

Chapter 16

Astroglia and Severe Mental Illness: A Role for Glutamate Microdomains

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Abstract Postmortem studies of schizophrenia have historically been focused on abnormalities of neurons, with only a small but growing body of work focused on astroglia. The limitations of this approach are reflected by the recent failed efficacy of several neuron-centric pharmacological treatments. In this chapter, we will review glutamate neurochemistry and present a novel hypothesis related to astrocyte dysfunction in schizophrenia. We posit that plasma membrane glutamate transporters expressed on astrocytes help partition extrasynaptic regions, where pools of extracellular glutamate may be tightly regulated. These so-called “glutamate microdomains” may impact excitatory neurotransmission via modulation of extrasynaptic glutamate receptors. This hypothesis is supported by structural, biochemical and electrophysiological evidence, which suggests that the fidelity of these domains could be altered in severe mental illness.

Keywords Schizophrenia · Psychosis · Glutamate transporter · Microdomain · Glutamate spillover

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16.1 Introduction

The study of severe mental illness includes materials/data obtained from living patients, animal models, and postmortem studies using brain tissues from afflicted patients. This review will primarily focus on postmortem studies that inform the hypothesis that remodeling of astroglial processes leads to glutamate spillover in glutamate circuits. We also argue that there are specialized extracellular regions, or microdomains, where glutamate levels are tightly regulated and contribute to astrocyte-neuron interactions via modulation of extrasynaptic glutamate receptors. We will present data that argue for an abnormality of extracellular glutamate microdomains secondary to astroglial deficits found in the brain in schizophrenia.

The focus of this chapter on astrocyte-associated changes in schizophrenia represents an important extension of the glutamate hypothesis of schizophrenia beyond the N-methyl-D-aspartate (NMDA) receptor signaling complex. For decades, the prevailing hypotheses of schizophrenia were centered on neurotransmitter receptor dysfunction, first with D2 dopamine receptors, and more recently NMDA subtype of glutamate receptors. Studies focused on these hypotheses explored how changes in neurotransmitter receptors, localized to the postsynaptic density (PSD) or presynaptic structures on neurons, contributed to the pathophysiology of severe mental illness (reviewed in (McCullumsmith et al. 2004a)). In general, the interpretation of the data in these studies is biased towards the view that observed changes in variables such as measures of receptor subunit gene expression reflect a primary deficit in pre- or postsynaptic neurons. In the sections below, we seek to develop a more balanced view of the role(s) of astroglia in the pathophysiology of this often devastating illness.

16.2 Glutamate Neurotransmission

Glutamate Release and Reuptake The process of release, activity as a ligand, and reuptake of glutamate involves three distinct cell types: the astrocyte, the presynaptic neuron and the postsynaptic neuron (Salt et al. 1996). In the presynaptic neuron, glutamine can be converted to glutamate by the enzyme glutaminase, and packaged into vesicles by a family of vesicular glutamate transporters (VGLUT1–3) for release into the synapse (Bellocchio et al. 2000; Takamori et al. 2000). Glutamate is released into the synapse and may occupy and activate ionotropic [NMDA, α -amino-3-hydroxy-5-methyl-isoxazole propionate (AMPA), and kainate] or metabotropic (mGluR1-8) glutamate receptors on both neurons and astrocytes (Hollmann and Heinemann 1994; Hollmann et al. 1994; Salt et al. 1996). Glutamate is rapidly removed from the synapse by a family of plasma membrane excitatory amino acid transporters localized to postsynaptic neurons and astrocytes (Masson et al. 1999b). Recovered glutamate may enter the tricarboxylic acid cycle via conversion to α -ketoglutarate by glutamate dehydrogenase, be converted to glutamine by glutamine

synthetase and transported back into the synapse, or be released into the extracellular space by a variety of mechanisms (Malarkey and Parpura 2008), including the cystine/glutamate antiporter. In astrocytes, glutamate may also be added to this cycle via *de novo* synthesis of glutamate in a pathway involving pyruvate carboxylase and transaminases (Brainard et al. 1989; Patel et al. 2001; McKenna 2011). Recovered glutamate can also contribute to formation of lactic acid. Lactate production is favored in astrocytes, while lactate breakdown is favored in neurons (Stobart and Anderson 2013). Lactate may be efficiently shuttled from astrocytes to neurons, suggesting that lactic acid may be a preferred energy source for neuronal structures enveloped by astrocytic processes (Stobart and Anderson 2013). Finally, several families of novel glutamate receptor and transporter associated molecules regulate glutamate release and reuptake through intracellular signaling mechanisms (Jackson et al. 2001; Lin and Maiese 2001; Marie et al. 2002; Watanabe et al. 2003).

Glial Plasma Membrane Glutamate Transporters The excitatory amino acid transporters (called EAATs in human) are expressed in the plasma membranes of neurons and glia throughout the brain in a region and cell specific manner (Arriza et al. 1993; Utsunomiya-Tate et al. 1996). EAATs mediate glutamate transport by an electrogenic exchange of 3 Na⁺, 1 H⁺, and 1 glutamate (monovalent anion at conditions in the brain) molecule into the cell and 1 K⁺ ion out of the cell, with the net inward movement of positive charges (Zerangue and Kavanaugh 1996; Levy et al. 1998). EAATs are likely homomers comprised of 2–3 non-covalently linked subunits that have 6–10 transmembrane domains (Danbolt 2001). The transporters have specific patterns of cellular localization. EAAT1 and EAAT2 have primarily been localized to astroglia. EAAT3–4 and EAAT5 are primarily localized to neurons and the retina, respectively (Arriza et al. 1997; Danbolt 2001). In the prefrontal cortex, glial transporters (EAAT1 and EAAT2) are predominately expressed in discrete subsets of astrocytes which account for approximately 90% of synaptic glutamate reuptake (Regan et al. 2007). The glial transporters are localized to perisynaptic processes facing the synaptic cleft (Tzingounis and Wadiche 2007). In the rodent, activation of the promoters and expression of EAAT1 (called GLAST in the rodent) and EAAT2 (GLT-1 in rodent) is generally nonoverlapping (Regan et al. 2007). In addition, the GLAST, but not GLT-1, promoter was activated and EAAT1 expressed in oligodendrocytes, suggesting that EAAT1 has a role in myelination and CNS connectivity (Regan et al. 2007). The functional importance, perisynaptic localization, and heterogeneity of expression of the glial glutamate transporters suggests that examination of the expression and function of these molecules may be a high yield target for studies of neuropsychiatric illnesses that involve alterations in glutamate transmission.

EAAT Expression, Processing, and Trafficking EAATs are synthesized in the endoplasmic reticulum and have extensive posttranslational modification in the Golgi, including N-linked glycosidation of at least two sites that are important for homomultimer formation (Kalandadze et al. 2004). EAATs are then trafficked to the plasma membrane where, localization and clustering are regulated by protein-protein interactions and phosphorylation (Conradt and Stoffel 1997; Figiel and Engele

2000; Gamboa and Ortega 2002; Schluter et al. 2002; Figiel et al. 2003). Ultrastructural studies indicate that most EAAT1-2 expression is in the plasma membrane (Chaudhry et al. 1995). EAATs may be removed from the plasma membrane via endocytosis and shuttled back to the cell surface via recycling endosomes, or be targeted for degradation in lysosomes (Nakagawa et al. 2008).

Extracellular Glutamate Glutamate released at the presynaptic terminal diffuses out of the synaptic cleft and may be transported into the astrocyte by EAATs or spillover to extrasynaptic areas (Masson et al. 1999a; Bridges et al. 2012). Extrasynaptic glutamate may also originate from astrocytes via vesicular release, the cystine/glutamate antiporter (also known as the system X_c^-), or other less prominent mechanisms (Montana et al. 2004; Haydon and Carmignoto 2006; Bridges et al. 2012). The cystine/glutamate antiporter releases glutamate and transports cystine into the astrocyte for glutathione synthesis. Glutamate levels in the extracellular milieu are postulated to be tightly regulated, as activation of extrasynaptic glutamate receptors has potent effects. For example, activation of extrasynaptic NMDA receptors promotes initiation of NMDA spikes, while long-term potentiation (LTP) and long-term depression (LTD) can be readily induced in the adult cortex by activation of extrasynaptic GluN2B containing NMDA receptors (Massie et al. 2008; Chalifoux and Carter 2011).

Removal of glutamate from the synaptic cleft may be conceptualized as a two-step process, involving first the high affinity binding of glutamate by perisynaptic transporters, and second the transport of bound glutamate by the transporter across the plasma membrane (Tong and Jahr 1994; Tzingounis and Wadiche 2007). Once bound, glutamate may be “unbound” or released, or alternatively, transported across the plasma membrane (Tong and Jahr 1994; Tzingounis and Wadiche 2007). The “capture efficiency” of the EAATs is defined as the ratio of the rate of unbinding of glutamate to rate of transport, which is reported to be about 0.5 (Tzingounis and Wadiche 2007). The relatively low rate of transport of bound glutamate compared to the capture efficiency suggests that the EAATs first act as buffers for released glutamate (Tzingounis and Wadiche 2007). Thus, glutamate molecules may bounce from one transporter binding site to another, until transported, limiting glutamate spillover from the synaptic cleft.

Glutamate Spillover The density of perisynaptic glutamate transporter protein, the amount of glutamate released, and the rate of glutamate transport determine, in part, the kinetics of glutamate diffusion away from the synaptic cleft. While several regions have well characterized glutamate spillover between excitatory synapses (such as the cerebellum and hippocampus), there is ongoing debate regarding the extent of glutamate diffusion in other regions, including the frontal cortex, where spillover of glutamate may detrimentally lead to loss of input specificity and activation of cell death pathways (Kullmann and Asztely 1998; Hardingham and Bading 2002; Hardingham et al. 2002; Lozovaya et al. 2004a; Tsvetkov et al. 2004; Marcaggi and Attwell 2007; Leveille et al. 2008). Under physiologic conditions, release of glutamate may exceed the capacity of cortical synapses to remove glutamate from the cleft (Weng et al. 2007; Drew et al. 2008). Mathematical models suggest

that glutamate may diffuse and activate NMDA receptors within a radius of 0.5 μm from the release point (Rusakov and Kullmann 1998). Thus, the spatial arrangement of glutamate synapses, their glutamate transporter buffering zones, and extrasynaptic glutamate receptors will determine the extent and effect of glutamate spillover (Lehre and Danbolt 1998; Sem'yanov 2005; Weng et al. 2007).

Glutamate Microdomains We posit that glutamate microdomains are formed by specialized protein clusters on the membranes of astrocytic processes apposed to extrasynaptic glutamate receptors expressed on specialized regions of neuronal membranes (Fig. 16.1) (Grosche et al. 1999; Genda et al. 2011). Diffusion of glutamate between domains or domains and synapses would be limited by the dense expression of glutamate transporters between these specialized structures (Lehre and Danbolt 1998).

Extrasynaptic Glutamate Receptors The G protein-linked mGluRs have a central role in regulating synaptic glutamate. mGluRs are expressed perisynaptically, and activation of mGluR2/3 receptors decreases presynaptic glutamate release (Schoepp 2001). Thus, activation of mGluRs may serve as a brake on glutamate spillover, preserving input specificity by diminishing synaptic glutamate levels (Huang and Bergles 2004). This mechanism has recently been exploited to develop a highly selective mGluR2 agonist that decreases glutamate release (Patil et al. 2007). This development suggests a role for the pharmacological modulation of glutamate spillover as a treatment for schizophrenia and other illnesses where psychosis is a central feature.

16.3 The Schizophrenia Syndrome

Schizophrenia is most properly considered to be a syndrome, as its specific causes are unknown, and the diagnosis is based on the presence of a heterogeneous collection of signs and symptoms. There is no blood test, specific genetic marker, or cognitive battery that can predict who will develop this illness. The schizophrenia phenotype is characterized by the presence of specific clinical findings, often divided into positive, negative, and cognitive symptoms (Buchanan et al. 2000). Positive symptoms include hallucinations, which are often auditory. Patients report that they hear voices that are clearly located outside of their heads, most often engaged in a running commentary on their thoughts and behaviors (Kay 1990; Badcock 2010). Other common positive symptoms include paranoid delusions and disorders of thought processes (Kay 1990; Badcock 2010). Much more debilitating are the negative symptoms, which are associated with the diminution of normal social behaviors, and include withdrawal, decreased speech, diminished eye contact, decreased or muted facial expression and vocal inflection, and diminished spontaneous movement (Buchanan et al. 2000; Fleischhacker 2000). Cognitive impairments in this illness include, but are not limited to, deficits in verbal fluency, executive function, and working memory (Rajji and Mulsant 2008; Szoke et al.

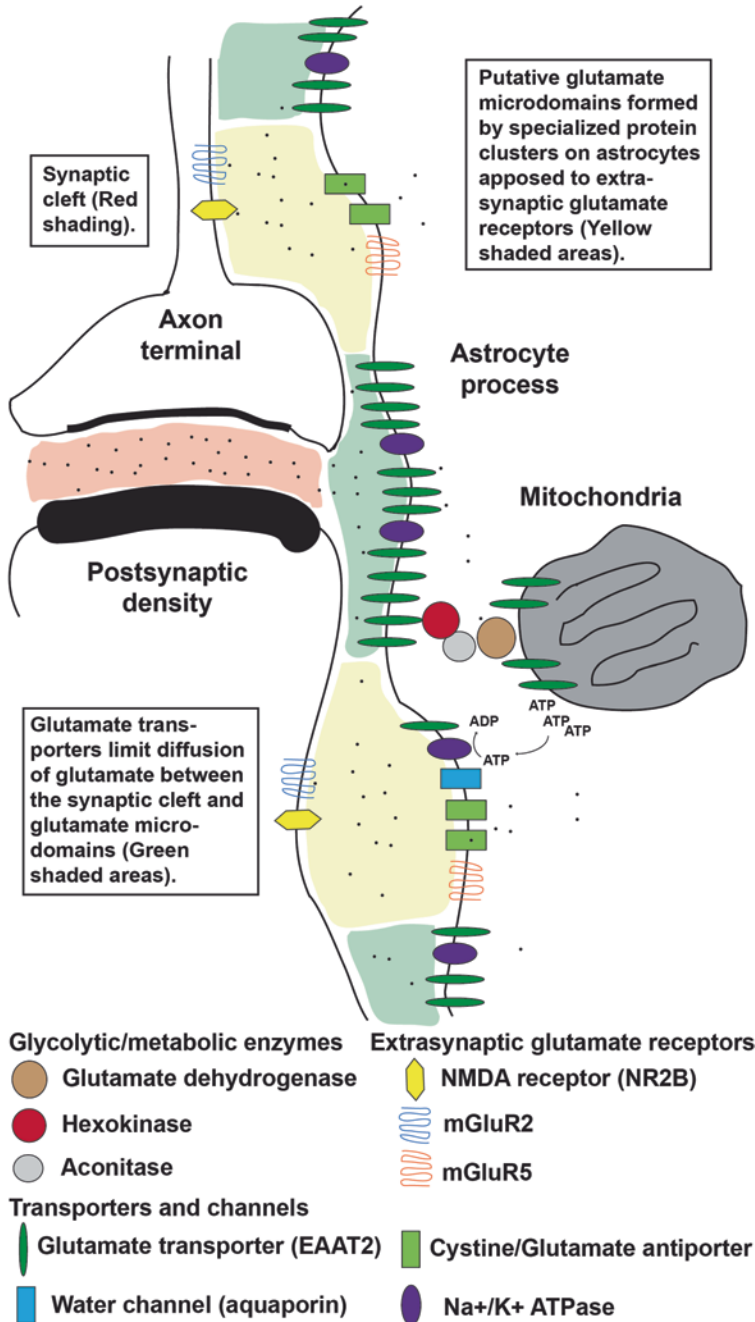


Fig. 16.1 Glutamate microdomains may be formed by specialized protein complexes found on plasma membranes in extrasynaptic regions where astrocytic and neuronal membranes are apposed to one another. Glutamate is released into the synapse (*red shading*) where it may bind and activate glutamate receptors. Plasma membrane glutamate transporters localized to perisynaptic astroglial processes bind and transport glutamate (*green shading*). Glutamate levels in extrasynaptic regions (*yellow shading*) may be regulated by glutamate spillover from synapses, release of glutamate from astrocytes, as well as reuptake of glutamate by glutamate transporters

2008; Wobrock et al. 2008; Potkin et al. 2009; Zanello et al. 2009). Few individuals suffering from schizophrenia have all of these symptoms, but the persistence of several characteristic symptoms, like auditory hallucinations, must be present in order for someone to be diagnosed with this disorder (Buchanan et al. 2000).

16.4 Prevailing Hypotheses of Schizophrenia

For decades, scientists have focused on the dopamine hypothesis of schizophrenia, which postulates that dysregulated dopaminergic neurotransmission is a key feature of the pathophysiology of the illness. The dopamine hypothesis is based on the observation that antipsychotics block D_2 receptors, and antipsychotic affinity for these receptors correlates with the ability to attenuate psychotic symptoms. Although numerous studies point to dopaminergic abnormalities in schizophrenia, dopamine dysfunction cannot completely account for all of the symptoms observed, since neuroleptics typically are effective only for the positive symptoms of the illness while negative symptoms and cognitive deficits are relatively refractory to treatment with typical antipsychotics (Joyce and Meador-Woodruff 1997; Laruelle et al. 1999). Consequently, alternative hypotheses that may help explain the pathophysiology of schizophrenia have been sought.

While the dopaminergic neurotransmitter system was implicated due to the effects of antipsychotic drugs, this system does, of course, not act in isolation. Dopamine receptors are found throughout the brain, where they modulate excitatory and inhibitory neurotransmission via G-protein signaling pathways. Blockade of dopamine receptors in corticolimbic circuits directly alters release of other neurotransmitters including glutamate and γ -aminobutyric acid (GABA). Not surprisingly, extensive postmortem studies have found changes in glutamatergic and GABAergic systems in this illness (Alda et al. 1996; Lewis et al. 2004; Deep-Soboslay et al. 2011; McCullumsmith and Meador-Woodruff 2011). However, evidence for involvement of glutamate receptor dysfunction, in particular the NMDA-subtype glutamate receptor, suggests a prominent role for glutamatergic abnormalities. NMDA receptor antagonists (but not GABA receptor antagonists) can induce both the positive and negative symptoms of schizophrenia, including cognitive deficits (Javitt and Zukin 1991; Tamminga 1999). Moreover, these compounds can exacerbate both positive and negative symptoms in schizophrenia (Lahti and Tamminga 1995). Chronic administration of phencyclidine (PCP)-like compounds induces a persistent psychotic symptomatology (Tsai and Coyle 2002) and reduces frontal lobe blood flow and glucose utilization, which is remarkably similar to the “hypofrontality” described in schizophrenia (Hertzmann et al. 1990).

Despite these observations, the complexity of schizophrenia is not readily explained by a static neurochemical model. The onset of schizophrenia is typically in late adolescence or early adulthood (Alda et al. 1996). The onset of positive and negative symptoms in a previously normally functioning person, coupled with a lifetime of waxing and waning symptoms, accompanied by the possibility of a steady

decline in social, occupational, and cognitive functioning, has led to longitudinal neurodevelopmental models that take into account genetic and environmental factors (Marenco and Weinberger 2000). Data supporting the neurodevelopmental hypothesis include studies suggesting that schizophrenia is associated with late winter births in urban environments, as well as a number of other prenatal, perinatal, and postnatal events (Marenco and Weinberger 2000; Lewis and Levitt 2002). Schizophrenia is perhaps best considered a disorder of neuroplasticity (McCullumsmith et al. 2004a). Plasticity refers to the ability of a system to affect reversible, long term changes in response to stimuli. Molecular correlates of learning and memory, including LTP and LTD, likely facilitate neuroplasticity in the brain. These processes are significantly impaired in severe mental illnesses, including schizophrenia (Malenka and Nicoll 1999; McCullumsmith et al. 2004b; Talbot et al. 2009). Glutamate transmission is a central component of LTP and LTD and, hence, has a central role in plasticity.

16.5 Glutamatergic Abnormalities in Schizophrenia

A number of studies have evaluated glutamate neurotransmission in schizophrenia using different approaches. In this section, we first discuss data from magnetic resonance spectroscopy (MRS), a technique which is largely focused on measuring glutamate, glutamine, and associated metabolites in specific brain regions of living patients. A strength of these studies is that the data are collected from afflicted individuals relatively close in time to the onset of the illness, while postmortem studies are examining brain tissues from more aged individuals who have had a lifetime of psychiatric illness. Next, we will briefly discuss data from postmortem studies, with a particular focus on extrasynaptic glutamate receptors and astroglial transporters.

Magnetic Resonance Spectroscopy Findings in Schizophrenia The balance of studies using MRS to examine glutamate, glutamine, N-acetylaspartate (NAA), and other metabolic intermediates have yielded mixed results (Bartha et al. 1997; Deicken et al. 1997; Theberge et al. 2002; Hutcherson et al. 2012; Kraguljac et al. 2012b). While a couple of studies have found changes in glutamate in schizophrenia, one large meta-analysis found decreases in the glutamate metabolite NAA in the basal ganglia and frontal lobe (Kraguljac et al. 2012a, b). Changes in NAA levels suggest abnormalities of glutamate synthesis and/or cycling in schizophrenia (Clark et al. 2006). A different meta-analysis found decreased glutamate and decreased glutamine in the medial frontal cortex in schizophrenia, suggesting that glutamate neurotransmission is diminished in this illness (Marsman et al. 2011). One interesting finding from these studies is the loss of correlation between NAA and glutamate levels in subjects with schizophrenia, compared to disease-free control subjects (Hutcherson et al. 2012; Kraguljac et al. 2012a). Taken together with the meta-analyses, these data suggest a significant abnormality in the glutamate/glutamine cycle in limbic circuits in schizophrenia. One limitation of the MRS approach

is that it measures all glutamate, glutamine or NAA without regard for it/them being intra or extracellular. For example, there may be a global increase in glutamine in the anterior cingulate cortex, with a large increase in intracellular pools and a small decrease in extracellular levels.

Abnormalities of Glutamatergic Enzymes in Schizophrenia There are several key enzymes involved in the glutamate/glutamine cycle as well as the synthesis or break-down of glutamate. Changes in enzymes levels may impact the amount of glutamate available for release from neurons and glial cells. Several studies have found decreased expression of carboxypeptidase II (binding and activity), glutaminase (mRNA), and glutamine synthetase (mRNA and protein) in limbic regions in schizophrenia (Burbaeva et al. 1999; Goff and Coyle 2001; Laruelle et al. 2003; Bruneau et al. 2005). Other studies have found increases in glutaminase (mRNA and activity) (Gluck et al. 2002; Bruneau et al. 2005). While these data support the hypothesis that glutamate synthesis and cycling may be impaired in schizophrenia, all of these studies were done at the regional level, and thus fail to capture the complexity of glutamate synapses at the cellular or subcellular level. For example, there may be diminished expression of glutamate enzymes in astrocytes, but increased expression in pyramidal neurons. Finally, one of the most interesting findings is a decrease in the dipeptidase glutamate carboxypeptidase II (GCP II; also known as NAALADase) activity in the frontal cortex and hippocampus in schizophrenia. GCPII catabolizes N-acetylaspartyl glutamate (NAAG) to glutamate and NAA (Ghose et al. 2004). These findings are consistent with the MRS data discussed above, which found decreased levels of NAA. NAAG antagonizes NMDA receptors, and increased levels (secondary to diminished GCP II activity) might contribute to NMDA receptor hypofunction. One strength of this study is that the authors measured enzyme activity, and not just transcript or protein levels, a technically demanding approach (Tsai and Coyle 2002).

Glutamate Receptor Abnormalities in Schizophrenia The observation that PCP may cause a schizophreniform psychosis in persons without a prior diagnosis of schizophrenia led to investigation of ionotropic glutamate receptor expression in schizophrenia. Initial hypotheses were focused on the idea that a loss or hypofunction of NMDA receptor activity would be reflected by diminished expression of NMDA receptor subunits as well as NMDA receptor binding sites. However, on balance, studies of NMDA receptor expression in the postmortem brain in schizophrenia have no clear or consistent pattern of findings (McCullumsmith et al. 2012). For example, there are over 18 studies of NMDA receptor subunit expression in the frontal cortex alone. Other than some changes in binding site expression, the hypothesis that there is deficient NMDA receptor expression stands largely unproven (McCullumsmith et al. 2012). Similar to NMDA receptors, AMPA and kainate receptor studies generally do not have a consistent pattern of abnormalities other than perhaps changes in AMPA receptor GluA2 subunit expression in the hippocampus (Tamminga 1999; Meador-Woodruff et al. 2001a; Harrison 2004). Interestingly, administration of PCP, which blocks the NMDA receptor channel, leads to increased glutamate release, which may lead to spillover of glutamate from

the synaptic cleft to extrasynaptic areas, activating extrasynaptic (non-NMDA) glutamate receptors. This mechanism may simulate/reflect the condition in schizophrenia, where impaired astrocyte function leads to diminished glutamate buffering and reuptake.

Metabotropic Receptor Expression in Schizophrenia While there are fewer post-mortem studies of mGluRs, compared to ionotropic receptors, the data are no less contradictory. For example, mGluR3 protein expression has been reported as increased, decreased and unchanged in the frontal cortex (Gupta et al. 2005; Corti et al. 2007; Ghose et al. 2009; Shan et al. 2012). Genetic linkage studies suggest that mGluR5 is involved in schizophrenia, and increased mGluR5 and mGluR1 transcript and mGluR1 protein expression have been found in prefrontal cortex in this illness (Ohnuma et al. 1998; Devon et al. 2001; Gupta et al. 2005; Volk et al. 2010).

Abnormalities of Glutamate Transporters in Schizophrenia Several studies have reported region-level changes in the expression of the glial glutamate transporters EAAT1 and EAAT2 in schizophrenia. EAAT1 protein expression was decreased and EAAT1 glycosylation altered in the dorsolateral prefrontal cortex (DLPFC) (Bauer et al. 2008, 2010). Decreased EAAT1 and EAAT2 protein was found in the superior temporal gyrus, while only EAAT2 protein was decreased in the hippocampus (Shan et al. 2013). In contrast to these protein studies, increased levels of EAAT1 mRNA were found in the anterior cingulate cortex and thalamus (Smith et al. 2001; Bauer et al. 2008; Rao et al. 2012), suggesting a compensatory response to diminished glutamate reuptake capacity. Alterations in EAAT2 mRNA have also been reported in the hippocampus (decreased) and neocortex (increased, decreased and unchanged) in schizophrenia (Ohnuma et al. 1998, 2000; Matute et al. 2005; Lauriat et al. 2006; Bauer et al. 2008; Rao et al. 2012).

The neuronal transporters have also been studied. We have previously reported increased expression of EAAT3 protein and mRNA in the anterior cingulate cortex, while other studies have measured EAAT3 mRNA expression in the frontal cortex (increased), DLPFC (no change) and striatum (decreased) (McCullumsmith and Meador-Woodruff 2002; Lauriat et al. 2006; Nudmamud-Thanoi et al. 2007; Bauer et al. 2008; Horiuchi et al. 2012; Rao et al. 2012). No changes in EAAT3 protein levels were detected in the superior temporal gyrus or hippocampus (Shan et al. 2013). These conflicting data for neuronal glutamate transporters mirror the findings of glutamate receptor subunit expression, and are limited by the likelihood that glutamate transporter expression changes may be cell-specific, and change in different directions in different populations of cells. These findings have contributed to reformulation of the glutamate hypothesis of schizophrenia, with the idea are changes in glutamate receptor and/or transporter expression in schizophrenia is not a problem of too much or too little protein expression, but a problem with protein trafficking or signaling. Localization and activity of astroglial-localized glutamate transporters is mediated, in part, by protein-protein interactions. Expression of some of these EAAT-interacting proteins has been assessed in severe mental illness.

EAAT-Interacting Proteins in Schizophrenia Several glutamate transporter-interacting molecules have been identified, including G-protein suppressor pathway 1 (GPS1), JWA, ARHGEF11, and KIAA0302 (also called beta III spectrin). These molecules can affect glutamate transport function through trafficking, anchoring, phosphorylation, glycosylation, and degradation of transporters in the brain. For example, GPS1 decreases EAAT2-mediated glutamate reuptake through a direct protein-protein interaction, and levels of GPS1 protein are elevated in the frontal cortex in schizophrenia. These data suggest that there may be normal levels, but decreased activity, of a specific transporter due to modulation of transporter function or localization to the plasma membrane (Bauer et al. 2008).

In summary, changes in glutamate neurotransmission in schizophrenia point towards complex abnormalities of protein localization and function, contributing to the molecular neuropathology that underlies the schizophrenia phenotype.

16.6 Glutamate Spillover in Schizophrenia

Glutamate Spillover in Schizophrenia As outlined above, several postmortem studies have found changes in EAAT expression in schizophrenia, as well as changes in the molecules that regulate EAAT localization and activity (Ohnuma et al. 1998, 2000; Smith et al. 2001; McCullumsmith and Meador-Woodruff 2002; Lauriat et al. 2006; Bauer et al. 2008). In general, these changes in gene expression are consistent with diminished regional expression of astroglial (but not neuronal) glutamate transporter expression and activity. Further, a recent genetic study analyzing copy number variants reported a subject with schizophrenia with a deletion of several EAAT1 exons (Cook and Scherer 2008). Finally, characterization of the complete GLAST (EAAT1) knockout found changes consistent with behavioral endophenotypes associated with schizophrenia, including locomotor hyperactivity and abnormal social behavior (Karlsson et al. 2008, 2009). Several of these abnormal behavioral findings were reversed by administration of antipsychotic medication or mGluR2/3 receptor agonist administration, which decreases presynaptic glutamate release. These data suggest that there are region-specific deficits in EAAT reuptake capacity in schizophrenia, which could lead to glutamate spillover.

Chronic Glutamate Spillover May Lead to Remodeling of Synapses In the prefrontal cortex (PFC) in schizophrenia, there are changes in the structure, composition, and numbers of excitatory synapses (Broadbelt et al. 2002; Lewis et al. 2003, 2008). Increased packing density, decreased numbers of dendritic spines and diminished expression of structural proteins suggest significant alterations of synapses in this region (Selemon et al. 1995; Rajkowska et al. 1998, 2002). Several reports have found specific alterations in layers III and IV of the PFC, including abnormalities of pyramidal cells and interneurons (Hashimoto et al. 2003; Dong et al. 2005; Huang and Akbarian 2007; Huang et al. 2007; Lewis et al. 2008). One well-replicated finding is a decrease in parvalbumin-positive interneurons in the middle cortical layers

(Lewis et al. 2001; Hashimoto et al. 2003). A lamina-specific deficit in inhibitory tone could lead to increased release of glutamate, which, combined with diminished reuptake capacity, could lead to increased glutamate spillover.

Accumulating evidence from postmortem gene expression studies suggests neurochemical alterations consistent with spillover. We have found increased mGluR2/3 protein in the PFC (Gupta et al. 2005), which may be interpreted as an attempt by synapses to decrease spillover by decreasing presynaptic release. Expression of the cystine/glutamate antiporter catalytic subunit (xCT) was also increased in the DLPFC (Baker et al. 2008). This molecule is expressed on glia and releases glutamate into the extrasynaptic space in exchange for uptake of cystine, which is required for glutathione synthesis. The effect of increased xCT protein expression on glutamate release is not known, because it is the activity, and not expression level, of this molecule that determines the rate of glutamate release. However, changes in the expression of xCT minimally suggest abnormalities of the regulation of extracellular glutamate levels (Baker et al. 2008). Finally, a number of studies have also described changes in ionotropic glutamate receptor binding site expression, suggesting a change in NMDA and AMPA receptor stoichiometry in the frontal cortex in schizophrenia (Akbarian et al. 1996; Healy et al. 1998; Ibrahim et al. 2000; Meador-Woodruff et al. 2001b; Beneyto and Meador-Woodruff 2006; Beneyto et al. 2007; Harney et al. 2008). Interestingly, preclinical studies have shown that glutamate spillover is associated with alterations in ionotropic receptor subunit composition and function (Lozovaya et al. 2004b; Harney et al. 2008).

We propose that there is remodeling of glutamate synapses in schizophrenia secondary to glutamate spillover (Fig. 16.2). Glutamate spillover may be secondary to increased release (in a misguided attempt to activate “sick” NMDA receptors), as well as deficits in glutamate reuptake capacity. In this setting, we would predict that perisynaptic localization of glutamate transporters on the plasma membrane of astrocytes is diminished, either as a primary deficit in transporter localization, as a compensation for increased extrasynaptic glutamate release, or both. Redistribution of glutamate transporters on astrocytes would lead or contribute to increased spillover, causing excitotoxicity and loss of input specificity. Further, we postulate that these deficits are initially relatively subtle but chronic in nature, leading to inappropriate remodeling of excitatory synapses, which do not function normally. This idea is supported by the phenotype of the GLAST/EAAT1 knockout mice, which have moderate cognitive and behavioral impairment, but no morbidity or mortality associated with seizures (Watase et al. 1998; Karlsson et al. 2008, 2009).

16.7 A Role for Glutamate Microdomains

Glutamate Microdomains Critical to cellular function of proteins is subcellular locality or microenvironment, in which proteins cluster and interact with numerous others. These biologically and morphologically discrete microdo-

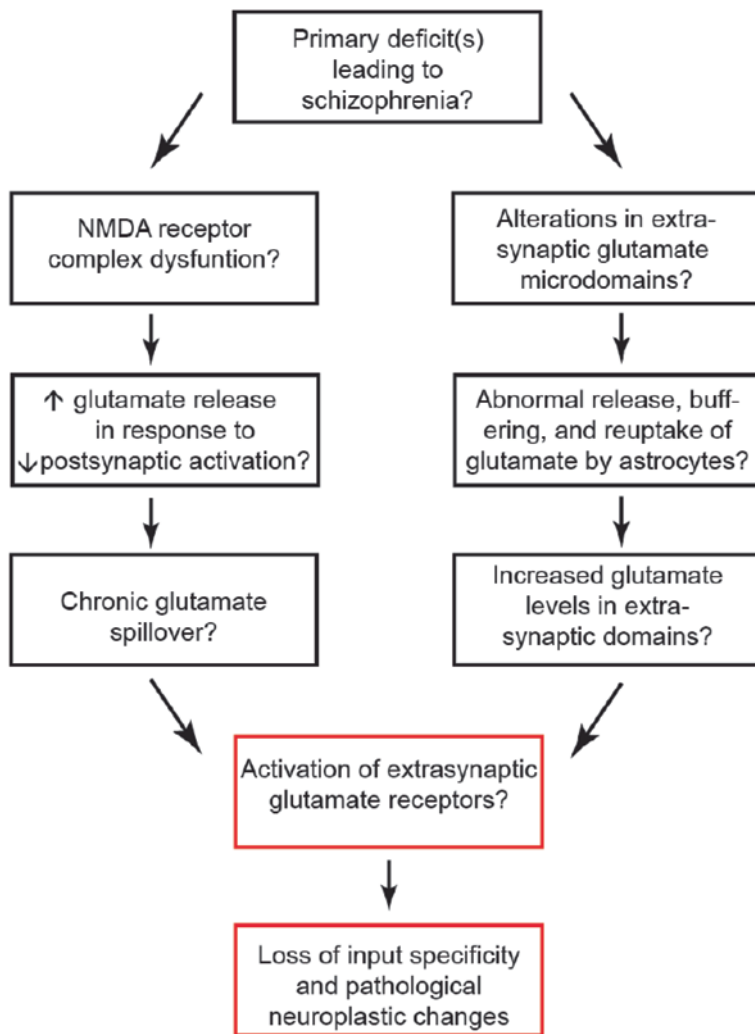


Fig. 16.2 There are likely a number of primary causes, which yield the constellations of symptoms found in schizophrenia. Proximate causes might include NMDA receptor dysfunction or abnormalities of glutamate microdomains. The *red boxes* may be a final common pathway that contributes to the phenotype of this syndrome

mains require tightly regulated trafficking of component proteins and thus organize the intermolecular environment for proteins and their interactions (Grosche et al. 1999; Hemler 2003; Füllekrug and Simons 2004; Tzingounis and Wadiche 2007; Farr et al. 2009; Newpher and Ehlers 2009). Evidence for cortical glutamate microdomains is based on several observations: (1) extracellular glutamate levels may vary between 0.2–7 μM in the extrasynaptic space in a region- and milieu-dependent manner; (2) a large body of work has described localization of

functional glutamate receptors outside of synapses; (3) the localization and buffering/transport properties of glutamate transporters strongly suggest partitioning of non-synaptic extracellular spaces; (4) several recent studies have characterized the functional coupling of protein complexes, structural proteins, organelles, and signaling pathways that converge on cellular processes involving glutamate; and finally, (5) glial cells may form electronically independent morphological structures that ensheath neuronal structures of unknown function (Grosche et al. 1999, 2002; Szabadkai and Rizzuto 2007; Spat et al. 2009; Genda et al. 2011). Taken together, these data suggest that glutamate microdomains are formed by specialized regions of astrocytic processes and neuronal membranes (Fig. 16.1) (Grosche et al. 1999; Genda et al. 2011). Glutamate levels within these domains may be determined by spillover from synapses or release from astrocytes (Fig. 16.1) (Lehre and Danbolt 1998).

The composition and function of specialized protein clusters in astrocytes has recently been investigated in rodent brain tissues. Using immunoisolation, one study found a complex comprised of GLT-1 (the rodent isoform of EAAT2), Na⁺/K⁺-ATPase, hexokinase, and mitochondria (Genda et al. 2011), while another found Na⁺/K⁺-ATPase, the water channel aquaporin 4, and mGluR5 (Fig. 16.2) (Illarionova et al. 2010). Other studies have found functional coupling of glutamate reuptake, cytosolic and mitochondrial sodium exchange, and glucose utilization in astrocytes (Hertz 2011; Skytt et al. 2012). Interacting partners and ultrastructural localization have not been determined for the cystine/glutamate antiporter.

Evidence for Abnormalities of Glutamate Microdomains in Schizophrenia: Structural Abnormalities In the frontal cortex in schizophrenia, there are changes in volume and cell density suggesting significant alterations in the spatial arrangement of synapses and microdomains in this illness (Selemon et al. 1995; Rajkowska et al. 1998, 2002; Broadbelt et al. 2002; Lewis et al. 2003; Lewis and Gonzalez-Burgos 2008). Specifically, there is thinning of cortical gray matter accompanied by decreased density of astrocytes as well as a loss of neuropil, while the balance of studies has typically found no changes in the number of neurons (Selemon et al. 1995; Rajkowska et al. 1998, 2002; Broadbelt et al. 2002; Lewis et al. 2003; Lewis and Gonzalez-Burgos 2008). These findings suggest a marked abnormality in the ultrastructural elements that account for the large volume of gray matter not occupied by cell bodies or synapses.

Abnormalities of Mitochondria A few studies have assessed the density of mitochondria in schizophrenia. One study found a decrease in the number of mitochondria per synapse in the striatum in treatment-responsive subjects with schizophrenia, while another found decreased volume fraction and area density of mitochondria in subjects with duration of disease more than 21 years (Uranova et al. 2001; Somerville et al. 2011, 2012). In addition, decreased expression of transcripts for a mitochondrial proton transporter were found in the frontal cortex in schizophrenia, and association of the glycolytic enzyme hexokinase 1 with mitochondria was decreased in the parietal cortex in this illness (Gigante et al. 2011; Regenold et al. 2012). These data suggest an abnormality of mitochondrial coupling in schizophrenia, which may

reflect an alteration in the association of mitochondria with proteins that comprise these domains.

Changes in Glutamate Receptor and Transporter Expression The changes in glutamate receptor and transporter expression detailed above are consistent with diminished glutamate reuptake capacity, indicating increased diffusion of glutamate between synapses and microdomains. In addition, changes in extrasynaptic expression of the cystine/glutamate antiporter (which releases glutamate from astrocytes) indicate an increased capacity for extrasynaptic release of glutamate (Kristiansen and Meador-Woodruff 2005; Baker et al. 2008). Both of these mechanisms could lead to increased activation of extrasynaptic glutamate receptors in extracellular glutamate microdomains.

Conclusions

We propose that there is increased activity of extrasynaptic glutamate receptors in schizophrenia secondary to increased levels of extrasynaptic glutamate (Fig. 16.2). Increased extrasynaptic glutamate may be due to increased presynaptic release and spillover (in a misguided attempt to activate sick NMDA receptors in the PSD), increased diffusion of glutamate out of the cleft secondary to deficits in glutamate buffering and reuptake capacity, and/or increased release of glutamate from astrocytes. In this setting, we would predict that integrity of glutamate microdomains is disrupted, either as a primary deficit in the assembly and localization of these domains, as a compensation for increased synaptic glutamate release and spillover, or both. Regardless of the mechanism, we hypothesize that the composition and localization of astrocytic proteins in glutamate microdomains are abnormal in schizophrenia, leading to pathological neuroplastic changes in the structure and function of glutamate circuits in schizophrenia.

Interestingly, several promising trials of various glutamate receptor modulators have failed to yield new pharmacological treatments for schizophrenia, including NMDA receptor glycine-site agonists (targeting NMDA receptors in the PSD), AMPA receptor modulators, called AMPAkinases (targeting AMPA receptors in the PSD), and more recently mGluR2 modulators (targeting presynaptic release of glutamate). The challenges of developing effective glutamatergic drugs may be harder to surmount because the pathophysiology of excitatory neurotransmission in schizophrenia has generally been viewed in a neuron-centric manner. Viewing abnormalities of glutamate transmission as a problem of astrocyte dysfunction may yield fresh insights for developing pharmacological targets to treat this devastating illness.

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