complex pattern of multigenic inheritance, or even genetic mosaicism, three possibilities which have long bedeviled the analysis of inherited disease. However a new alternative has arisen from an important insight made over the past decade, and promises a steep increase in our ability to isolate genetic risk of epilepsy in individuals, even in small families – namely, the detection of de novo mutation of single genes or copy number variants of even larger chromosomal regions that encompass them. De novo splice site or nonsense mutations that impair function by removing critical portions

of the encoded protein have been identified at convincingly high frequency within specific epilepsy phenotypes, in particular the SCN1A sodium channel linked to the severe myoclonic epilepsy known as Dravet Syndrome. This realization, along with the recent ability to rapidly sequence and assess gene variation in a large list of candidate genes, will greatly contribute to the personal identification of causative genes in the epilepsy clinic.

### 25.2.2 Phenotype Complexity

Large scale genotype-phenotype studies within monogenic populations have determined that the simple correspondence between genotype and phenotype can break down, resulting in different ages of onset and clinical seizure severity (phenotypic heterogeneity) within those bearing mutations in the same gene. This poor correlation may be due to the many possible structural alterations in the mutant protein leading to either gain or loss of function. However, even in families with single gene inheritance of an identical gene mutation, a degree of complexity remains, as evidenced by ‘unaffected carriers’ of the ‘causative’ mutation. In these cases, the phenotypic variability in such families can be attributed to the presence of polymorphisms in modifier genes influencing the phenotypic expression or, alternatively, to environmental factors.

Conversely, identical clinical phenotypes may be due to different underlying genotypes (genetic heterogeneity). Most of the broad phenotypic categories of seizure disorders are now recognized to arise from mutations in more than a single gene. In some cases, the different genes for a clinical epilepsy syndrome all contribute a single functional heteromeric unit, such as the different receptor subunits ( $\alpha, \beta, \gamma, \delta$ ) contributing to a functional GABA<sub>A</sub> receptor in generalized epilepsy, the pore forming ( $\alpha$ ) and regulatory ( $\beta 1$ ) subunits of the sodium channel in Dravet Syndrome, the different pore forming subunits (KCQ2/3) of the M-current in Benign Neonatal Infantile Epilepsy, or the nicotinic cholinergic receptor subunits ( $\alpha 2, \beta 2, \beta 4$ ) in ADNLFE. In other cases, entirely

separate gene pathways may contribute to a very similar phenotype. This property may ultimately explain not only clinical differences in the seizures attached to each gene and their neurological severity in affected patients, but also their pharmacoresistance in clinical subsets of the disorder.

Pharmacoresistance can itself be considered as a phenotypic trait whose intrinsic mechanisms are, at least in part, influenced by genetic variation. Pharmacogenetic studies have attempted to investigate whether drug resistance is influenced by single nucleotide variants in genes for drug targets, or in other genes related to drug uptake and metabolism, which might explain resistance to drugs [16]. These studies, however, are hampered by serious methodological difficulties, since they do not take into account the causal heterogeneity of 'epilepsy' in the populations studied. This oversimplification is reflected in the assumption that a single mechanism would influence drug efficacy in relation to different mechanisms of epileptogenicity, which should be replicated across multiple studies. However results from such studies have not been consistent. For example, a single intronic nucleotide polymorphism in the *SCN1A* gene was associated with higher prescribed doses of phenytoin and carbamazepine in a UK based study [29], but not in subsequent studies in Austria [34] and Italy [20]. Likewise, studies exploring how gene variants may influence AED penetration into the brain have provided conflicting results [2]. Very large studies on etiologically and ethnically homogeneous populations would be necessary to fully explore the real influence of specific genes on pharmacoresistance.

### 25.2.3 Discovery of Novel Comorbidity Syndromes

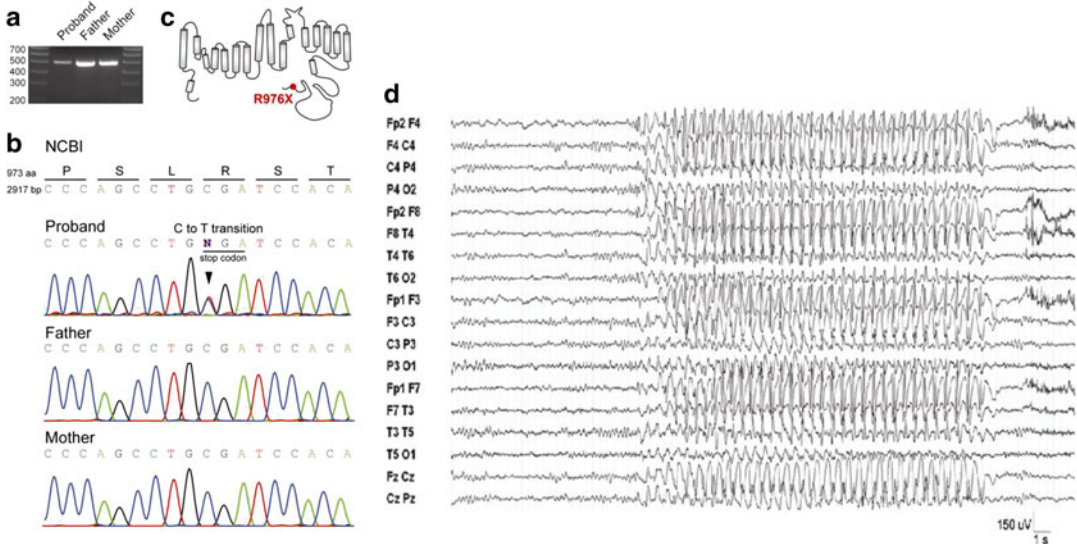
Epilepsy clinicians have long recognized the frequent association of a variety of cognitive and neuropsychiatric symptoms with seizures, and understood that these occur more often than would be predicted by chance. Whether these result as a direct developmental effect of the cause of the epilepsy, the seizures themselves, or

their treatment will always be under debate. Since their co-expression is usually incomplete, other genes may also contribute to the relative penetrance of the co-morbidity.

In the laboratory, mouse models of apparently unrelated disorders, such as Alzheimer's disease, have delivered firm evidence that single genes can produce both epilepsy and cognitive defects unrelated to antiepileptic treatment, and suggest that antiepileptic treatment may be especially neuroprotective in carriers of these genes [24, 25]. Sudden unexpected death (SUDEP) is another important comorbidity, affecting individuals with idiopathic epilepsy. Recent evidence has confirmed the hypothesis that genes underlying cardiac arrhythmias are co-expressed in brain and produce epilepsy [10]. These genes may prove clinically useful in predicting SUDEP risk and exploring treatments to prevent premature lethality in epilepsy patients.

A great deal of interest is now devoted to understanding the developmental causes of epileptic encephalopathies, in relation to autism spectrum disorders (ASD) and intellectual disability. Exome sequencing in a large cohort of individuals with epileptic encephalopathies and subsequent protein-protein interaction analysis revealed a high interconnectivity between genes carrying de novo mutations, with a much greater probability of overlap with ASD and intellectual disability exome sequencing studies [7].

Finally, the ability to probe the full genomic variant profile of unrelated epilepsy patients with and without comorbidities holds enormous promise in understanding the genetic roots of comorbidity. Recently, one such study identified a de novo truncation of the skeletal muscle chloride channel, *CLCN1* in a young woman with a childhood writer's cramp and longstanding pharmacoresistant seizures. This genomic analysis led to the unexpected discovery that *CLCN1* is expressed not only in skeletal muscle, but in thalamocortical and cerebellar brain networks, where disruption of chloride-mediated membrane repolarization could lead to hyperexcitability and seizures [5]. This hypothesis-generating study is a harbinger of the kind of novel candidate gene discovery that awaits the widespread use of next generation sequencing (Fig. 25.1).



**Fig. 25.1** Detection of heterozygous de novo nonsense mutation in *CLCN1*, encoding a premature stop codon in the CLC-1 chloride channel protein in a proband with generalized tonic clonic and absence seizures with a subtle myotonic phenotype. (a) PCR amplification of the final coding exon (exon 23) in the trio yielded a 550 bp product. (b) Sequence chromatograms for the trio

shows the heterozygous base pair substitution encoding a premature stop codon in the proband, but not in either parent. (c) Schematic diagram of a single alpha-subunit of the CLC-1 channel protein showing the location of the C-terminal truncation R976X mutation in the proband. (d) Typical absence seizure in the proband (From Chen et al. [5])

### 25.2.4 Gene Testing

Recent identification of causative genes for a number of early-onset severe epilepsies has created the opportunity for diagnostic genetic testing in this population. Some examples include brain malformations and epileptic encephalopathies of infancy. At present, the clinical impact of genetic testing in these syndromes is by itself limited, due to the small percentage of patients in whom a single, causative gene mutation can be identified and the lack of specific, gene-directed, treatment options. However, genetic counseling can certainly be improved by recognizing a specific etiology. It also sets the stage for further research advances in understanding how each of the genes give rise to epileptogenic defects, and discovering which of these may be reversible. It has been claimed that, in some cases, discovery of a single causative gene defect may reduce the need for further diagnostic investigation at the biochemical level. However, from a practical standpoint, since genomic variants require time to analyze, this information typically arrives after

reversible causes have been clinically excluded. It is also essential to understand that recent profiling studies of whole genomes and large sets of candidate exomes such as ion channel genes have determined that patients with sporadic epilepsy often carry more than one potentially causative mutation [17], complicating the interpretation. Furthermore, all individuals carry, on average, 50–100 loss of function variants in disease genes that for the most part produce no apparent clinical effects [30], signifying that the mere presence of a variant does not predict clinical status. This is likely explained by the presence of other ‘protective’ modifier genes. Thus, as we gain access to a broader view of the genetic landscape in individuals with epilepsy, we expect to routinely encounter patients with a complex genetic basis for their seizure phenotype.

The next steps toward increasing the power of genetic testing in epilepsy include identifying more genes for monogenic epilepsies, and learning to understand the contribution of specific genes in epilepsies with complex inheritance. This will require continued genotype-phenotype



correlations, coupled with functional studies of the abnormal proteins to more accurately understand the pathophysiological implications of each new mutation and how they combine to create neural excitability phenotypes. While bioinformatic analysis offers an increasingly powerful way of categorizing the potential damage a gene variant may inflict on protein function, it cannot conclusively predict its actual effect upon a neuron, and indeed, despite being expressed in multiple cell types, it may not affect them all equally. However we are now entering the era of 'personalized' mutation analysis, where the mutant functional defect can be determined directly in the patient's own cells, sometimes finding that it is counter to the expected result. For example, a currently held hypothesis for the mechanism of epilepsy in Dravet Syndrome, as studied in a mouse model of *Scn1a* haploinsufficiency, is based on the failure of interneurons to fire adequately in the face of reduced sodium current through *Scn1a* ion channels [31]. Analysis of membrane excitability in stem-cell derived neurons transformed from a Dravet Syndrome patient showed that the mutation, predicted to reduce the density of functional sodium channels, resulted in increased sodium current and hyperexcitability in cells classified as both excitatory and inhibitory, implying a distinctly different pathogenic mechanism [19]. These studies may have practical implications for diagnosis, genetic counseling and possible treatment, as well as increasing our knowledge of normal brain function and mechanisms of epileptogenesis.

We can now consult lists of epilepsy syndromes in which a chromosomal locus or loci have been mapped, and those in which one or more gene mutations or variants have been identified. These lists are constantly expanding as new loci and genes are identified. Our view on the correlations between phenotype and genotype in genetic epilepsies is also rapidly changing in relation to new findings emerging from exome sequencing and the use of diagnostic panels as unexpected phenotypes become associated with mutations of specific genes and vice-versa. A constantly updated database will be essential to establish all the known gene mutations and

polymorphisms and their clinical correlates, so that genotype-phenotype correlations can be determined. This is the objective of the Human Variome Project [13] and an achievable goal for epilepsy genetics. For example, over 700 mutations in *SCN1A* have now been reported in the *SCN1A* Variant Database [14] and offer the possibility of predicting the onset, if not the severity, of the related phenotype with a reasonable likelihood.

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### 25.3 Does Knowledge of a Specific Genetic Cause Influence Treatment?

The appropriateness or inadvisability of a given treatment in a specific condition has in some instances been acquired in clinical practice and then scientifically justified by genetic knowledge. For example, the potential for lamotrigine, a sodium channel blocker, to aggravate seizures in Dravet syndrome was initially reported well before the discovery that loss of function *SCN1A* mutations were the cause of the syndrome [11]. However from a clinical perspective, a number of specific conditions provide evidence that improved understanding of epilepsy genetics, together with enhanced knowledge of molecular pathology and electroclinical characteristics, substantiate more rational and effective treatment choices resulting in better patient management. In cases where genotype-phenotype correlations suggest that the epilepsy may have a benign course, gene testing may support the decision to withhold antiepileptic drug therapy during critical periods of brain maturation. Examples of this are mainly related to benign familial epilepsies starting in the first years of life due to *PRRT2*, *KCNQ2* and *SCN2A* gene mutations [33].

Only in very rare conditions, however, do the treatment choices specifically target the inherited pathophysiological mechanism. An interesting example is represented by autosomal dominant nocturnal frontal lobe epilepsy (ADNFLE) due to mutation of neuronal nicotinic acid acetylcholine receptor alpha subunit in which the therapeutic effect of nicotine patch treatment on refractory

seizures was elegantly albeit anecdotally demonstrated using an N-of-1 trial [32]. Other studies have confirmed that transdermal nicotine administration may be a suitable treatment option for patients with ADNFLE and severe seizures [4]. However, the translational dimension of these observations is limited as nicotine is highly addictive and may cause cardiovascular effects. More specific examples derive from clinical conditions in which a rationale therapeutic approach is prompted by administering a substance that can correct a metabolic defect that causes epilepsy. Pyridoxine-dependent epilepsy, for example, is an autosomal recessive disorder in which seizures manifesting in the neonatal period or in infancy can only be controlled after administration of high doses of pyridoxine [12]. If untreated, the disorder can lead to life threatening status epilepticus. Affected patients require lifelong pyridoxine supplementation but antiepileptic medication is usually unnecessary. While prognosis for seizure control is excellent in most patients, neurodevelopmental impairment is often present and although it has been suggested that children who are treated early have a better outcome, this is not always the case [9]. Pyridoxine-dependent epilepsy is likely underdiagnosed and for this reason in many centers pyridoxine administration is part of a treatment protocol for neonatal seizures. Pyridoxine dependent epilepsy is caused by mutations in the *ALDH7A1* gene, which encodes for an aldehyde dehydrogenase (antiquitin) acting in the cerebral lysine catabolism pathway [21]. Affected individuals have  $\alpha$ -amino adipic semialdehyde (AASA) levels, which cause an intracellular reduction in the active vitamin B6 co-factor pyridoxal-5'-phosphate (PLP) and a concomitant imbalance of glutamic acid and  $\gamma$ -aminobutyric acid (GABA). Folinic acid-responsive seizures are very similar to PDE [8]. Early seizures can also be caused by deficient pyridox(am)ine 5'-phosphate oxidase (PNPO), which respond to pyridoxal-5'-phosphate supplementation [9].

A third important example of a direct link between genetic diagnosis and effective treatment choice is represented by the use of the ketogenic diet in the treatment of the GLUT1 deficiency

syndrome, an autosomal dominant disorder due to a mutation in the *SLC2A1* gene. Brain glucose transport occurs by facilitated transport, predominantly via GLUT1, located on the blood-brain barrier endothelium, *SLC2A1* mutations result in insufficient transport of glucose into the brain. Patients with GLUT1 deficiency had originally been described as exhibiting a severe neurological syndrome with early intractable seizures, followed by developmental delay, microcephaly and paroxysmal or continuous dyskinesia. This condition was initially identified and subsequently diagnosed in clinical practice, based on low glucose levels in the CSF (hypoglycorrachia) in the setting of normal serum glucose or of abnormal CSF/serum glucose ratios [6]. However, a wide range of variants have been described, resulting in variable degrees of impairment of glucose transport [23], complicating the utility of the genetic information in clinical practice [18]. Neurologic consequences of GLUT1 deficiency presumably arise from disordered brain energy metabolism, secondary to reduced transport. D-glucose is the main fuel for the brain, although alternative fuels such as ketone bodies can be used. The treatment of choice for GLUT1 deficiency syndrome is a diet that mimics the metabolic state of fasting and provides ketones as an alternative fuel for the brain, effectively restoring brain energy metabolism. The ketogenic diet is a high-fat, adequate protein, low carbohydrate diet that provides 87–90 % of daily calories as fats and is used in the treatment of drug resistant childhood epilepsy. As the developing brain requires substantially more energy in young children, patients with GLUT1 deficiency syndrome should be started on the diet as early as possible and should remain on the diet at least until adolescence. Although some patients with milder seizure disorders may respond to antiepileptic medication, most do not and seizure response to the ketogenic diet is remarkable. Also, some pharmacological agents such as phenobarbital and diazepam, impair GLUT1 function and should be avoided [3]. In the past few years, the range of clinical epilepsy phenotypes where GLUT1 mutations and a positive response to the ketogenic diet have been identified is expanding

[22, 26–28], raising the possibility that the gene acts as a modifier of other coexisting abnormalities. The usefulness of the ketogenic diet in GLUT1 deficiency syndrome was demonstrated before the syndrome was linked to mutations of the GLUT1 gene, based on the expected pathophysiological consequences of low levels of glycorrachia, however, the possibility of uncovering GLUT1 mutations in patients with atypical clinical presentations of GLUT1 deficiency and even borderline or normal levels of glycorrachia, has brought about invaluable advantages for the diagnosis and treatment of this disorder.

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## 25.4 Conclusion

In the epilepsy clinic, genetic analysis has revealed not only the presence of more monogenic epilepsy syndromes, but, thanks to continually emerging genome-phenome correlations, is also pointing the way to earlier and more accurate diagnosis. Gene-specific classification of patients will aid clinical stratification to tailor more relevant diagnostic testing and better characterization of the natural history of the disease, enabling improved outcome prediction and genetic counseling for family planning. Future clinical treatment trials will almost certainly include genomic characterization to enhance the detection of a drug response signal as well as any gene-linked adverse effects. The relative contributions of major categories of genetic influence, including inherited monogenic epilepsies, de novo mutations, and sporadic individuals with complex multigenic inheritance are under exploration in many different seizure types and are beginning to inform genetic counseling in the neuropediatric setting. The definitions of classical epilepsy syndromes have been enlarged to account for multiple genetic etiologies, and novel comorbidity syndromes.

In the epilepsy neurobiology laboratory, genetics continues to reveal mechanistic insight into the rich biological diversity of gene defects leading to epilepsy phenotypes. Genes linked to epilepsy have opened the door to understanding the neurobiology and pathology of epilepsies and localizing the vulnerable neural pathways at the

molecular, cellular, and functional levels. Defects in ion channels and a broad range of other cellular signaling pathways including receptors, transporters, and proteins for exocytotic release of neurotransmitters now constitute primary classes of epileptogenic mechanisms. A second major category of epilepsy genes involves transcription factors regulating the early migration and maturation of interneurons, and a third group controls metabolic functions within the cell and neuron-glia relationships.

Mouse models bearing mutations in each of these gene-delineated pathways allow us to examine the fine details of how they alter developmental plasticity in the epileptic brain, to learn when and where cellular pathology arises, and how it spreads to alter excitability in cortical networks through remodeling of gene expression and synaptic reorganization. Mice engineered to conditionally express gene mutations in specific circuits provide information on which circuits are necessary or sufficient to produce the seizure phenotype. Finally we can learn whether there are critical developmental stages for correcting or reversing the gene defect, exactly what the desired drug effect should be at the cellular level, and which molecular targets are most effective in preventing the epileptic disorder. Taken together, these advances hold great promise for improving the clinical management of seizure disorders.

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# How Might Novel Technologies Such as Optogenetics Lead to Better Treatments in Epilepsy?

# 26

Esther Krook-Magnuson, Marco Ledri, Ivan Soltesz, and Merab Kokaia

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## Abstract

Recent technological advances open exciting avenues for improving the understanding of mechanisms in a broad range of epilepsies. This chapter focuses on the development of optogenetics and on-demand technologies for the study of epilepsy and the control of seizures. Optogenetics is a technique which, through cell-type selective expression of light-sensitive proteins called opsins, allows temporally precise control via light delivery of specific populations of neurons. Therefore, it is now possible not only to record interictal and ictal neuronal activity, but also to test causality and identify potential new therapeutic approaches. We first discuss the benefits and caveats to using optogenetic approaches and recent advances in optogenetics related tools. We then turn to the use of optogenetics, including on-demand optogenetics in the study of epilepsies, which highlights the powerful potential of optogenetics for epilepsy research.

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## Keywords

On-demand • Responsive • Channelrhodopsin • Halorhodopsin • Arch • AAV • Optrode • Seizure

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## 26.1 Introduction

By enabling unprecedented possibilities for studying the cell populations and networks involved in seizure initiation, propagation, and termination, recent technological advances open exciting avenues for improving the understanding of mechanisms in a broad range of epilepsies. Through optogenetics, modulation of select cell populations is possible at specific times, providing the opportunity to not only record neuronal activity during seizures, but also to manipulate neuronal activity. In this way, it is possible to probe critical networks and circuits, and identify potential new therapeutic approaches. This chapter focuses on the development of optogenetics and on-demand technologies for the study of epilepsy and the control of seizures.

Optogenetics is a rapidly evolving field providing powerful tools for neuroscience [19, 72], including the study of epilepsies [12, 46]. Optogenetics is a technique in which light-sensitive proteins, called opsins, are introduced into cells. In this way, it is possible to control the activity of neuronal populations by shining light and activating the opsins. Opsins can be light-sensitive channels, pumps, G-protein-coupled receptors, or even transcriptional effectors [47]. We focus on light-sensitive channels and pumps whose activation can inhibit or excite neurons, and first discuss the benefits and caveats to using these optogenetic approaches, as well as recent advances in related tools. We then turn to the use of optogenetics in the study of epilepsies specifically.

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## 26.2 Optogenetics: Development and Technical Advances

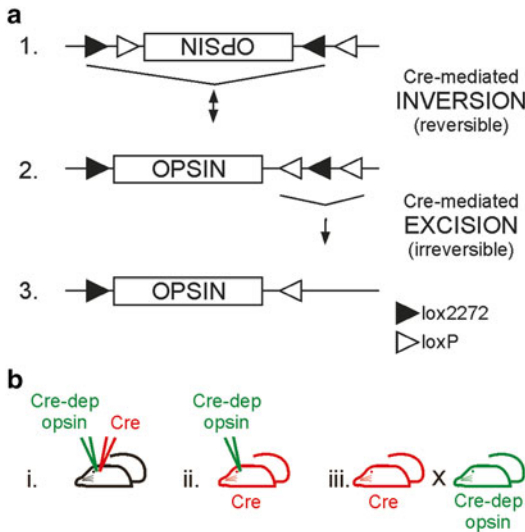
Cell-type and temporal precision are two key strengths to optogenetic approaches. Temporal precision is achieved by appropriately timed light delivery (though, of course, this can present its own challenges, as discussed below for on-demand approaches). Selective opsin expression is less straightforward, and is achievable through

distinct methods. In general, expression is often achieved through the use of viral vectors, (including adeno-associated virus (AAV) or lentivirus), electroporation [2], or the use of transgenic animals. Inducible expression [79, 80, 103] and selective expression of opsins can be achieved for specified populations of neurons defined by their neurochemical profile (e.g., expression of parvalbumin), developmental origin [20, 25, 59, 81], their date of birth (e.g., through the use of retroviruses which only infect actively dividing cells [83]), their location (e.g., by injecting virus in a restricted region), levels of activity at a specific time [36], or their long-distance projections (e.g., through the use of WGA-Cre, which is retrogradely transported transsynaptically [34]).

### 26.2.1 Selective Cell-Type Expression

To achieve selective expression in neurons defined by their neurochemical profile, two broad methods are used. The most straight forward approach is to place the expression of the opsin under a specific promoter (or even enhancer [88]). However, this approach has three disadvantages. First, especially when used with viruses, leaky expression is often noted (that is, expression in other cell populations). Second, long promoters do not fit in small vectors (e.g. adeno-associated viruses (AAV)). Third, in cases where the promoter is a relatively weak promoter, the expression of opsins can be insufficient to achieve strong light-induced currents and alter the activity of the neurons.

In order to overcome these drawbacks, a second method was developed: the opsin is instead placed under a strong promoter, and selectivity is achieved through the Cre/loxP system. Cre can mediate either inversion (flipping) or excision (removal) of DNA, depending on the relative orientations of the loxP sites. For viruses, attempts at selective expression through the introduction of a floxed STOP cassette (which would be excised by Cre) can produce leaky expression (expression even in cells not expressing Cre). Additionally, attempting selective expression through a single inversion (which is then flipped



**Fig. 26.1 Strategies for selective opsin expression.** (a) The FLEX system makes use of two pairs of loxP sites (*triangles*), including the mutated lox2272 (*dark triangles*). Cre mediates inversion using one set of loxP sites (for simplicity, only the inversion using lox2272 sites are illustrated), flipping the opsin sequence into the correct orientation (stage 2). Cre-mediated excision of one of each loxP sites locks the vector in an active state (stage 3) (Based on Figure 1 from Ref. [9]). (b) Three potential ways to achieve selective opsin expression include (i) injecting a Cre-dependent virus (as in a) and a Cre-delivering virus (e.g., WGA-Cre, as further discussed in the text [34]), (ii) injecting a Cre-dependent virus into a mouse expressing Cre in a subset of neurons, or (iii) crossing a mouse line expressing Cre in a subset of neurons with a mouse line expressing opsins in a Cre-dependent manner

by Cre to allow transcription) can produce weak expression, as Cre can continue to mediate flipping, re-inverting the sequence and inhibiting transcription. Therefore, a FLEX system ('flip-excision' [9, 71], also referred to as DIO – double-flxed inverse open reading frame [99]) was implemented (Fig. 26.1a). In this scenario, two sets of loxP sites are used. For one, a mutated sequence is used – lox2272. This sequence is still recognized by Cre, but is only paired with a similarly mutated sequence [51]. Therefore, two distinct sets of loxP pairs can be achieved (one set carrying the mutation, and one set not). One round of Cre-mediated recombination flips the sequence, and another excises one of each type of loxP site, preventing future recombination and

locking the virus in its activated state. This method has proven effective in achieving specific opsin expression, as well as sufficient levels of opsin expression [9]. Cre can be introduced by several methods, including virus injection (note that only low levels of Cre expression are needed). WGA-Cre, mentioned above, can be used to achieve selective expression based on axonal projections [34]. For example, a FLEX-opsin virus can be injected into the hippocampus contralateral to WGA-Cre virus injection, to achieve opsin expression selectively in hippocampal neurons projecting contralaterally, e.g., mossy cells [34]. Alternatively, Cre-dependent virus can be injected into a transgenic mouse (or rat) line expressing Cre in a select population of neurons.

There is a wealth of transgenic mouse lines available, including an ever-growing resource of Cre lines [80], many of which are commercially available (e.g., the Jackson laboratory Cre Repository: cre.jax.org). In addition to being useful in combination with Cre-dependent viral-based opsin expression methods, Cre lines can be crossed with lines expressing opsins in a Cre-dependent fashion [56] (Fig. 26.1b). For example, the Ai32 line developed at the Allen Institute expresses the excitatory opsin channelrhodopsin fused to an enhanced yellow fluorescent reporter protein (ChR2(H134R)-EYFP) from the endogenous *Gt(ROSA)26Sor* locus (a locus active in most cells) with expression enhanced with a CAG promoter [56]. Cre mediates removal of a floxed STOP cassette, and allows expression of the opsin.

An important caveat for Cre-mediated selectivity is that excision of DNA (e.g., removal of the STOP cassette) is permanent, even if Cre-expression itself is transient. This means that opsins can be expressed in cells that are not (currently) expressing Cre. Indeed, even if the cell is simply descended from a cell in which recombination has occurred, opsins will be expressed. This caveat can have significant experimental consequences. For example, following seizures, somatostatin (a neuropeptide whose expression is often used as a biochemical marker for populations of inhibitory interneurons) is transiently expressed in principal cells

[26]. If selective opsin expression in somatostatin-expressing interneurons is being achieved through a Cre-dependent mechanism, selectivity of expression will be (permanently) lost following a seizure.

Another major limitation of available methods for achieving opsin-expression selectivity is the current inability to achieve selectivity in a population defined by multiple characteristics. For example, within a broad neuron population defined by a single neurochemical marker, there are several distinct cell-types. In the hippocampus alone, axo-axonic (also referred to as chandelier cells), dendritically targeting bistratified cells, and a subset of basket cells (which target the perisomatic region of postsynaptic cells) all express the calcium binding protein parvalbumin [8, 30, 41, 45]. Therefore, selective opsin expression in parvalbumin-expressing neurons still results in expression across multiple cell-types. Additionally, there are interneurons that are defined in part by expression of proteins which are also expressed by principal cells. For example, subsets of interneurons express the neuropeptide cholecystokinin (CCK) [30, 45, 53]. However, as principal excitatory cells can also express CCK, selective expression in interneurons cannot be achieved through a Cre-mediated mechanism alone.

Importantly, this is a limitation of current methods which can be overcome through intersectional transgenics [80]. By combining the powerful Cre/loxP system with the Flp/Frt system (an analogous, but distinct, recombination system), it is possible to require expression of two markers for opsin expression. For example, Cre expression could be placed under the CCK promoter (and thus expressed in CCK-expressing cells) and Flp placed under an interneuron-specific marker. Indeed, selective expression of fluorescent proteins has already been achieved in CCK interneurons by using such an approach and a RCE-dual reporter mouse line [80]. However, in order for such an approach to be used for selective expression of opsins, mouse lines or viral vectors requiring both Cre and Flp for opsin expression will need to be generated. Additionally, while there is a vast resource of Cre lines, Flp-lines

are markedly scarcer, and the field would certainly benefit from an increase in this resource. Note that beyond allowing access to relatively selective expression in more interneuron types (including neurogliaform and ivy cells, the numerically most dominant interneuron cell type in the hippocampus [7, 16, 31]), intersectional transgenic approaches could also overcome the loss of selectivity for somatostatin interneurons following seizures (described above).

In order to apply optogenetics in humans, a viral-based approach will clearly have to be used. Note that viral vectors have been used in humans, including in the brain [10, 58, 62], and gene-delivery in general is being considered for a range of neurological diseases [10, 87]. Beyond optogenetics, gene-delivery itself may be a new approach in epilepsy [67, 74, 94]. Optogenetic tools to modify gene transcription may also one day be used therapeutically [47]. Note that insertional mutagenesis (and the risk for tumor generation) can be avoided by using vectors which remain extrachromosomal (e.g. recombinant AAV).

For animal studies, however, transgenic mouse methods offer several benefits over viral-based approaches. First, injection of virus is an invasive process, which is avoided through a transgenic-only approach. Second, for viral-based expression methods, the level of opsin expression varies depending on the number of copies of viral vector in the cell. Therefore, there can be great cell-to-cell variability in the amount of opsin expression. In some cases expression can be so high that light induces toxic levels of current. Of course, high levels of expression can also be a benefit of viral-based methods, when transgenic lines do not produce strong enough photocurrents. Third, if the site of virus injection and the placement of the optical fiber delivering light to the tissue are improperly aligned, insufficient light may reach the opsin-expressing neurons. In contrast, in the transgenic lines, variability is reduced, even expression is achieved in the select cell population throughout the brain, and spatial selectivity is achieved through the location of light delivery.

In addition to the Cre-dependent opsin expressing mouse lines described above, there



are several mouse lines expressing opsins directly under a specific promoter. This avoids the need for crossing strains, and leaves open the door for other Cre-based manipulations. However, a strong promoter must be used to achieve sufficiently high levels of opsin expression. The currently available lines include mice expressing the excitatory opsin channelrhodopsin under the Thy1 promoter [5]. Many of these mice are commercially available. Finally, there have been recent developments in achieving transgenic optogenetic rats [82], further expanding the possibilities for using optogenetics in epilepsy research.

### 26.2.2 Direction of Modulation of Neuronal Activity

In addition to cell-type specificity, another benefit of optogenetic approaches, over for example electrical stimulation, is the control of direction of modulation of neuronal activity (e.g., excitation versus inhibition).

#### 26.2.2.1 Activation

Two main classes of opsins are available, allowing cell-specific activation or inhibition. Most of the optogenetic tools used for neuronal activation derive from Channelrhodopsin-2 (ChR2), a naturally-occurring, non-selective cation channel expressed by the algae *Clamydomonas reinhardtii*. Upon exposure to blue light (470 nm absorption peak), ChR2 opens and allows passive movement of Na<sup>+</sup>, K<sup>+</sup>, Ca<sup>2+</sup> and H<sup>+</sup> following the electrochemical gradient [63], depolarizing cell membranes, and if the cell is depolarized to threshold, generating action potentials. ChR2 possesses fast activation kinetics, and is able to trigger single action potentials in expressing cells following 1–2 ms light exposure, making it a particularly attractive tool for precisely timed stimulation of neuronal populations. ChR2 was the first opsin successfully expressed in mammalian neurons [19]. Since then, researchers have focused on improving its expression levels and its ON/OFF kinetics, and developed different variants with largely diverse properties. Several com-

prehensive reviews about Channelrhodopsin variants are available in the literature (see references [14, 54, 60]), and new variants are constantly being developed. Here we will focus on some important variants most often used in epilepsy research applications.

The first modification to the original ChR2 sequence, an amino acid substitution at position 134, produced a variant with improved expression levels and larger photocurrents in neurons (ChR2-H134R), but presenting slightly lower deactivation kinetics [33]. The lower deactivation kinetics produces lower fidelity of light pulse to action potential generation at high light stimulation frequencies, such that cell firing may not accurately follow the stimulus (missed spikes and/or multiplet spikes per light pulse). Further research therefore then focused on improving kinetics, to allow activation of neuronal populations at higher frequencies (above 40 Hz) with better fidelity of spike generation. The first variant producing higher consistency of high frequency spike generation was developed by a chimeric combination of ChR1 (another channelrhodopsin from *C. reinhardtii*) and ChR2, and was named ChIEF [55]. ChIEF displayed reduced inactivation during persistent light stimulations and improved fidelity at frequencies higher than 25 Hz. Similarly, one amino acid substitution at position 123 of the original ChR2 sequence, led to the development of ChR2(E123T), or ChETA [37], a ChR2 variant displaying dramatically improved activation/deactivation kinetics, allowing consistent and reliable action potential generation at frequencies up to 200 Hz. However, photocurrents generated by ChETA were somewhat smaller than wild-type ChR2, posing a potential drawback for its successful application *in vivo*. To solve this issue, an additional modification of the amino acid sequence at position 159 resulted in the development of ChR2(ET/TC), an improved ChETA variant combining high temporal fidelity with large photocurrent generation [14]. ChR2(ET/TC) still represents to date the channelrhodopsin with the best performance in terms of spike fidelity generation and amplitude of photocurrents.

All of the ChR2 variants described above display an excitation maximum at around 470 nm, and require blue light for their activation. However, the propagation of light in tissue is directly proportional to its wavelength, with blue light presenting high scattering and low penetration compared to higher wavelengths such as red light. Additional penetration through brain tissue is achieved by avoiding wavelengths absorbed by hemoglobin. For experiments requiring coverage of large brain areas, channelrhodopsin variants with red-shifted absorption maxima are therefore preferred, as they allow activation of an increased number of neurons with lower light stimulation intensity. The first attempt towards generating red-shifted activating opsins was made by cloning VChR1, a channelrhodopsin naturally expressed by the spheroidal alga *Volvox carteri*. VChR1 presented an excitation maximum at 550 nm, but significantly lower photocurrents and expression levels in mammalian neurons when compared to ChR2 [100]. To improve VChR1 photocurrents, researchers created a chimera by substituting helices 1 and 2 of VChR1 with their analogs in ChR1, thereby developing C1V1 [97]. Subsequent modification of glutamic acid residues at positions 122 and 162 (resulting in C1V1-T/T) further improved its photocurrents, and resulted in a channelrhodopsin variant with photocurrents comparable to ChR2(H134R) and excitation maximum at 550 nm. C1V1(T/T) also presented vastly increased light sensitivity, allowing its activation with lower light power, making it especially attractive for *in vivo* studies.

The ChR2 variants described above enable fast and precise activation of neuronal populations, but are not optimal for experiments requiring activation of specific neuronal population over longer time windows (minutes). At expression levels typically achieved in neurons, long time activation would require constant delivery of high power to the tissue, with potential and undesirable heating effects. To allow neuronal activation for longer time periods, a separate class of activating opsins was developed, where a single brief pulse of blue light is sufficient to trigger the channel into its

active state. Channelrhodopsins with these properties were named Step-Function Opsins (or SFOs), and caused depolarization of cell membranes for periods of 30–60 s after 10 ms blue light exposure [15]. Even slower deactivation kinetics were achieved with a Stabilized SFO [97], displaying dramatically improved light sensitivity and a channel deactivation time constant of about half an hour. A major advantage of SFOs is that they can be used to slightly alter the network contribution of different cell types, as the depolarization they provide following light is generally sub-threshold, and therefore does not directly activate expressing cells, but only increases cell sensitivity in responding to physiological network activity.

### 26.2.2.2 Suppression

The second major class of optogenetic tools available for the study of neuronal networks is constituted by opsins able to hyperpolarize the cell membrane and, if strong enough, silence action potential generation. The first opsin shown to inhibit neuronal activity was halorhodopsin (NpHR), a chloride pump driven by orange light and naturally expressed by the bacterium *Natromonas pharaonis*. When expressed in neurons, exposure to orange light (570 nm absorption maximum) causes active pumping of chloride ions into the cell, thereby hyperpolarizing the membrane potential and inhibiting action potential generation [98]. However, expression of NpHR in neurons was not optimal, and it formed aggregates in the endoplasmic reticulum that could lead to cellular toxicity [33]. Further development of the NpHR sequence focused on decreasing aggregates, improving photocurrents and promoting membrane localization. Several rounds of substantial mutagenesis of the original NpHR sequence allowed researchers to develop a variant (named eNpHR3.0) displaying a threefold increase in photocurrents and twofold increase in membrane hyperpolarization effects, together with a significant red shift of its excitation wavelength, making eNpHR3.0 ideal for a varied range of studies involving neuronal silencing [34].

Although halorhodopsin chloride pumps are able to reduce neuronal activity with high efficiency, actively pumping chloride ions into the neurons could have effects on chloride homeostasis, with potential shifts in the effect of GABAergic inhibition via chloride-permeable GABA<sub>A</sub> receptors (i.e., shifting  $E_{\text{GABA}}$ ) [66].  $E_{\text{GABA}}$  is already compromised in epileptic tissue [35]. Increasing the intracellular concentration of chloride by its active pumping via NpHR activation could further exacerbate this phenomenon, and cause a shift in  $E_{\text{GABA}}$  to the point where GABA<sub>A</sub> activation becomes depolarizing [66].

Together with halorhodopsins, a separate class of tools to inhibit neuronal activity was developed from naturally-occurring proton pumps derived from different strains of the bacterium *Halorubrum sodomense*. In contrast to NpHR and its variants, proton pumps hyperpolarize cell membranes by actively transporting protons to the extracellular environment, upon exposure to orange/yellow light. The most widely used proton pumps include Archaeorhodopsin-3 (also called Arch [22]) and ArchT [38]. Both have been shown to be able to successfully inhibit neuronal activity *in vitro* and *in vivo*, including when expressed in the brain of non-human primates [38]. Recently, Arch3.0 and ArchT3.0 variants were developed, using modifications similar to those made to the original NpHR sequence, and yielded proton pumps displaying large photocurrents in neurons and increased action potential silencing effects [60]. Due to the fact that these pumps rely on active transport of protons for hyperpolarizing cell membranes (rather than chloride transport), they would not contribute to the disturbance in chloride reversal potential and GABA<sub>A</sub>-mediated inhibition [66], but may have alternate effects, such as altered pH.

Channelrhodopsins and halorhodopsins or proton pumps can also be expressed simultaneously in the same cells to allow bidirectional control of the cell population of interest [34, 39, 40]. ChR2 and most of its variants are activated by blue light, and are therefore spectrally compatible with NpHR or Arch variants, which are activated by orange/yellow light. Moreover, if particular experimental conditions require simul-

taneous activation of one population and silencing of another, a combination of red-shifted channelrhodopsins could be used together with NpHR or Arch. This could be used, for example, to study the effects of simultaneous pyramidal cell silencing and GABAergic interneuron activation on seizure activity.

Although the expression of opsins can be specific and directed to desired cell populations using the strategies described above, the outcome of neuronal activation and/or silencing in intact networks can be more intricate than perhaps initially expected, due to the extremely complex nature of neuronal circuits. For example, results from *in vivo* experiments using ChR2-mediated light stimulation show some cells being activated (as expected), while others are silenced, likely due to network interactions [38–40]. Similarly, in a study using ArchT activation (expected to inhibit cells), a substantial number of neurons responded to light instead by increasing their firing rate [38]. As epileptic circuits often undergo considerable changes, including axon sprouting and changes in network connections, potentially unexpected network roles should also be considered when using optogenetics with epileptic tissue. Indeed, optogenetics provides a powerful means to explore these changes and their consequences on the functioning of the network in epilepsy. Provided opsin expression remains specific, optogenetics provides the ability to examine the role of specific neuronal populations in health and disease in a manner previously unachievable with techniques such as electrical stimulation.

### 26.2.3 Light Delivery

In experimental conditions, light is delivered by using a variety of different systems, depending on the needs. Sources able to generate light with suitable wavelength and power include lasers and light emitting diodes (LEDs). Laser sources have the advantage of providing light with narrow wavelengths, and therefore do not require filtering. Additionally, lasers can provide high power, even when coupled to small diameter fibers which

are routed through optical commutators. Lasers can be used with mechanical shutters for light on/off switching to avoid delays in reaching maximum power. However, shutters can be expensive, sensitive, and have relatively short life expectancies. A major disadvantage of lasers is their cost. LED light sources are generally more affordable and are becoming increasingly powerful. While LEDs have the disadvantage of delivering light with typical “tails” in excitation spectrum, these can be adequately filtered to ensure proper wavelength excitation. LED sources typically reach maximum power in less than 200  $\mu\text{s}$  even at very high frequency. Therefore, light can be switched on or off by delivering external voltage pulses (rather than via a mechanical shutter) without sacrificing light power.

For *in vitro* preparations light is typically delivered through the lens of the microscope [48, 84, 102], although other methods are also used, including optical fibers positioned in close proximity to the tissue area of interest [50]. For example, small diameter fibers [50] or laser-scanning photostimulation [95] can be used to activate specific regions in the slice, allowing for example circuit mapping and investigation of network alterations occurring after seizures. When light is delivered through the lens of the microscope, filtered light from a mercury or xenon lamp source can also be used, similar to epifluorescence applications.

For *in vivo* situations, light is most commonly delivered through an optical fiber implanted in the region of interest and connected to the light source of choice. Sophisticated light delivery options have also been designed, including multi-waveguides capable of delivering light of different wavelengths to different locations along the guide [104]. Additionally, the optical fiber can be combined with a recording electrode. The combination is termed an optrode, and a number of designs and protocols exist [1, 4, 42, 69, 75, 78, 89, 90, 101], including a recent protocol for simple and relatively low cost optrodes designed for chronic (months long) recordings in rodents [6]. Optical fibers can be directly implanted [6] or guided into the tissue by a cannula previously fixed to the animal’s skull [99]. For long-term *in*

*vivo* applications, an optical commutator is often used to reduce torque on the optical patch cord connecting the animal and the light source. There also exist wireless options for light delivery, including headborne LED devices [90] and injectable  $\mu$ -LEDs [44].

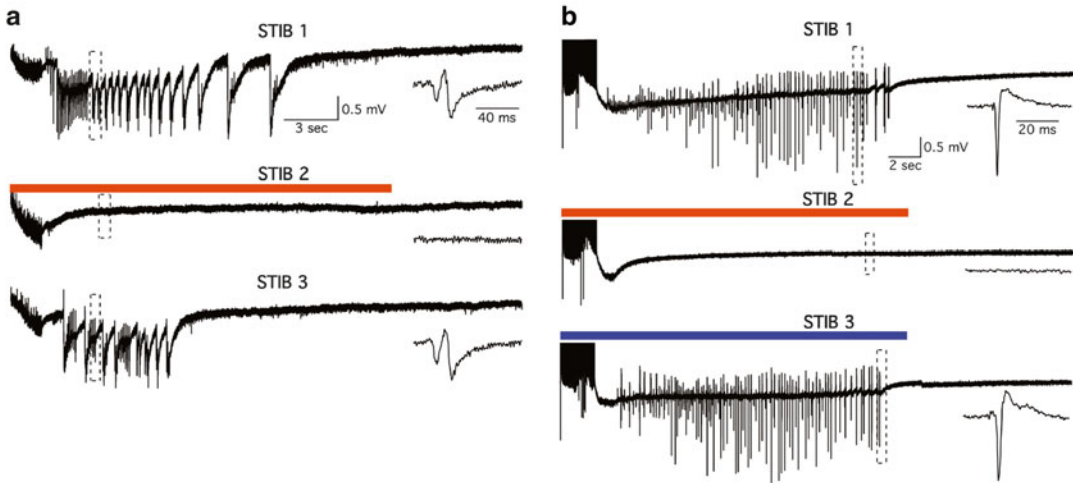
A major caveat to consider while planning *in vivo* optogenetic experiments is that brain penetration by light is rather limited, as described above, and progressively reduces with decreasing wavelengths. Therefore, the spatial distance between the light and the cells expressing the opsin can be critical, and will determine the minimal required power for adequate activation of the transgene. The choice of opsin is also important, as some are several fold more sensitive to light than others, or have red-shifted excitation maximum allowing simultaneous activation (or inhibition) of a large number of neurons while maintaining a small diameter optical fiber (reducing tissue damage).

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## 26.3 Optogenetics: Shedding Light on Epilepsy

### 26.3.1 Review of Recent Studies

The first attempt at using optogenetic approaches for suppressing abnormal hypersynchronized activity involved expression of eNpHR (a slightly improved NpHR protein) in pyramidal cells of the hippocampus [84]. The inhibitory opsin was introduced in excitatory principal cells of organotypic hippocampal slices by using a lentivirus carrying the NpHR transgene under the control of the CaMKII alpha promoter, which is expressed in excitatory cells and absent in inhibitory interneurons. Organotypic hippocampal slices are proposed to represent an *in vitro* model of epileptic tissue, as they exhibit network reorganization, such as cell death, axonal sprouting and synaptic formation, leading to hyperexcitability [3, 11]. The ability of NpHR to inhibit epileptiform activity in such “epileptic” tissue was tested by applying orange light during stimulation train induced bursting (STIB), a stimulation protocol that reliably evokes afterdischarges in



**Fig. 26.2 Optogenetic inhibition of epileptiform activity in vitro.** NpHR expression in excitatory principal cells of organotypic hippocampal slices is efficient in inhibiting stimulation train induced bursting (STIB) when acti-

vated by *orange light*, in both CA3 (a) and CA1 (b) areas. Stimulation with *blue light* failed to alter STIB-induced bursting (b, bottom) (Reproduced with permission from Ref. [84])

the CA1 and CA3 area. Orange light application effectively suppressed STIB-induced activity, while blue light application was ineffective, indicating the specificity of the approach used (Fig. 26.2) [84].

Optogenetic approaches have also had success in inhibiting seizures *in vivo* across a range of epilepsies, including induced (acute) seizures [76], focal cortical seizures [94], temporal lobe seizures during the chronic (spontaneous seizures) phase of the disease [48], and thalamocortical epilepsy in a model of cortical stroke [65].

Using the rat pilocarpine model of acute induced seizures in awake behaving male rats, Sukhotinsky and colleagues examined the ability to inhibit seizures using optogenetic inhibition of the hippocampus [76]. The inhibitory opsin halorhodopsin (eNpHR3.0 [34]) was expressed in principal excitatory cells in the hippocampus using adeno-associated virus (AAV) and a CamKII $\alpha$  promoter. Animals receiving light and expressing the opsin showed an increase in time to seizure onset from the time of pilocarpine injection compared to controls (time to seizure onset with opsin activation:  $21 \pm 1.8$  min versus  $15.2 \pm 1.1$  min in controls). Controls included animals not injected with virus and not receiving

light, animals injected with virus but not receiving light, and animals receiving light but not expressing the opsin. Therefore, the activation of opsins (and inhibition of hippocampal principal excitatory cells) delayed the time to seizure onset. This study supports the notion that optogenetics can be used to inhibit seizures. Moreover, it indicates that targeted inhibition of principal cells in the hippocampus can delay the onset of pilocarpine induced seizures.

Wkyes and colleagues demonstrated the successful use of an optogenetic approach to inhibit focal cortical seizures [94]. Neocortical epilepsy is frequently drug-resistant, and new therapeutic approaches are being actively sought. Focal cortical epilepsy was induced in rats by focal injection of tetanus toxin into the motor cortex. Lentivirus was co-injected with the tetanus toxin, in order to transduce excitatory pyramidal neurons in the epileptic focus with the inhibitory opsin halorhodopsin (NpHR2.0, under a CamKII $\alpha$  promoter). Seven to 10 days after the injection of tetanus toxin, the ability of an optogenetic approach to inhibit seizures was investigated. EEG was recorded for a 1,000 s baseline period, then intermittent light (20 s on, 20 s off) was delivered for 1,000 s, and then a final 1,000 s of post-light EEG

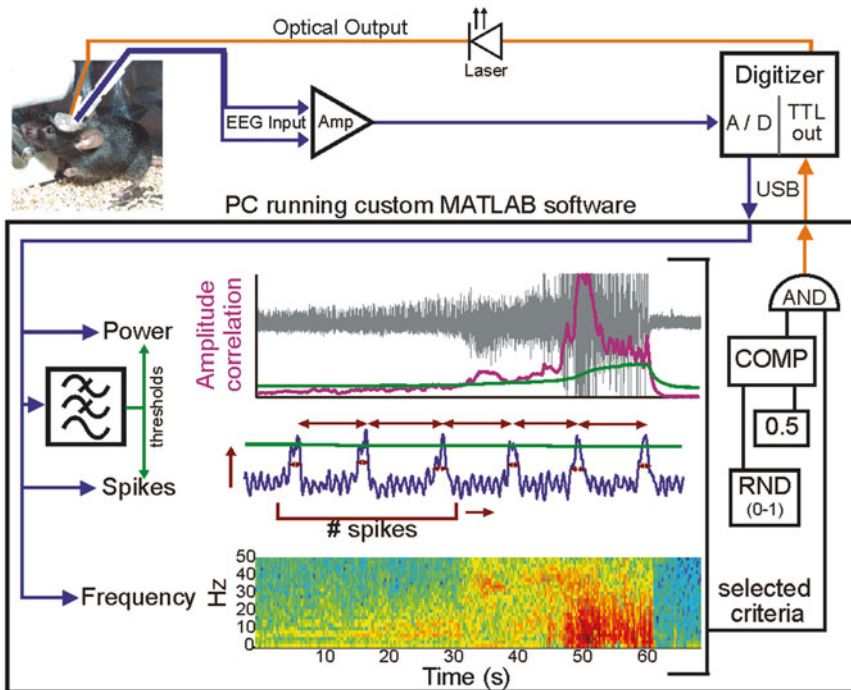
was recorded. Compared to the periods of no-light delivery, opsin activation by light delivery attenuated recorded EEG epileptiform activity. In animals not expressing opsins, light delivery did not affect the high-frequency power of the signal, supporting the conclusion that the light-effect observed in opsin-expressing animals was due to the activation of opsins, rather than of light delivery *per se*. Not only does this study indicate that an optogenetic approach can inhibit focal cortical epileptiform activity, but also that inhibition of a portion of excitatory cells at the focus is sufficient to do so.

These two studies support the potential for an optogenetic approach for diverse epileptic activity, and make use of the power of optogenetics to selectively target specific populations of cells. An additional major benefit of optogenetics is the temporal precision which it can provide. That is, an optogenetic approach could be employed in an on-demand or responsive fashion, such that intervention only occurred either immediately before a seizure would occur (seizure prediction) or early during a seizure onset (seizure detection). In addition to the experimental benefits of an on-demand approach, by limiting intervention to only those times when it is needed, an on-demand approach may reduce negative side effects associated with chronic treatments.

On-demand optogenetics have been used in two models of epilepsy – thalamocortical and temporal lobe epilepsy. Using a cortical stroke model of thalamocortical epilepsy, and line-length threshold crossing for automated seizure detection, Paz and colleagues demonstrated the successful inhibition of seizures [65]. The inhibitory opsin halorhodopsin (eNpHR3.0 [34]) was expressed under a CamKII $\alpha$  promoter in the ventrobasal thalamus ipsilateral to the site of induced cortical stroke. On-demand light activation of opsins interrupted seizures. In addition to illustrating the potential for on-demand optogenetics to stop seizures, these findings supported the theory that the cortical strokes produced thalamocortical seizures; that is, optogenetics can provide insight into the mechanisms of seizures, including critical brain regions and networks.

On-demand optogenetics has also been used successfully in a mouse model of chronic temporal lobe epilepsy [48]. Seizures were detected on-line with custom-designed, tunable, multi-algorithm based detection software (Fig. 26.3). This software, and instructions on how to use the software, is available for download through Nature Protocols [6]. The intra-hippocampal kainate mouse model used mimics unilateral hippocampal sclerosis, and displays both spontaneous electrographic-only seizures (that is, seizures with little or no overt accompanying behavior) as well as seizures that progress to overt behavioral seizures. Seizures were detected early, prior to overt behavior. Selective expression of the inhibitory opsin halorhodopsin (eNpHR3.0) was achieved by crossing mice expressing halorhodopsin in a Cre-dependent fashion with mice expressing Cre under the CamKII $\alpha$  promoter [56]. On-demand light delivery to the hippocampus, inhibiting excitatory cells, dramatically truncated seizures (Fig. 26.4).

Krook-Magnuson et al. [48] then went on to try a second approach. Rather than inhibiting excitatory cells directly through optogenetics, the authors instead used optogenetics to excite a subpopulation of inhibitory neurons. Selective expression of the excitatory opsin channelrhodopsin (ChR2) was achieved by crossing mice expressing ChR2 in a Cre-dependent manner with mice expressing Cre selectively in parvalbumin-expressing neurons. In the hippocampus, parvalbumin-expressing interneurons represent less than 5 % of the total neuronal population [16, 30, 92]. Remarkably, seizures were significantly inhibited through this approach. Seizures were also significantly inhibited when light was delivered to the contralateral hippocampus. Finally, light delivery reduced the number of seizures progressing to overt behavioral seizures. These data indicate that focal light delivery can have a significant effect on temporal lobe seizures, that an on-demand approach can work in temporal lobe epilepsy, and that a strategy directly targeting only a small fraction of cells (that is, parvalbumin-expressing interneurons) can significantly inhibit temporal lobe seizures.



**Fig. 26.3 Schematic of online seizure detection for on-demand optogenetics.** EEG input (blue) recorded from the animal is amplified (Amp), digitized (A/D), and relayed to a PC running real-time seizure detection software. This software is tuned for each animal, with user-defined thresholds (green). Seizure detection algorithms utilize features of signal power (top), spikes (middle), or frequency (bottom). Once a seizure has been detected using the selected criteria, the software can activate, via a TTL signal from the digitizer

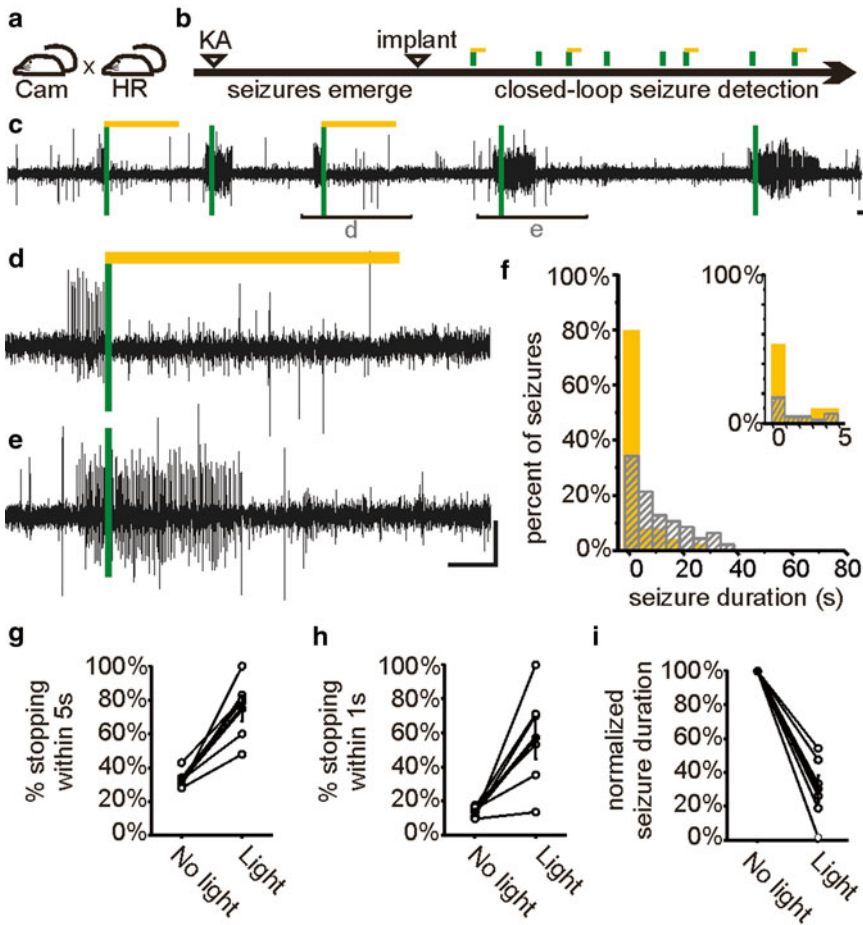
to the laser, the optical output (orange) delivered to the animal. Signal power related calculations (purple, during an example seizure shown in grey), spike characteristics (e.g., amplitude, rate, regularity, and spike width, shown in red), and frequency characteristics (shown for the same seizure, with warmer colors representing higher energy) are illustrated. COMP: digital comparator. This on-line seizure detection software is available for download through reference [6] (Figure reproduced with permission from Ref. [48])

### 26.3.2 Obstacles and Future Potential in Epilepsy Research

While the results from these studies are promising, a number of hurdles need to be overcome before optogenetics, and hopefully on-demand optogenetics, can be realized in the clinical setting. These include demonstration of safe and stable opsin expression in humans, as well as a safe implantable device for on-line seizure detection and light delivery. However, on-demand optogenetics, with its cell-type, spatial, and temporal-specificity, may one day aid patients currently suffering from uncontrolled seizures and the negative side-effects of systemic treatment options. An example patient population that could benefit from the clinical realization of an

on-demand optogenetic therapeutic is patients with refractory bilateral temporal lobe epilepsy for whom surgical resection is not an option.

Optogenetics additionally presents a powerful tool for expanding our understanding of mechanisms of epilepsy. While the studies discussed here have demonstrated a wide potential for optogenetics in the field of epilepsy, there is much more to be gained from fully harnessing the power of optogenetics. Through optogenetics it is possible to test hypotheses regarding critical cell-types and networks involved in the initiation, continuation, propagation, and (natural or induced) cessation of seizures. The studies described above inhibited seizures using optogenetic techniques, but it is also possible to use optogenetic approaches to study mechanisms



**Fig. 26.4 Seizure control in vivo in mice expressing HR in principal cells in a model of temporal lobe epilepsy.** (a) Crossing CamK-Cre and Cre-dependent halorhodopsin (HR) mouse lines generated mice expressing the inhibitory opsin HR in excitatory cells (Cam-HR mice). (b) Experimental timeline. (c–e) Example electrographic seizures detected (vertical green bars), activating amber light (589 nm) randomly for 50 % of events (light: amber line, example in d; no-light example in e). (f) Typical example distribution of post-detection seizure durations (5 s bin size) during light (solid amber) and no-light inter-

nal control conditions (hashed gray). Inset: first 5 s bin expanded, 1 s bin size. Note that most seizures stop within 1 s of light delivery. (g–i) Group CamHR data showing the percent of seizures stopping within 5 s of detection (g), within 1 s of detection (h), and the average post-detection seizure duration (normalized to average no-light post-detection duration for each animal) (i). Note that in one animal (shown in c–e), all seizures were stopped within 1 s of light delivery. Averaged data: filled circles. Error bars represent s.e.m. Scale bars in c–e: 100  $\mu$ V, 5 s (Reproduced with permission from Ref. [48])

of epilepsy through the induction (rather than inhibition) of seizures [64]. The information gained from optogenetic experiments can in turn open the door for new therapeutic approaches beyond optogenetics, including new drugs targeting key cell types or electrical stimulation targeting key brain regions.

While the field is benefiting greatly from recent technological advances, there is a continuing need

for additional developments. A reliable and inexpensive long-term EEG monitoring system, with fully computerized analysis of EEG and video for automated detection and analysis of electrographic and behavioral seizures, would push the field forward dramatically. For example, this would increase the feasibility (and statistical power by allowing more animals to be monitored and analyzed) of studies with mild or moderate head



injury for which only a small subset of animals go on to develop epilepsy. Recent advances in wireless devices, including those capable of delivering light [44, 90], and improvements in seizure detection [6, 91] are paving the way for such future advances. These advances will additionally improve the utility of optogenetics for epilepsy research by allowing chronic on-demand light delivery to freely moving, untethered, animals.

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## 26.4 Other Technical Advances: New Avenues, New Insights

This chapter has focused on optogenetics. Clearly, however, the field takes advantage of a large range of new technological advances, several of which are being rapidly integrated with optogenetics. For example, on-demand approaches (which as described above can be successfully integrated with optogenetics) have the potential to provide both experimental and therapeutic benefits. While electrical stimulation lacks the cell-type specificity of optogenetics, it can provide temporal precision, and thus can also be used in an on-demand fashion. Previously, on-demand electrical stimulation was found to provide superior seizure control in rats [32]. More recently, on-demand transcranial electrical stimulation (TES) was used to reduce spike-and-wave episodes in absence seizures in rats [13]. There is also intense clinical interest in an on-demand therapeutic option, and clinical trials have shown promise (reviewed in reference [93]).

A step beyond early seizure detection is seizure prediction. A recent study in patients with drug-resistant partial-onset epilepsy was able to predict for a subset of patients periods of high seizure risk and periods where the chances of having a seizure were relatively low, based on an analysis of the frequency bands recorded from intracranial EEG [24]. Further supporting the possibility of seizure prediction, changes in multi-unit activity are reported in human patients prior to seizure onset [18, 85]. Unfortunately, there is considerable variability in this early activity from seizure to seizure [17, 18], which may limit the ability to have accurate seizure pre-

diction. However, detecting seizures early (prior to overt behavioral manifestations) and intervening (optogenetically or otherwise) to truncate seizures to this pre-clinical stage could have a large impact on patient quality of life.

Imaging techniques are an additional example of the wide-range of expanding techniques that are being increasingly applied to the study of epilepsy, and include diffusion tensor imaging (DTI, reviewed in reference [28]), magnetic resonance imaging (MRI, which can be combined with optogenetics [52]), positron emission tomography (PET), single-photon emission computed tomography (SPECT, for a review see reference [57]), the new clarity brain [23], calcium imaging and voltage sensitive dyes (for recent reviews see references [28, 77]). Anatomical imaging techniques of neuronal projections in intact brains allow examination of network connections between brain regions in health and disease. Appreciating long-distance network connections, and how these shape local network connections [49, 86], will undoubtedly provide crucial information on seizure propagation mechanisms, as well as potentially mechanisms behind seizure initiation and termination. Functional imaging can reveal local as well as long-distance network dynamics, and is contributing substantially to our understanding of mechanisms in epilepsy. For example, a recent study using calcium imaging of epileptic tissue found not only variability in firing between neurons during epileptiform events, but also variability between epileptiform events, with each event comprised of different patterns of co-activated clusters of neurons [29].

Advances are certainly not limited to seizure detection or imaging techniques. Whole-genome sequencing, which is providing ever-expanding information on the genetics of epilepsies (reviewed in reference [61]), is an excellent example of the driving force that new technological advances can provide to the field. Additional diverse technological advances, including uncaging of GABA [96] and devices allowing focal cooling [68], are introducing unique new opportunities for studying and treating epilepsy. Advances in

recording techniques are providing unprecedented information regarding the activity of neurons during epileptiform events. It is now possible to record from hundreds of units in human epileptic patients (for a discussion of the spike sorting techniques involved see reference [27]), providing a wealth of information on the involvement of neurons in seizures [18, 43, 85]. The novel information gained from these new techniques can aid in seizure detection and prediction discussed above. Importantly, this data can also be incorporated into “big data”-driven large-scale computational models [16, 70]. Hypotheses can then be tested *in silico*, and new hypotheses in turn generated to be tested *in vitro* or *in vivo* (for reviews of computational neuroscience in epilepsy, see references [21, 73]).

From the genetics, to the proteins, to the cell-types and networks critical in epilepsy, advances are being made and insights gained. Optogenetics, together with a vast array of novel technological developments, is expected to continue to light new avenues for studying the mechanisms of the epilepsies.

**Acknowledgements** This chapter on optogenetic approaches to epilepsy highlights the fundamental veracity of Phil’s overarching conceptual framework that placed a major emphasis on the critical importance of rigorous, quantitative mechanistic understanding of epileptic neuronal circuits in order to develop new generations of temporally and spatially selective, more effective seizure control strategies.

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