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Neville Vassallo *Editor*

Natural Compounds as Therapeutic Agents for Amyloidogenic Diseases

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Natural Compounds as Therapeutic Agents for Amyloidogenic Diseases

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Preface

A large group of human diseases are characterised by the deposition of insoluble proteinaceous lesions, termed amyloid. The underlying pathogenic process in these disorders involves the misfolding and self-assembly of native monomeric proteins into toxic, multimeric aggregates. Indeed, “protein misfolding” is one of the most researched topics in *cell biology* and *sub-cellular biochemistry* in recent years. To date, more than 20 amyloidogenic proteins have been identified, including amyloid-beta and tau in Alzheimer’s disease, alpha-synuclein in Parkinson’s disease, huntingtin in Huntington’s disease, superoxide dismutase in amyotrophic lateral sclerosis, prion protein in Creutzfeldt-Jakob disease and islet amyloid polypeptide in type 2 diabetes mellitus. Collectively, these disorders inflict an enormous economical burden on societies, whilst being devastating at a personal level. It is therefore of paramount importance to understand their origin and learn how to treat, or prevent, them. At present, there are no approved therapies that target amyloid formation directly, but a wealth of experimental and epidemiological evidence now indicates that various naturally derived compounds have beneficial anti-amyloid effects.

This book volume was written with the aim of disseminating the state-of-the-art and most recent knowledge of the field, highlighting the most promising nutra-therapeutics derived from natural compounds or extracts. The focus is mainly on Alzheimer’s disease and related dementias; however, Parkinson’s disease and other forms of cerebral amyloidosis are also discussed. Evidence is presented based upon published studies that have undergone rigorous scientific peer review. Researches include a combination of *in vitro* (biophysics, biochemistry, molecular and cellular biology), *in vivo* (fruit fly, transgenic mice and rats) and epidemiological (human) studies, so as to provide the reader with a holistic scientific approach. Among the therapeutic agents that are discussed in detail, we find Mediterranean diet and olive oil polyphenols, teas and catechins, natural phenolic compounds and bioflavonoids. Given the recent surge of public interest in herbal medicines, a chapter is devoted to evidence-based traditional Chinese medicine in brain ageing. Individual chapters are written by established scientists in their relevant fields; sufficient background is given in each chapter by way of introduction to the non-expert reader. Hence, this book should be of interest to a wide audience, including academia, various health professionals, the nutraceutical industry and undergraduate students reading for pharmaceutical, medical, biology and chemistry degrees.

I conclude by stating that, on a personal level, the most noble goal of this book is that with our collective contributions to scientific progress, we aim to provide hope for the future and hope for the many people suffering from as yet incurable amyloid diseases. This book is, after all, dedicated to them.

Msida, Malta

Neville Vassallo

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Olive Oil Phenols as Promising Multi-targeting Agents Against Alzheimer's Disease

1

Stefania Rigacci

Abstract

Amyloid diseases are characterized by the deposition of typically aggregated proteins/peptides in tissues, associated with degeneration and progressive functional impairment. Alzheimer's disease is one of the most studied neurodegenerative amyloid diseases and, in Western countries, a significant cause of dementia in the elderly. The so-called "Mediterranean diet" has been considered for long as the healthier dietary regimen, characterised by a great abundance in vegetables and fruits, extra virgin olive oil as the main source of fat, a moderate consumption of red wine and a reduced intake of proteins from red meat. Recent epidemiological studies support the efficacy of the Mediterranean diet not only against cardiovascular and cancer diseases (as previously demonstrated) but also against the cognitive decline associated with ageing, and several data are highlighting the role played by natural phenols, of which red wine and extra virgin olive oil are rich, in such context. In the meantime, studies conducted both *in vivo* and *in vitro* have started to reveal the great potential of the phenolic component of extra virgin olive oil (mainly oleuropein aglycone and oleocanthal) in counteracting amyloid aggregation and toxicity, with a particular emphasis on the pathways involved in the onset and progression of Alzheimer's disease: amyloid precursor protein processing, amyloid-beta (A β) peptide and tau aggregation, autophagy impairment, neuroinflammation. The aim of this review is to summarize the results of such research efforts,

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showing how the action of these phenols goes far beyond their renowned antioxidant activity and revealing their potential as multi-targeting agents against Alzheimer's disease.

Keywords

Alzheimer's disease • Mediterranean diet • Polyphenols • Extra virgin olive oil • Oleuropein aglycone

1.1 Introduction

Age-associated cognitive decline, in the relatively benign form of mild cognitive impairment (MCI) and in the more severe one of Alzheimer's disease (AD), has become a considerable social and clinical problem during the last decades, particularly in Western countries. With the important exception of familial AD, which is associated with one of several genetic mutations favouring the early onset of the disease, such a condition develops over a considerable period of time, maybe even decades. Moreover, in spite of the great efforts spent by researchers to identify early and accessible markers of this disease, AD is still diagnosed very late, when the neurological symptoms appear and the neuropathology is already in an advanced stage. For these reasons (life-long development – late diagnosis – neuronal loss) two important branches of the research in this field are early diagnosis and prevention.

With regards to prevention, a lot of attention has been placed on the role played by different lifestyles in favouring this disease. Epidemiological evidence points to a lower incidence of MCI and AD in populations adhering to the Mediterranean diet (MD), a dietary regimen that has already been strongly associated with a reduced risk for cardiovascular diseases and cancer (Martinez-Gonzalez et al. 2011; Sofi et al. 2010; Benetou et al. 2008; Lopez-Miranda et al. 2010). MD is characterized by extra virgin olive oil (EVOO), high intake of plant-based foods, relevant consumption of seafood, low-to moderate intake of dairy products, low intake of meat and a regular but moderate intake of red wine. In spite of regional and cultural variations (for example,

fish or cheese can be more-or-less abundant), one ingredient should never be absent from the Mediterranean table: EVOO, probably the most typical component of MD. EVOO consists of 98 % glycerides, mainly monounsaturated fatty acids (MUFA). In any case, it is the remaining 2 %, including various so-called “minor compounds”, that deserves much of our attention in this context. These consist of α -tocopherol and of several specific phenolic compounds including phenolic acids (caffeic, vanillic, syringic, *p*-coumaric, *o*-coumaric, protocatechuic, sinapic, *p*-hydroxybenzoic and gallic), phenolic alcohols (tyrosol and hydroxytyrosol), lignans (acetoxypinoresinol and pinoresinol), flavones (apigenin and luteolin) and, last but not least, secoiridoids (oleuropein aglycone, oleochantal and their derivatives). The latter are the most abundant and typical phenolic components of EVOO; hence investigations aimed at identifying the active principles responsible for the specific healthy benefits of EVOO have been mainly focused on them (Servili et al. 2009).

The concentration of these substances in oil is highly variable, depending on a number of different factors: olive cultivar, ripening stage at harvesting, geographic origin of olives, olive trees irrigation, operative conditions applied during crushing, malaxation and oil separation, oil storage modalities. The last two factors are particularly relevant. In fact, the highest phenolic content is present in oil immediately after its cold mechanical extraction from olives (a mandatory procedure to obtain an EVOO) and progressively declines with oil ageing, particularly if it is exposed to air and light that promote phenols oxidation and degradation. With some notable exceptions, total phenols in olive oil generally

range between 130 and 350 mg/kg, with EVOO at the highest end of this concentration interval.

Regarding the protection provided by EVOO consumption against age-associated cognitive decline, the results provided by the studies on the PREDIMED (PREvención con DIeta MEDiterránea) cohort are particularly interesting. The aim of these clinical trials was to test the efficacy of MD in counteracting cardiovascular disease events in asymptomatic people at high cardiovascular risk. In one of these studies, 578 subjects reasonably conforming to the traditional MD (except for a higher intake of meat and dairy products) were evaluated for their cognitive performance and total urinary polyphenol excretion (Valls-Pedret et al. 2012). Results showed that higher intake of both total olive oil and its virgin variety, coffee, walnuts, and wine were associated with better memory function and global cognition. Interestingly, the consumption of different foods and beverages was found significantly associated with the improvement in specific cognitive capabilities. Thus, EVOO and coffee were found to associate with better delayed verbal memory, walnuts with improved working memory, and red wine with higher Mini-Mental State Examination scores. Urinary phenols excretion was dose-dependently associated with improved memory, supporting the hypothesis that the common denominator of all those different foods, that is their high phenolic content, was the main factor responsible for the observed benefits.

The more recent PREDIMED-Navarra randomized trial included a cohort of 268 subjects (74.1 ± 5.7 years old, 44.8 % men with no cardiovascular disease but at high vascular risk because affected by type 2 diabetes mellitus or by three or more vascular risk factors) which were randomly assigned to three groups receiving for 6.5 years a low-fat diet (control), or a typical MD containing either EVOO (1 L/week) or mixed nuts (30 g/day) (Martinez-Lapiscina et al. 2013). The intervention with the EVOO-diet was associated with better cognitive performances, especially across fluency and memory tasks, and less MCI as compared to controls, while no significant differences were associated with the nuts-diet. Moreover, those who received the EVOO supplementation

had significantly better performances on both visual and verbal memory domains compared to those who received nuts supplementation. These results support an inverse association between the consumption of EVOO and amnesic cognitive impairment, the deficit most commonly associated with the risk for AD. These data further suggest that, in contrast to what was postulated to justify the reduced cardiovascular disease risk associated with MD, it was not the unsaturated lipid component of EVOO that was the major determinant factor for its beneficial effect in this context; indeed, nuts are a valuable source of polyunsaturated fatty acids too. These and previous data strongly indicate that EVOO phenolic content is the main ingredient responsible for the reduced risk for age-related cognitive impairment (Jacomelli et al. 2010).

But what exactly is the mechanism by which EVOO phenols exert their protection? These molecules have for long been considered just for their antioxidant activity (Tasset et al. 2011; Pierno et al. 2014; Farr et al. 2012) but accumulated data now suggest that other properties play an even more important role when we look at the maintenance of a good cognitive performance. For example, recent research conducted on C57Bl/6 J mice fed with an EVOO rich in phenols (10 % EVOO weight/weight of dry diet for a daily total polyphenol dose of 6 mg/kg), showed improved contextual memory and reduced age-related impairment in motor coordination with respect to controls. While the latter effect was associated with reduced lipid peroxidation in the cerebellum, the former did not correlate with oxidation or inflammation parameters, suggesting the involvement of other mechanisms not relying on increased antioxidant protection (Pitozzi et al. 2012). In support of this, are studies on the determination of the antioxidant activity of the main phenolic derivatives in human biological fluids after dietary EVOO intake, i.e. hydroxytyrosol (HT), tyrosol and homovanillyl alcohol glucuronide derivatives, at concentrations that covered all biologically relevant ranges (0.01–0.1 μM). The results of such assays show that glucuronidation rapidly decreases the protective activity of

EVOO phenols against Cu-mediated low-density lipoprotein (LDL) oxidation and in 2,2-diphenyl-1-picrylhydrazyl (DPPH) radical scavenging, confirming that the biological activity of a phenol-rich olive oil cannot merely be an antioxidant one (Khymenets et al. 2010).

Actually, the investigation on individual EVOO phenolic compounds has uncovered a lot of specific biochemical activities in different biological contexts. Here I will review the most relevant ones for what concerns AD counteraction, but first of all I will provide a brief description of this disease, particularly for those that are not completely familiar with its biochemical traits.

1.2 General Traits of Alzheimer's Disease

AD is characterised by a progressive decline of several cognitive domains including memory, visuospatial skills and executive function (Sa et al. 2012). The histopathological analysis of affected brains reveals selective neuronal degeneration and synaptic loss, particularly in the hippocampus, amygdala and temporal neocortex (Serrano-Pozo et al. 2011), accompanied by extracellular senile plaques and intraneuronal neurofibrillary tangles. The main constituent of plaques is a fibrillar network of polymeric unbranched fibrils made of A β peptides, fragments of different length; the most represented are A β (1–40), A β (1–42) and the highly aggregation prone N-terminally truncated A β (3–42) and A β (11–42) pyroglutamylated-species. A β fragments originate from the sequential activity of two proteases, γ -secretase and β -secretase (BACE) that act on the extracellular N-terminal domain of the transmembrane amyloid precursor protein (APP). These natively unfolded peptides aggregate into increasingly ordered structures (oligomers – protofibrils – fibrils) that become insoluble and resistant to the clearing activities of cells and tissues (Mohamed et al. 2011), thanks to the acquisition of a peculiar β -sheet rich fold called “amyloid structure” (Fig. 1.1) (Serpell et al. 2000).

Neurofibrillary tangles are amyloid in nature too, being mainly formed by the cytoskeletal protein tau, a member of the microtubule-associated protein family normally concurring to the assembly and stabilization of microtubules, physiologically involved in axonal transport and neurite outgrowth (Maccioni et al. 2001). Tau exists in six different isoforms (45–65 kDa) originating from alternative splicing; it is a hydrophilic cationic protein, unfolded under native conditions (Jeganathan et al. 2008), whose normal function is regulated by phosphorylation, glycosylation, ubiquitination, truncation, and nitration (Farias et al. 2011). In AD, abnormal phosphorylation occurs on specific tau residues (Ser202, Thr205, Ser235, and Ser404) (Alvarez et al. 1999). These post-translational modifications are catalyzed by two main protein kinases: the Cyclin-dependent kinase (Cdk)5/p35 system and glycogen synthase kinase-3 beta (GSK-3 β) (Farias et al. 2011), and are thought to promote tau aggregation.

More than two decades ago, Hardy and Higgins (1992) put forward the “amyloid cascade hypothesis” of AD, posing the insoluble A β fibrils as the primary toxic species. More recently, widespread support has been provided for the “toxic oligomer hypothesis” which proposes that pre-fibrillar intermediates are the toxic determinant of several amyloid diseases, including AD (Benilova et al. 2012; Brorsson et al. 2010; Walsh et al. 1999, 2002; Walsh and Selkoe 2004; Cleary et al. 2005; Billings et al. 2005; Koffie et al. 2009). Accordingly, such oligomers have been retrieved in diseased tissues both from animal models and humans, and *in vitro* assays have repeatedly confirmed their toxicity to cells. Several mechanisms determine oligomer cytotoxicity: membrane destabilization and derangement of ion homeostasis, particularly of Ca²⁺; oxidative damage; overload and dysfunction of the pathways devoted to protein quality control (proteasome, unfolded protein response, autophagy). With regards to tau, its oligomers are also now considered to be the most toxic species, possibly inducing neurodegeneration by affecting mitochondrial and synaptic function, both of which are early hallmarks of AD

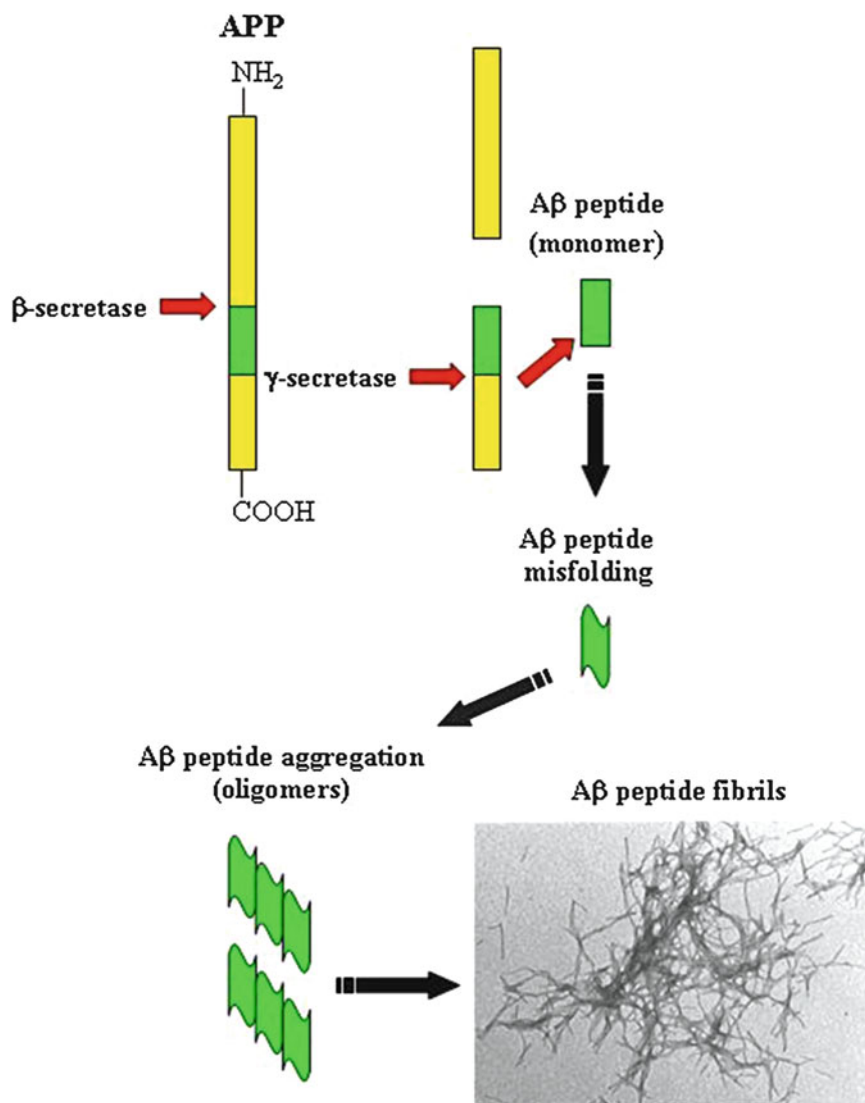


Fig. 1.1 Aβ peptide generation from APP by γ-secretase and β-secretase, followed by aggregation into oligomers and fibrils

and other tauopathies (Lasagna-Reeves et al. 2011; Guzman-Martinez et al. 2013).

Recently, the complexity of the amyloid scenario has been further increased by the discovery of different species of oligomers, varying in their structural features as well as in their toxicity (the so-called “amyloid polymorphism”) (Lee et al. 2007; Meinhardt et al. 2009; Stefani 2012). Moreover, a reappraisal of the role of fibrils in amyloid toxicity is taking place in recent years, particularly concerning AD, supporting a role for

Aβ fibrils not only as a possible reservoir of toxic oligomers but also as a neurotoxic species in their own right (Pan et al. 2011; Gilbert 2013).

As efficaciously summarized in a recent review, inflammation seems to play an important role in AD pathogenesis (Meraz-Rios et al. 2013). While initial activation of microglia and particularly astrocytes may be a positive response in the attempt to clear protein aggregates through phagocytosis and intracellular degradation, this process soon becomes dysfunctional leading to a

worsening of the pathology through the production of nitric oxide (NO), reactive oxygen species (ROS), pro-inflammatory cytokines (tumor necrosis factor- α , interleukin-1 β and interleukin-6), and prostaglandin-E2, which eventually promote neuronal death. Moreover, pro-inflammatory cytokines mediate an increase in tau phosphorylation, APP synthesis and A β generation through BACE-1 transcription (Lyman et al. 2014; Chen et al. 2012; Krstic et al. 2012).

This brief description of AD, which does not pretend to be exhaustive, can give an idea of the complexity of this pathology; accordingly, the search for possible therapeutic interventions must face such a highly multiform scenario. The investigation of EVOO phenols in the context of amyloid diseases has started relatively later with respect to other extensively-studied polyphenols like curcumin, epigallocatechin-gallate (EGCG) and resveratrol. Nevertheless, the results obtained with oleuropein aglycone (OLE) and oleocanthal (OLC) are promising. Here, I will critically review published reports, placing an emphasis on the multi-targeting potential of such compounds which, in my opinion, constitutes their main strength.

1.3 Oleuropein Aglycone

OLE or 3,4-dihydroxyphenylethanol elenolic acid (3,4-DHPEA-EA) (Fig. 1.2), is a secoiridoid derived from oleuropein by the activity of a β -glucosidase released from olive fruits during crushing (Brenes et al. 1999). Its content in EVOO is highly variable, depending on several factors: olive cultivar (Romani et al. 1999; Franconi et al. 2006), ripening (Brenes et al. 1999), method of oil production (Tripoli et al. 2005) and storage conditions, with oxygen and light being the most detrimental factors by promoting oxidation (Cicerale et al. 2009). Moreover, the method of extraction of OLE from oil samples and the procedure followed to determine its concentration seem to considerably affect the reported values. To give a general idea, I will refer in this review to the work of Servili *et al.* who analyzed 263 EVOO samples and found

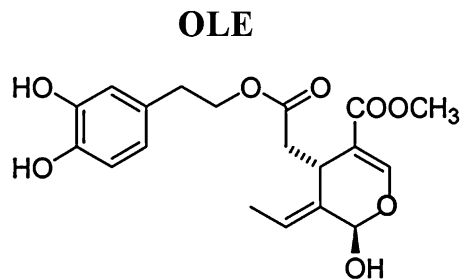


Fig. 1.2 Oleuropein aglycone (OLE)

a median value of 137 mg/kg for OLE and of 308 mg/kg for its dialdehydic form 3,4-DHPEA-EDA, also called Oleacein (Servili et al. 2009). Multiple beneficial effects were demonstrated for OLE, ranging from anti-atherogenic and anti-hypertensive to anti-cancer, anti-microbial and anti-inflammatory (Rigacci and Berti 2009) and in the last years promising anti-amyloid, neuroprotective and gerosuppressant activities have started to emerge for this secoiridoid.

Preliminary Mass Spectrometry (MS) analysis performed on its glycoside, oleuropein, revealed that it can associate the monomeric form of A β 1–40 and the oxidized form A β Met³⁵(O) with a 1:1–2:1 oleuropein:peptide stoichiometric ratio. Such interaction is non-covalent but possesses a remarkable binding energy, since the complexes are still observable when an orifice potential of 100 V is applied to the Electrospray ionization (ESI)-MS apparatus (Bazoti et al. 2006, 2008). Through enzymatic cleavage of the A β :oleuropein complex prior to ESI-MS analysis, three peptide regions were identified as being implicated in the interaction between the two and, of these, the hydrophobic (17–21) one was the best candidate for the association with the non-polar moiety of oleuropein and hence also with its aglycone derivative OLE (Galanakis et al. 2011). The importance of this peptide region for the interaction of A β with oleuropein has been further confirmed using NMR (Benaki et al. 2009; Kallberg et al. 2001). Interestingly, the A β sequence critical for peptide fibrillization overlaps the putative OLE-binding region (Kallberg et al. 2001; Tjernberg et al. 1996, 1999);

thus, an interference of such secoiridoid with peptide aggregation could somehow be anticipated.

Actually, most of the studies on oleuropein as an aggregation inhibitor were performed on its aglycone OLE, which was shown to interfere with both A β 42 and human islet amyloid polypeptide (hIAPP) aggregation *in vitro* (Rigacci et al. 2010, 2011). Through a combination of structural analysis by Thioflavin-T (ThT) and Anilinonaphthalene-8-sulfonate (ANS) binding, Circular Dichroism (CD) analysis, Atomic Force Microscopy (AFM), Transmission Electron Microscopy (TEM), and toxicity assays on cultured cells, it was demonstrated that OLE is capable of redirecting the aggregation pathway, thereby avoiding the formation of toxic oligomers and triggering peptide precipitation into amorphous aggregates from which non-harmful protofibrils eventually evolve. Concerning hIAPP, the amorphous aggregates that originate during the first phases of incubation in the presence of OLE are unable to interact with, and to damage, the cell membrane (Rigacci et al. 2010). Pre-existing A β fibrils can also be remodeled by the addition of OLE, with a reduction in the density of the fibrillar deposit and, most importantly, no release of toxic fragments (Rigacci et al. 2011).

Such preliminary *in vitro* results paved the way to experimentation on model organisms, which have corroborated the idea that OLE can be protective against amyloid aggregation and aggregate toxicity not only in cultured cells but also *in vivo*. In fact, administration of OLE to the CL2600 *Caenorhabditis elegans* (*C. elegans*) strain, constitutively expressing A β (3–42), resulted in significantly lower plaque deposition and toxic oligomer formation, with a reduction in the extent of worm paralysis and an increase in survival (Diomede et al. 2013). When A β (1–42) was aggregated in the presence of OLE and then injected in the *nucleus basalis magnocellularis* of adult male Wistar rats, it was not toxic to cholinergic neurons and did not raise an inflammatory reaction. This is in contrast to what happened when A β aggregates, grown under the same conditions but in the absence of OLE, were injected. The much lower amount of soluble

toxic oligomers (recognized by the conformation-specific A11 antibody) in the injected rat brain suggested that OLE stably hinders the formation/release of toxic species, also when the aggregated peptide is introduced in a complex tissue environment (Luccarini et al. 2014).

A significant step forward in the *in vivo* research on the anti-amyloid potential of OLE was made by using the TgCRND8 transgenic mouse model of AD, encoding a double-mutant form of APP and showing cognitive impairment and amyloid plaque deposition from the age of 3 months. A robust improvement in cognitive performance and a remarkable reduction in A β plaque number, size and compactness was evident in mice fed for 8 weeks with an OLE-supplemented diet (50 mg/kg of diet). Improvement occurred even when the treatment was started at 4 months, when amyloid deposits are already present, hence suggesting that OLE can not only prevent amyloid deposition but also disaggregate preformed plaques. In this model, other relevant biological effects of this secoiridoid were uncovered: microglia migration to the plaques for phagocytosis was increased, astrocyte reaction was reduced, hippocampal neurogenesis was stimulated and, most interestingly, an intense and functional (i.e. leading to substrates degradation following vacuoles fusion with lysosomes) autophagy induction was elicited. Indeed, data obtained with cultured N2a mouse neuroblastoma cells confirmed that OLE dose-dependently activates autophagy acting on the “classical” pathways which involves mTOR inhibition (Grossi et al. 2013, 2014).

Notably, OLE antioxidant activity does not seem to be relevant in these systems. In fact, OLE does not significantly reduce either lipid peroxidation in the cortex of TgCRND8 mice (Grossi et al. 2013), nor intracellular superoxide level in the transgenic CL2600 *C. elegans* strain (Diomede et al. 2013). Rather, OLE acts both as an inhibitor of amyloidogenic peptide aggregation (possibly through direct binding, as suggested by the previously cited MS and NMR studies) and as a signaling molecule promoting cellular protective responses like autophagy stimulation and inflammation reduction. The latter property sounds particularly attractive since

the pathogenetic role of neuroinflammation in AD seems relevant – perhaps, even more so than the role of amyloid toxicity (McGeer and McGeer 2013). Though this position is debatable, it is a fact that AD progression is accompanied by a widespread neuroinflammation. Accordingly, incidence-based, population-based and case-control studies in humans suggest that some non-steroidal anti-inflammatory drugs (NSAIDs) could provide a degree of protection against AD (Vlad et al. 2008; in t' Veld et al. 2001). In any event, the ineffectiveness of different NSAIDs in several clinical trials leads us to hypothesise that, maybe, they work as preventative agents only if their administration is started early on – at least 2 years before the clinical diagnosis of AD (McGeer and McGeer 2013). This, together with the adverse gastrointestinal and cardiovascular effects that were recorded in some clinical trials, strongly reduce the potential of NSAIDs as anti-AD drugs (Scharf et al. 1999; Group 2006). Also for these reasons, research on the anti-inflammatory properties of natural compounds is particularly active and has led to substantiate at molecular level the already documented efficacy of EVOO and of its phenolic component in this respect. For example, significant reductions in serum leukotriene B4 and thromboxane B2 concentrations at 2 and 6 h after consumption of EVOO, but not after consumption of either olive oil or corn oil (both poor in phenols) were observed (Bogani et al. 2007); a 3-month intervention with a MD characterised by EVOO as the main source of fats determined a decrease in the expression of inflammation related genes (INF- γ , Rho GTPase-activating protein-15 and interleukin-7 receptor) higher than that coming from the adoption of a MD containing an olive oil poor in phenols (Konstantinidou et al. 2010); 50 μ M OLE (a concentration that was supposed to be reached in plasma by individuals consuming an EVOO-rich diet) inhibited tumour necrosis factor- α (TNF- α) induced matrix metalloproteinase 9 (MMP-9) expression and secretion in THP-1 human monocytic leukaemia cells by impairing NF- κ B-mediated genes transcription (Dell'Agli et al. 2010); similar oleuropein and HT concentrations inhibited

PMA-stimulated COX-2 and MMP-9 expression and activities thus reducing inflammatory angiogenesis in cultured endothelial cells (Scoditti et al. 2012); in this case too, a reduction in the ROS-sensitive NF- κ B transactivation was observed. However, the opposite result was obtained in a different cell type (embryonic kidney HEK293 cells stably expressing APP) where oleuropein seemed to increase MMP-9 gelatinolytic activity in the culture medium (Kostomoiri et al. 2013). The authors observed that treatment with oleuropein modified APP processing, increasing the formation of the non-amyloidogenic and neuroprotective sAPP α fragment, and they attributed this result to an increase in the α -secretase activity of MMP-9. The discrepancy between the two studies could derive from the different conditions employed (different cell lines, PMA-stimulated or basal MMP-9 secretion) and the different approaches to MMP-9 evaluation (in the first case MMP-9 mRNA level, protein expression and enzymatic activity were evaluated and found to be decreased (Scoditti et al. 2012)) while in the second study only MMP-9 gelatinolytic activity was determined (Kostomoiri et al. 2013). OLE dose-dependently inhibited the production of the pro-inflammatory chemokine CCL2 by human endothelial cells too, thus reducing endothelium inflammation (Sindona et al. 2012), and attenuated the inflammatory response in a carrageenan-challenged mouse model of inflammation (Impellizzeri et al. 2011).

The potential of OLE in the context of AD is corroborated also by its ability to inhibit the fibrillization of tau (both wild-type and carrying the P301L mutation, a mutation that increases its aggregation propensity and leads to frontotemporal dementia and parkinsonism in carriers) leading to reduced deposition of fibrillar material in the form of short rods (TEM analysis) (Daccache et al. 2011). The IC₅₀ of OLE for the inhibition of P301L tau aggregation was 1.3 μ M, as determined by the Thioflavin-S (ThS) fluorescence assay; in contrast, HT was poorly effective, inducing the growth of fibrils that were less dense but structurally similar to those originated in the absence of any inhibitor, thus confirming the

superiority of the secoiridoid as an anti-amyloid agent. Data concerning the toxicity of tau aggregates obtained in the absence or in the presence of OLE are still lacking.

Olive oil phenols are renowned for their antioxidant activity. Nevertheless, the relevance of this property following ingestion is questionable because of phenols modification (this point will be more thoroughly discussed in Sect. 1.4). Nevertheless, an increase in cellular antioxidant defense can be observed following exposure to these molecules: a 6-month oleuropein administration to aged rats (50 mg/kg body weight/day) significantly increased superoxide dismutase, catalase and glutathione peroxidase activities in the brain, thus reducing the lipoperoxidative damage. Interestingly, a concomitant increase in dopaminergic neurons in *substantia nigra* was also observed. This could derive from the increase in enzymatic antioxidant defense, since these neurons are particularly sensitive to oxidative stress because of their abundance in iron ions and in free radicals, generated during dopamine metabolism (Sarbishegi et al. 2014).

With regards to the molecular mechanisms by which OLE produces its cellular effects, these are far from being clearly defined. A step forward in this direction was made by investigating the whole-genome transcription profile of human breast cancer cells exposed for 6 h to an EVOO extract containing 25 % OLE and 49 % its dialdehydic form 3,4-DHPEA-EDA (Mendez et al. 2013). This extract induced a marked increase in the expression of several genes; bioinformatic analysis of the global transcriptomic profiles revealed the induction of the endoplasmic reticulum (ER) stress chaperones and of the unfolded protein response (UPR), both relevant against the accumulation of protein aggregates. Age-related changes in cell size, morphology and senescence associated β -galactosidase (SA- β -gal) staining were significantly counteracted by the EVOO extract, and these effects correlated with the increase in the expression of the histone deacetylase Sirtuin1 and the activation of the AMP-activated protein kinase (AMPK). These two proteins regulate a multitude of signaling pathways in cells and one of them, significantly

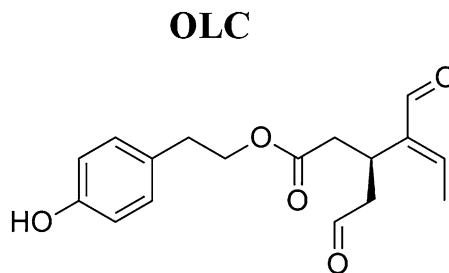


Fig. 1.3 Oleocanthal (OLC)

involved in the maintenance of cellular homeostasis, is autophagy. AMPK can activate autophagy by both positively regulating the Atg1/(ULK1) complex and inhibiting mTOR (Cai et al. 2012), while Sirtuin1 can deacetylate and activate the pro-autophagic Atg5, 7, 8 proteins and the LKB1 kinase, which in turn can activate AMPK (Chung et al. 2010). AMPK activity deregulation in AD seems to be involved in the perturbed brain energy metabolism, in A β generation and accumulation and in altered tau phosphorylation (Cai et al. 2012). Although the precise mechanism by which OLE activates autophagy still needs to be defined, these pieces of evidence reporting AMPK activation by a phenol-rich EVOO extract support the hypothesis of an involvement of this pathway.

1.4 Oleocanthal

OLC (3,4-HPEA-EDA) is the dialdehydic form of the decarboxymethyl derivative of ligstroside aglycone 3,4-hydroxyphenylethanol elenolic acid (3,4-HPEA-EA) (Servili et al. 1999) (Fig. 1.3). The median value of its concentration in EVOO, according to Servili et al. (2009), is 85 mg/kg depending on several variables (Brenes et al. 2000, 2001; Tovar et al. 2001), as already mentioned for OLE.

Regarding OLC effects on amyloid aggregation, currently available data support its ability to convert peptide monomers and oligomers (but not fibrils) into high-molecular-weight (HMW) species. Regarding A β (1–42), the latter's immunoreactivity to the conformation-specific an-

tibody NU1 (recognizing oligomers) as well as to the sequence-specific M89, 4G8, and 6E10 antibodies, increased when it was incubated in aggregation conditions together with OLC. This suggests the adoption of a different conformation, while no effect was evident when OLC was added to pre-formed fibrils (Pitt et al. 2009). Such an outcome could be beneficial only if the newly-formed HMW species lost the potential to induce cytotoxic effects. The ability of OLC-modified aggregates to bind the synapses of hippocampal neurons was not significantly altered (a reduction was observed, but it was not statistically relevant); however, unfortunately, cytotoxicity of the OLC-modified aggregates was not assessed by the authors. For such reasons, these data do not, as yet, convincingly support a beneficial effect of OLC in redirecting A β aggregation. However, something different occurs if neurons are pre-treated with OLC before coming in contact with toxic A β oligomers: in such situation, a reduced binding of oligomers to the cell membrane and a protection against synaptic deterioration was observed (Pitt et al. 2009). Moreover OLC, once present in the culture medium, increases the immunoreactivity of subsequently added toxic oligomers, making them more prone to clearance by NU1 antibodies.

Collectively, these data suggest a complex interaction of OLC with amyloid peptides, with structural remodeling (and subsequently altered immunoreactivity) varying depending on the preliminary aggregation state of the amyloid species. Such preliminary pieces of evidence lead us to hypothesize that OLC could work as a preventative agent since its presence in the cellular environment prior to the appearance of amyloid deposits seems particularly efficient in inhibiting the toxic outcome. An *in vivo* experimentation is required to assess OLC actual potency in reducing toxic amyloid deposition in the brain.

Conversion of monomers and oligomers into non-fibrillar HMW aggregates was also shown for tau in the presence of OLC (Li et al. 2009). An interesting comparison of the different chemical behavior of OLE and OLC as tau aggregation inhibitors was made by Daccache et al. (2011) also on the basis of other data (Li et al. 2009;

Monti et al. 2011): a structure–activity relationship study based on a series of derivatives of OLC pointed to an anti-fibrillization pharmacophore comprising both the saturated and unsaturated aldehyde moieties. Such a dialdehyde was proposed to cross-link two lysine residues within the third repeat of tau that would initiate the fibrillization process, rapidly followed by the modification of a single lysine, thus producing a cyclic adduct that evolves towards a more stable pyridinium-like complex by rearrangement of the skeleton. This would lock tau into a random coil conformation, preventing the transition to the β -sheet rich amyloid form and thus favoring the formation of non-amyloid HMW aggregates (Li et al. 2009; Monti et al. 2011). On the contrary, the methoxycarbonyl group of OLE increases the acidity of the hydrogen on the adjacent carbon thus inducing an intramolecular rearrangement leading to the dihydropyran form as the main isomer of OLE in solution. This form is in equilibrium with a monoaldehydic one that, when reacting with a lysine residue, would yield an aliphatic Schiff base that is unstable in aqueous media. OLE, therefore, does not work as a stable crosslinker and it does not promote the precipitation of tau into amorphous macro-aggregates; rather it favours the maintenance of a soluble form of tau so that the amount of fibrillar species is highly reduced (Daccache et al. 2011).

Although OLC does not seem to significantly affect the ability of tau to promote microtubule assembly in an *in vitro* tubulin polymerization assay (Li et al. 2009), some concerns still remain about retention of this important tau physiological function following OLC crosslinking, and an *in vivo* confirm is warranted.

Apart from its interference during amyloid aggregation, OLC usefulness against AD was recently supported by additional findings. A study conducted on cultured bEnd3 mice brain endothelial cells and C57BL/6 mice suggests that intraperitoneal administration of OLC to mice can increase the clearance of ^{125}I -A β (1–40) from the brain by up-regulating both A β degrading enzymes (Nephrilysin and Insulin degrading enzyme) and A β transporters (P-glycoprotein and LDL receptor related protein-1) at the Blood

Brain Barrier (BBB) (Abuznait et al. 2013). These effects are highly relevant because the expression of such transporters declines with ageing and in AD, possibly contributing to amyloid accumulation in the brain parenchyma (Silverberg et al. 2010; Vogelgesang et al. 2004). OLC was also found to activate AMPK (Khanal et al. 2011). As already mentioned in the previous section, this kinase not only represents a node in the complex network regulating energy supply to the cell, but also participates in the quality control of proteins and organelles *via* autophagy. Autophagy is particularly beneficial and overexploited during neurodegenerative diseases, where protein amyloid deposition eventually overwhelms the cellular buffering capacity. The possibility that OLC activates autophagy, and the relevance of this in the context of AD models, has not been directly investigated yet. Last but not least, OLC possesses a striking ibuprofen-like activity: it dose-dependently inhibits COX-1 and COX-2 but has no effect on lipoxygenase *in vitro* (Beauchamp et al. 2005), and it decreases lipopolysaccharide-induced nitric oxide synthase in chondrocytes (Iacono et al. 2010). This strong anti-inflammatory activity is probably, at the moment, one of the most appealing properties of OLC, potentially useful also in counteracting AD-associated neuroinflammation.

1.5 Bioavailability

Bioavailability of phenolic compounds represents a critical issue that has to be addressed by answering three fundamental questions: (1) to what extent are these compounds absorbed following ingestion? (2) How are they modified following ingestion? (3) How do they (or their modification products) distribute in tissues? Unfortunately these topics have been to date investigated with very different and non-standardized approaches, as it has been outlined in a recent review (D'Archivio et al. 2010), so that it is not easy to derive conclusive answers.

Concerning the phenolic compounds OLE and OLC, the main variables are represented

by the form in which they are ingested (pure compounds, extracts containing different percentages of the compounds, whole EVOOs with different phenolic composition), dosing, duration of the treatment, and association with different foods. Moreover, the great majority of the studies have been performed employing oleuropein, which can be extracted from olive leafs or olive mill wastewater and so is more convenient (and commercial) than OLE, which is enriched in EVOO.

In spite of all these variables, a general consensus has emerged regarding the fact that EVOO phenols are in fact absorbed and metabolized by humans, because their degradation and modification products are retrieved in urine following ingestion (Vissers et al. 2002; Weinbrenner et al. 2004; Miro-Casas et al. 2003). Nevertheless, absorption profiles vary depending on the source of such phenols: when they are ingested as an oleuropein-rich olive leaf extract, mainly sulfate and glucuronide derivatives are detected both in plasma and in urine, with HT glucuronide being the most abundant one, hence suggesting extensive degradation of the secoiridoid component (Garcia-Villalba et al. 2014; de Bock et al. 2013). On the other hand, when phenols are introduced with EVOO, OLE, ligstroside aglycone and their Phase II metabolites are mostly abundant in urine (Garcia-Villalba et al. 2010; Suarez et al. 2010). These results support the view that secoiridoids aglycones are better absorbed than their glycosylated counterparts and suggest that EVOO matrix can contribute to phenols stability in the gastrointestinal tract and favour their absorption. Indeed, when EVOO is mixed with acidified water (pH 2.0) in a 1:1 ratio at 37 °C (a condition simulating the stomach environment), the secoiridoid aglycones present remain stable for 4 h (Romero et al. 2007). Under these conditions, 50 % of total phenols diffuse from EVOO into the simulated gastric juice; this percentage increases as the pH rises, whilst the EVOO:water ratio decreases to 1:2 (the latter two conditions mimic those present in the duodenum).

Oleuropein is far more stable, both in the stomach and in the intestine (over 12 h), if it is

ingested with a meal (Markopoulos et al. 2009). Conversely, oleuropein is strongly hydrolyzed in the upper gastrointestinal tract, with the rapid appearance of HT glucuronide in the plasma, when it is taken as an olive leaf extract under fasting conditions (Garcia-Villalba et al. 2014). Moreover, OLE is absorbed more efficiently than oleuropein, probably because its greater apolarity favours passive transport across the cell membrane. In fact, when rabbits are fed an EVOO containing a 9-fold concentration of OLE vs oleuropein, a 60-fold concentration of OLE vs oleuropein is reached in the plasma (Coni et al. 2000). A recent report further shows that, after oleuropein administration to rats, OLE is retrieved both in faeces and urine (together with hydrolysis and modification products) (Lin et al. 2013). Particularly relevant in the context of AD is experimental evidence suggesting that, in rat and humans, orally-administered olive oil phenols, including OLE, oleuropein and/or one of its derivatives arising from tissue metabolism, cross the BBB and are found inside brain parenchyma (Serra et al. 2012; Vissers et al. 2002). Finally, OLE and 3,4-DHPEA-EDA seem to associate to membranes as a consequence of their hydrophobicity (Paiva-Martins et al. 2010); this implies that they may accumulate at the cellular level, reaching a local concentration higher than that expected on the basis of their plasma concentration alone.

Considering that HT and tyrosol are the main degradation products of OLE and OLC, respectively, either in oil (increasing with EVOO ageing) or in the organism following ingestion, a few words should be spent on these phenols too. HT penetration into the brain has been demonstrated in a pharmacokinetic study assessing the metabolic fate of intravenously injected [^{14}C]HT (D'Angelo et al. 2001). Anti-atherogenic, anti-inflammatory, anti-microbial, anti-proliferative and pro-apoptotic effects have been attributed to HT (Granados-Principal et al. 2010). The antioxidant potency of HT *in vitro* is very high but its biological importance as a ROS scavenger has been questioned of late because this seems to be greatly affected by HT modification (i.e. glucuronidation)

following ingestion (Khymenets et al. 2010). Nonetheless, when mice are fed for 12 days with 100 mg of HT/kg of body weight, basal and Fe^{2+} -induced malondialdehyde (MDA) formation are significantly reduced in explanted brain cells, in spite of no systemic increase in antioxidant capacity. This suggests that, besides direct ROS and iron-scavenging activity of HT, cytoprotection would derive also from different mechanisms (Scharf et al. 1999). Indeed, several experiments have shown that HT increases the expression of antioxidant enzymes, an ability potentially relevant in the context of AD. This happens as a result of a hormetic mechanism by which HT, behaving like a mild pro-oxidant, induces the activation of cellular defenses: in fact, in the presence of peroxidases, HT can undergo a catechol-semiquinone-quinone redox cycling generating superoxide, which in turn increases MnSOD expression. As a result, age-associated mitochondrial ROS accumulation is counteracted and the chronological lifespan of normal human fibroblasts is extended (Sarsour et al. 2012). Hormesis seems to be implied also in the increase in lifespan and stress resistance of tyrosol-fed *C.elegans*, subsequent to the activation of components of the heat shock response and the insulin pathway (Canuelo et al. 2012). The up-regulation of several antioxidant enzymes was also observed in keratinocytes exposed to HT: glutaredoxin, thioredoxin reductase, glutathione peroxidase-3, heme oxygenase-1, biliverdin reductase and ferritin, the latter three participating in the degradation of heme to biliverdin (a potent antioxidant) and in storage of the pro-oxidant free iron which is produced as a heme degradation by-product (Rafehi et al. 2012). Interestingly, in vascular endothelial cells the HT-mediated up-regulation of catalase expression and the associated protection against ROS increase are dependent on the activation of the AMPK-FOXO3a pathway, since they are abrogated when AMPK expression is suppressed by siRNA (Zrelli et al. 2011). On the basis of such HT-mediated AMPK activation we could speculate whether HT is an autophagy activator, too. However, the available data seem suggest the opposite: exposure of prostate cancer PC-

3 cells to 80 μM HT resulted in a significant increase in superoxide production accompanied by a defect in autophagy (Luo et al. 2013), while HT supplementation to rats counteracted autophagy activation induced by intense physical exercise (Feng et al. 2011). In a context closer to AD, HT was found to reduce the cytotoxicity of A β (25–35) to N2a neuroblastoma cells by decreasing NF- κ B nuclear translocation and cell death, but no effect was observed on A β – or H $_2$ O $_2$ -induced decrease in cellular glutathione (GSH) (St-Laurent-Thibault et al. 2011). Limited data are available concerning HT activity as an amyloid aggregation inhibitor: as already mentioned, the *in vitro* aggregation of tau was affected by HT (as demonstrated by the reduction in ThS binding) but amyloid fibrils structurally similar to those grown in the absence of the phenol were eventually formed (Daccache et al. 2011).

From these data the superiority of OLC and OLE, with respect to HT, as potential drugs against amyloid diseases seems evident. Accordingly, studies are currently under way to improve the bioavailability and preserve the integrity of such secoiridoids following ingestion. A promising approach is represented by encapsulation: simple olive oil phenols (tyrosol, HT, homovanillic acid, 3,4-dihydroxyphenylacetic acid, and protocatechuic acid) have been successfully enclosed into the hydrophobic cavity of β -cyclodextrin (Rescifina et al. 2010). Oleuropein, too, was found to enter the β -cyclodextrin particle with its phenolic portion, and this seemed to protect it from oxidation and to increase its aqueous solubility (Mourtzinos et al. 2007). Further suggestions come from trials performed with other polyphenols: for instance, promising results were obtained when curcumin was encapsulated in polyethylene glycol (PEG) nanoparticles stabilised with β -cyclodextrin. Such nanoparticles (<80 nm in size) did not tend to aggregate, were fairly stable and performed well both in an *in vitro* cell monolayer permeability assay (mimicking BBB transit) and after oral administration to TgCNRD8 mice, allowing an increase in curcumin penetration into the brain (Cheng et al. 2013).

1.6 Conclusions and Perspectives

The beneficial effects of MD in counteracting human diseases and aging are widely recognised, and multiple evidence points to EVOO as one of the most valuable ingredients of such a dietary regimen. The main differences in the composition of EVOO with respect to other edible oils are the prevalence of oleic acid as the main lipid (up to 85 % of total fatty acids, as compared with 14 to 59 % in sunflower, soybean and peanut oils) and the minor unsaponifiable fraction (0.5–2 %) containing squalene, tocopherol, various sterols and some peculiar phenols which contribute significantly to the unique properties of EVOO, not least oxidative stability and distinguished flavour. Though oleic acid has been endorsed with multiple beneficial activities following its enrichment in the membrane lipid bilayer, as summarised in a recent review (Lopez et al. 2014), it is the minor phenolic constituents which have generated increased attention, and are now considered as predominantly responsible for most of the health-promoting properties of EVOO. Accordingly, the accumulating data on OLC and OLE reviewed in this paper collectively support the hypothesis that such phenols are capable of eliciting multiple biochemical and biological responses, that might be harnessed in counteracting several age-associated diseases and, particularly, cognitive impairment and AD (Table 1.1).

To date, the activities of EVOO phenols potentially useful against AD can be grouped into several main categories: (i) inhibition of the formation of amyloidogenic A β fragments from APP, (ii) promotion of A β peptide clearance, (iii) inhibition of A β and tau toxic amyloid aggregation, (iv) reduction of inflammation, (v) autophagy activation, (vi) neurogenesis stimulation and (vii) antioxidant defence activation. OLE has until now given more unequivocal positive results than OLC with regards to the inhibition of toxic amyloid aggregation and the stimulation of cellular protective mechanisms like autophagy. Moreover, the specificity and selectivity of OLC crosslinking activity still

Table 1.1 Summary of OLE and OLC activities useful in counteracting AD

	Activity	References
OLE and oleuropein	Inhibition of the formation of A β toxic oligomers and plaques and remodelling of pre-existing A β deposits <i>in vitro</i> and <i>in vivo</i>	Rigacci et al. (2011), Diomedea et al. (2013), Luccarini et al. (2014), and Grossi et al. (2013)
	Improvement in survival, cognitive function and motor performance in AD transgenic models (<i>C. elegans</i> , mouse)	Diomedea et al. (2013) and Grossi et al. (2013, 2014)
	Inflammation reduction	Luccarini et al. (2014), Grossi et al. (2013, 2014), Dell'Agli et al. (2010), Scoditti et al. (2012), Sindona et al. (2012), and Impellizzeri et al. (2011)
	Autophagy activation	Grossi et al. (2013, 2014)
	Stimulation of hippocampal neurogenesis and increase in dopaminergic neurons	Grossi et al. (2014) and Sarbishegi et al. (2014)
	Inhibition of tau aggregation	Daccache et al. (2011)
	Promotion of sAPP α fragment production	Kostomoiri et al. (2013)
	Antioxidant enzymes activation	Sarbishegi et al. (2014)
OLC	Reduction of A β oligomers binding to the membrane, protection against synaptic deterioration	Pitt et al. (2009)
	Conversion of tau monomers and oligomers into non-fibrillar aggregates	Li et al. (2009)
	Up-regulation of A β -degrading enzymes and A β transporters at the BBB	Abuznait et al. (2013)
	AMPK activation	Khanal et al. (2011)
	Inflammation reduction	Beauchamp et al. (2005) and Iacono et al. (2010)

needs to be confirmed *in vivo*, to exclude negative side effects on functional proteins. Nevertheless, the potent anti-inflammatory activity of OLC and its ability to promote A β degradation and clearance are valuable properties.

The molecular mechanism/s by which OLE and OLC evoke those multiple cellular responses must still be clarified. Do they enter cells and directly bind to target molecules or do they mainly interact with membranes thereby inducing a signalling cascade? Data obtained with unilamellar phosphatidylcholine vesicles using fluorescence anisotropy of probes and fluorescence quenching studies led to conclusions that both OLE and OLC remained at the surface of the lipid bilayer (Paiva-Martins et al. 2003). Yet experimental evidence provided more recently by the same authors using erythrocytes suggested that OLE not only interacts with the cellular membrane but may also penetrate into the bilayer to

reach the radicals formed *in situ* (Paiva-Martins et al. 2010). One explanation for this discrepancy could be that biological membranes present microdomains of peculiar lipid composition that may facilitate phenols uptake. Accordingly, OLE (as well as oleuropein and other phenols and derivatives) was recently found inside the cytoplasm of human breast carcinoma cells exposed to an olive leaf extract (Quirantes-Pine et al. 2013) and of colon cancer cells exposed to EVOO extracts (Fernandez-Arroyo et al. 2012). These results leave the possibility of a direct intracellular signalling activity by EVOO phenols open to discussion.

Finally, in spite of the documented low bioavailability of these compounds, the positive results already obtained following *in vivo* administration confirm their efficacy, at least in model organisms. It is hoped that OLE and OLC will soon enter clinical trials to

rigorously assess their efficacy against AD during the different phases of disease progression, especially preclinical stages. Investigations employing different encapsulation strategies aimed at increasing the stability, bioavailability and brain targeting of these phenols are very promising and critical in this phase of the research.

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Marlene Jimenez-Del-Rio and Carlos Velez-Pardo

Abstract

Alzheimer's disease (AD) is an insidious neurological disorder that affects memory, one of the human brain's main cognitive functions. Around 5.2 million Americans currently have AD, and the number threatens to climb to 7 million by 2020. Our native country, Colombia, is no exception with an estimated 260,000 individuals to be affected by AD in 2020. A large, genetically-isolated community in Antioquia, Colombia, with early-onset familial Alzheimer's disease due to a presenilin-1 mutation is ideally suited for the study of molecular mechanisms of AD, and hence accelerate the discovery of new or alternative treatment approaches. In this regard, polyphenols – also known as polyhydroxyphenols – have shown antioxidant activity, gene regulation, metal chelator and anti-amyloidogenic aggregation effects. However, further *in vitro* and *in vivo* investigations are warranted to validate their use in clinical trials. *Drosophila melanogaster* is increasingly being used as a valid *in vivo* model of AD. Here, we summarise data published within the past 16 years (1998–2014) on the molecular biology of AD and the use of polyphenols in the fly to understand the molecular actions and feasibility of these compounds in the treatment of AD.

Keywords

Alzheimer's disease • Amyloid-beta • *Drosophila melanogaster* • Lymphocytes • Oxidative stress • Polyphenols

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2.1 Global Alzheimer's Madness: Is This the End of the Road?

In 1906, Dr. Alois Alzheimer reported the clinical symptoms and neuropathology findings of a “peculiar disease” from his patient Augusta

Deter (Stelzmann et al. 1995). The patient, a woman of 51 years, was completely disoriented in time and space and was incapable of understanding anything or of recognizing her surroundings. Most importantly, she lost her capacity to memorize. At autopsy, the brain was atrophic. After histological examination with Bielschowsky's silver stain, Alzheimer observed fibrils aggregated into dense bundles and miliary foci, which were caused by deposition of an unusual "substance" in the cerebral cortex (Alzheimer (Munich): About a peculiar disease of the cerebral cortex 2000). Although his report ignited no discussion at the time, it has fascinated the medical and research community for more than 100 years. Alzheimer's patient not only presented cognitive and language deficits but also suffered from non-cognitive features such as hallucinations, delusions, paranoia and aggressive behavior, typical of psychiatric symptoms; and presented two prominent post-mortem brain features: the dense bundles of fibrils and deposits in the cerebral cortex, later recognized as neurofibrillary tangles (NFT) and senile plaques (SP), respectively (Graeber et al. 1998). These features are the hallmarks of what is presently known as Alzheimer's disease (AD, OMIM entry 104300), and have changed little since then (Esiri 2001). Further biochemical studies have shown that NFT represent the intracellular accumulation of hyperphosphorylated tau (τ) protein (Iqbal et al. 1993; Grundke-Iqbal et al. 1986), and SP are extracellular deposits due to aggregation of the hydrophobic amyloid- β ($A\beta$) peptide (Glennner and Wong 1984; Masters et al. 1985). The disorder was named after Alzheimer by Emil Krapelin in 1910 (Schottyky 1932), not without controversy however (Cipriani et al. 2011; Ramirez-Bermudez 2012). Alzheimer's disease has become the most frequent neurodegenerative disease worldwide. Indeed, an estimated 5.2 million Americans have AD (Fargo and Bleiler 2014) with estimated healthcare costs spiraling to \$172 billion per year (Thies et al. 2013). By 2050, the total number of people with AD dementia is projected to be 13.8 million, with 7.0 million aged 85 years or older (Hebert et al. 2013). A similar trend is observed all over the world (Chan et al. 2013).

It was estimated that 35.6 million people lived with dementia worldwide in 2010, with numbers expected to almost double every 20 years, to 65.7 million in 2030 and 115.4 million in 2050 (Prince et al. 2013). Colombia is no exception. By 2020, it is estimated that about 260,000 and 1,500 individuals will be affected by sporadic and familial AD in Colombia, respectively (Unicesi 2013).

Most disturbing, however, is the fact that despite many decades of intensive research no effective and/or definitive therapeutic treatment aimed at reducing or retarding the clinic and pathologic symptoms induced by this devastating neurologic disorder is at present available. These still unmet needs call for ever-more concerted and focused efforts in basic, clinic and applied research to save human lives from a death without *memoir* (Weuve et al. 2014). Since direct study, *in vivo*, of the human brain is either technically challenging or at best extremely limited, the biochemical mechanisms involved in neuronal loss have mainly been studied in biological models.

The fly *Drosophila melanogaster* has been used as valid *in vivo* model of AD. For the scope of this review, PubMed and Scopus searches were conducted to find data within the past 16 years (1998–2014) involving the molecular biology of AD and the use of polyphenols in the fly to understand the molecular actions and feasibility of these compounds in the treatment of AD. Additionally, the authors propose a cell-death molecular mechanism induced by oxidative stress (OS) as a minimal working *in vitro* and *in vivo* model of AD to better understand neuronal demise and to effectively design antioxidant therapeutic strategies against AD.

2.2 Alzheimer Is Knocking on Paisas' Doors

In 1991, Goate and colleagues discovered the first mutation (Val717Ile) linked to an inherited form of AD, in the amyloid precursor protein ($A\beta$ PP) gene on chromosome 21q21.2. The mutation was found to be linked to inherited cases of early-onset AD (EOAD) (Goate et al. 1991). By 1995,

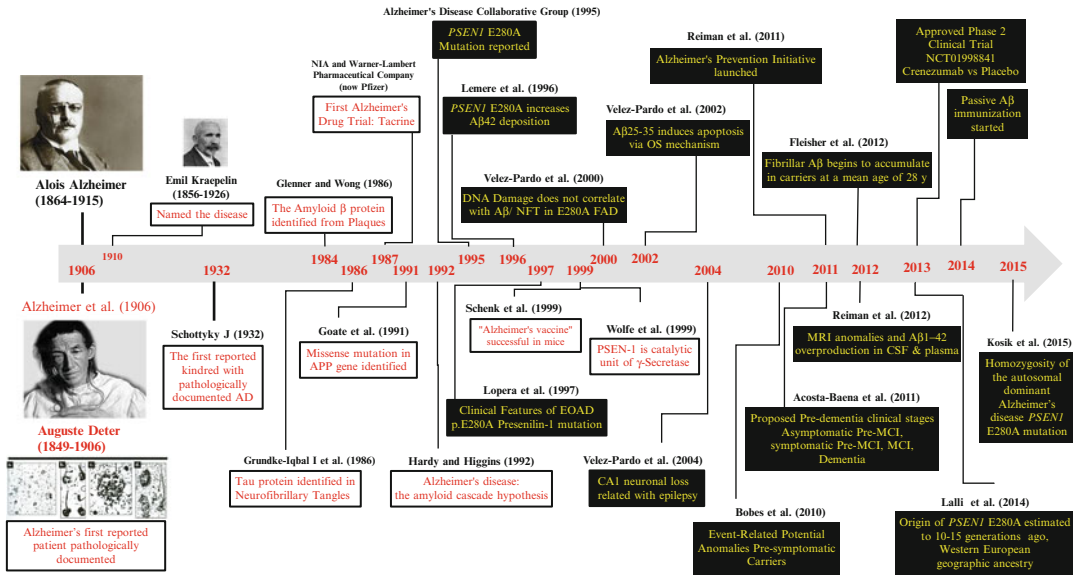


Fig. 2.1 More than 100 years of solitude. Historical timeline of important discoveries in Alzheimer's disease and the presenilin 1 E280A (g.50024 A>C, c.839A>C, p.E280A) autosomal-dominant familial mutant

Sherrington and co-workers (Sherrington et al. 1995) reported for the first time 5 mutations in the presenilin 1 (*PSEN 1*) gene located on chromosome 14q24.3 which were associated with EOAD. That same year, the Alzheimer's Disease Collaborative Group reported the structure of *PSEN 1* and identified the mutation p.E280A, a glutamic acid-to-alanine mutation at codon 280 (g.50024 A>C, c.839A>C, Ex 8) (Alzheimer's Disease Collaborative 1995). Similar to the APP mutation, the *PSEN 1* p.E280A mutation causes EOAD. This finding clarified previous descriptions of a family affected by Alzheimer-type presenile dementia with familial segregation in Antioquia, Colombia (Cornejo et al. 1987; Lopera et al. 1994). These initial observations in turn paved the way for important genetic, molecular and cellular, clinical, cognitive and neuropathological characterization of the *PSEN 1* mutation in the Antioquian kindred, also known as the "Paisas" people (Lopera et al. 1997; Velez-Pardo et al. 2004; for a review see Sepulveda-Falla et al. 2012). The "Paisas" are a genetically isolated community with more than 5,000 individuals (Arcos-Burgos and Muenke 2002). It is worth mentioning that the age and geographic origin of

c.839A>C mutation are consistent with a single founder (i.e. a *de novo* mutation event 15 generations ago) dating from the time of the Spanish Conquistadors who began colonizing Colombia during the early sixteenth century (Lalli et al. 2014). Interestingly, the search for a *PSEN 1* mutation in DNA extracted from a histological section of Auguste Deter's brain showed a T>C substitution at position 526 (c.526 T>C) resulting in a phenylalanine-to-leucine amino acid change at codon 176 (p.F176L) (Müller et al. 2013). This finding, a mutation in *PSEN 1* as in the Colombian study, revealed the cause of the disease and explained the histopathological observations previously noted by Alzheimer in this notable historical case. The most important findings associated with the *PSEN 1* p.E280A mutation and AD are outlined in Fig. 2.1.

By 2003, it was clear that *PSEN 1* (an aspartyl protease at its catalytic core (Wolfe et al. 1999; for a review, see De Strooper et al. 2012)) together with nicastrin (Nct), anterior pharynx defective 1 (Aph-1) and presenilin enhancer 2 (Pen 2) generate an active γ -secretase complex (Kimberly et al. 2003b; De Strooper 2003). This multimolecular complex (Haass 2004)

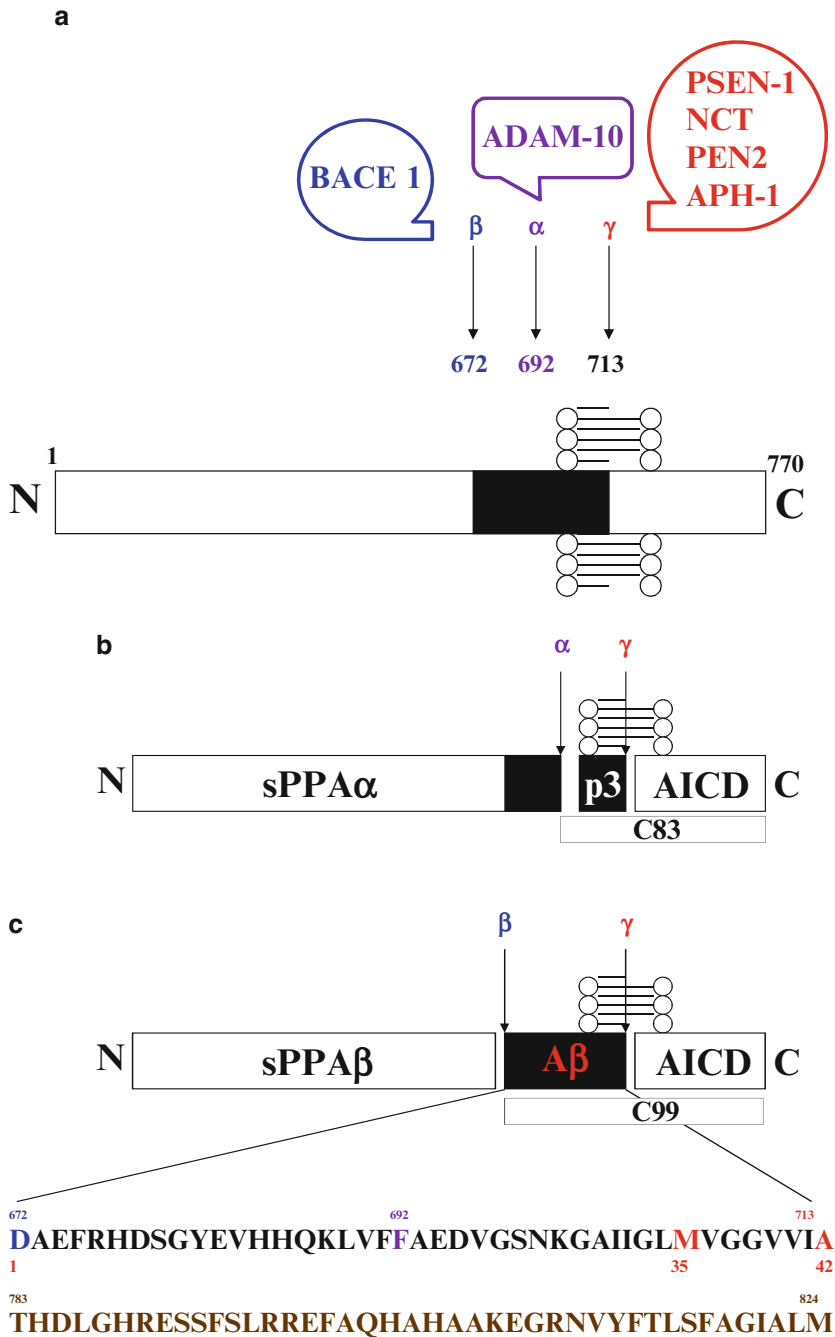


Fig. 2.2 Amyloid- β precursor protein (APP) metabolism. The APP is initially cleaved at either the α - (692 aa) or β - (672 aa) site (**a**) by α -secretase (i.e. ADAM-10) and β -secretase (i.e. BACE-1) to release the large ectodomains, designated APP α (**b**) and APP β (**c**), respectively. The ectodomain shedding leaves a membrane-embedded fragment, C83 or C99, which is a substrate for γ -secretase complex cleavage (i.e. PSEN-1, PEN-2, NCT, APH-1 complex). Proteolysis by γ -secretase complex releases the p3 (**b**) and A β fragments

(from 673 to 713 aa) at γ -site cleavage (**c**) as well as the carbonyl terminal- γ (ϵ -site cleavage), which forms the AICD. Protein aminoacid alignment between *Homo sapiens* A β (1–42) (in red) and *Drosophila melanogaster* (in brown) shows neither similarity nor identity. (**d**) The lonely pair electrons of methionine-35 residue in A β can react with oxygen to form both a sulfuranyl cation radical ($-S^{\bullet+}$) and anion superoxide radical ($O_2^{\bullet-}$), which can then either initiate the process of lipid peroxidation or H_2O_2 generation, respectively

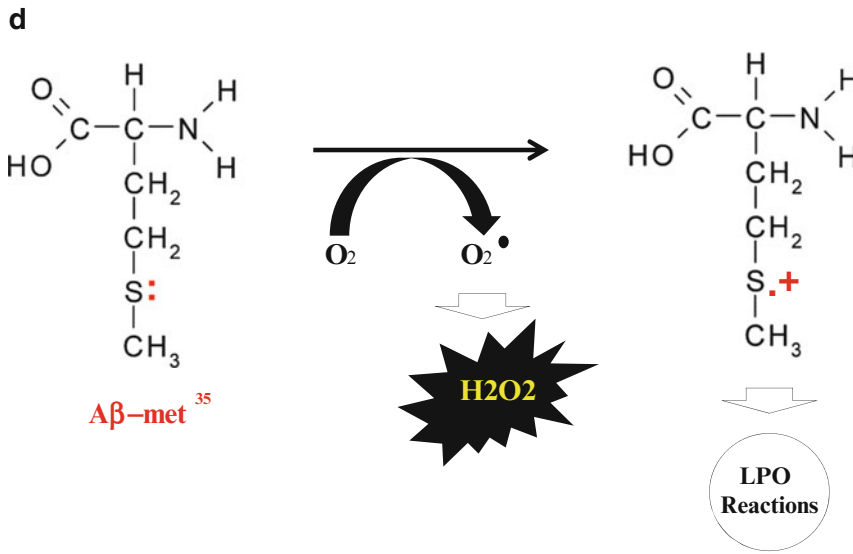


Fig. 2.2 (continued)

cleaves many type-I transmembrane proteins including AβPP (De Strooper et al. 1998) and Notch (Kimberly et al. 2003a) (Fig. 2.2a). It turns out that the AβPP is first cleaved by two other secretases – the α-secretase (also known as ADAM-10, for α-disintegrin and metalloprotease, (Kuhn et al. 2010)) or β-secretase (commonly known as BACE 1, for beta-site APP-cleaving enzyme, (Vassar et al. 1999)) to form the C83 (Fig. 2.2b) and C99 (Fig. 2.2c) residue membrane-bound fragments, respectively. C83 and C99 are substrates for a second cleavage (at the so-called γ-site) by the γ-secretase activity to form the p3 (non-amyloidogenic) and amyloid-β (Aβ) peptides, mainly Aβ(1–40) and Aβ(1–42), the latter being amyloidogenic (Fig. 2.2c). Further γ-secretase cleavage along the transmembrane domain at the so-called ε-site (A⁴²TVIVITL⁴⁹) generates the APP intracellular domain (AICD) (Fig. 2.2c). The function of AICD is unknown (Beckett et al. 2012). At present, several mutations in *PSEN1* (including the c.839A>C), *PSEN 2* (a homologue gene localized on chromosome 1q42.13) and *APP* gene (www.molgen.vib-ua.be/ADMutations/) have been demonstrated to modulate the position of this ε-site and to increase the ratio of the 42-residue form of Aβ versus the 40-residue

Aβ (Aβ₄₂:Aβ₄₀), thus defining a common AD phenotype caused by those mutations (Sato et al. 2003). Despite this advanced knowledge, exactly why the γ-secretase reaction results in more Aβ(1–42) and less Aβ(1–40) is not yet clear.

Nevertheless, these exciting developments have provided a biochemical framework for the emergence of the “amyloid cascade hypothesis” (Hardy and Higgins 1992). The hypothesis posits that deposition of the Aβ peptide in the brain parenchyma initiates a sequence of events that lead to the formation of paired helical filaments (PHFs) of tau aggregates and, ultimately, result in NFT formation, neuronal loss and AD dementia. This hypothesis has been modified and/or re-examined over the years (e.g. Hardy and Selkoe 2002; Golde et al. 2006; Hardy 2006, 2009; Korczyn 2008; Karran et al. 2011; Golde et al. 2011; Giacobini and Gold 2013; Drachman 2014). The amyloid cascade hypothesis, as it currently stands, suggests that synaptotoxicity and neurotoxicity may be mediated by transient Aβ peptide forms lying between free, soluble Aβ monomers and/or insoluble amyloid fibrils (Nimmrich and Ebert 2009; Mucke and Selkoe 2012; Hubin et al. 2014). The intermediary Aβ forms most probably damage neuronal synapses either via Aβ-receptor interactions (e.g. leukocyte

immunoglobulin-like receptor B2, LirB2 (Kim et al. 2013)) or by A β pore-membrane formation (Parodi et al. 2010; Sepúlveda et al. 2014).

Such observations imply that A β might lead to early brain atrophy in familial Alzheimer disease (FAD) mutation carriers (Cash et al. 2013). Recent clinical studies from the Colombian “Paisas” families bearing the *PSEN 1* E280A mutation support this view. Indeed, Acosta-Baena et al. (2011) have proposed that clinical deterioration in p.E280A carriers can be detected as measurable cognitive impairment around two decades before dementia onset. Remarkably, the median age at onset was 35 years for asymptomatic pre-mild cognitive impairment (MCI), 38 years (37–40) for symptomatic pre-MCI, 44 years (43–45) for MCI, and 49 years (49–50) for dementia. It was also calculated that the median time of progression from asymptomatic to symptomatic pre-MCI was just 4 years (95 % CI 2–8). This data was further supported when it was found that fibrillar A β began to accumulate in *PSEN 1* mutation carriers at a mean age of 28 years, about 16 years and 21 years before the predicted median ages at MCI and dementia onset, respectively (Fleisher et al. 2012). Furthermore, Reiman and co-workers reported that, although the carrier and non-carrier groups did not differ significantly in their dementia ratings, neuropsychological test scores, or proportion of apolipoprotein E (*APOE*) ϵ 4 carriers, the *PSEN 1* carriers had greater right hippocampal and parahippocampal activation, less precuneus and posterior cingulate deactivation, and less gray matter in several parietal regions (Reiman et al. 2012). Further A β (1–42) quantification in cerebrospinal fluid (CSF) and plasma showed that carriers had higher CSF concentrations and plasma A β (1–42) concentrations than non-carriers. These data clearly showed that young adults at genetic risk for autosomal dominant Alzheimer’s disease have functional and structural MRI findings, as well as CSF and plasma biomarker findings, consistent with A β (1–42) overproduction. Assessment of specific brain metabolites levels by two-dimensional proton magnetic resonance spectroscopy (¹H-MRS) in the posterior cingulate gyrus and precuneus are potentially sensitive and specific noninva-

sive biomarkers of subclinical emergence of AD caused by the *PSEN1* p.E280A mutation (Londono et al. 2013). Further, cognitively normal carriers of the p.E280A mutation have a thinner cerebral cortex in angular gyrus, precuneus and superior parietal lobule regions with trend-level effects in the medial temporal lobe compared to non-carriers, as measured by T1-weighted volumetric MRI (Quiroz et al. 2013; Bobes et al. 2010). Collectively, these data suggest that early overproduction of A β by *PSEN 1* E280A (Lemere et al. 1996) induces a pre-clinical stage in individuals genetically determined to develop AD (Acosta-Baena et al. 2011) that is detectable either by noninvasive techniques that evaluate the brain’s chemistry/structure *in vivo* such as by ¹H-MRS and volumetric/functional MRI imaging, or by biochemical measurements of A β (1–42) in CSF/plasma.

These observations imply that A β -induced neurodegeneration precedes aggregation in presymptomatic *PSEN 1* mutation carriers, and that the neuronal injury might be the result of either free, soluble A β species and/or insoluble amyloid aggregates not detectable at very early stages of the disease (Burgold et al. 2014). Furthermore, it is anticipated that presymptomatic, pre-MCI *PSEN 1* mutant carriers might develop a “pre-plaque-only” or “plaque-only” form of AD similar to the histopathology findings in the brain of Johann F., Alzheimer’s second patient, who lacked NFT (Klünemann et al. 2002). This clearly implies that A β might cause neuronal dysfunction at an early stage of AD independently of tau protein pathology (Palop and Mucke 2010). We propose that complete corroboration of these assumptions might come from functional, structural and biochemical brain studies in adolescents, or even children (6–10 years old) that are carriers of the *PSEN 1* c.839A>C mutation (Dean et al. 2014).

How A β induces memory loss and neuronal death is still unknown. However, at least three different mechanisms have been proposed to explain the A β -induced neurotoxicity: an extracellular mechanism (i.e. the amyloid cascade hypothesis), an intracellular mechanism (Capetillo-Zarate et al. 2012; Takahashi et al. 2013) and

A β -induced oxidative stress (Butterfield et al. 2013). Elucidating which mechanism is truly operative, or which is hierarchically relevant *in vivo* is essential for therapeutic success (Golde 2009). To further complicate matters, the co-occurrence of Alzheimer's A β and tau at synaptic terminals has been demonstrated (Takahashi et al. 2010). This observation implies tau as a mediator for Alzheimer's disease-related synaptic deficits (Liao et al. 2014). Therefore, AD is a genetically complex, slowly progressive, and irreversible disorder of the brain involving multiple pathways and molecular mechanisms, yet to be clearly delineated.

Nonetheless, several therapeutic approaches have been considered, including (i) inhibition of either β - or γ -secretase that generate A β from β APP (Ghosh and Osswald 2014; De Strooper et al. 2010; Ran et al. 2014); (ii) prevention of A β oligomerization or enhancing its clearance by active or passive immunization (Lambracht-Washington and Rosenberg 2013; Fettelschoss et al. 2014; Schenk et al. 1999); (iii) the use of metal chelators to avoid metal-induced rapid A β aggregation (Sharma et al. 2012; Squitti 2012); (iv) anti-inflammatory agents (Morales et al. 2014); (v) use of cholesterol-lowering drugs (Barone et al. 2014); and (vi) use of antioxidant agents (Aliev et al. 2008), among other therapeutics (<http://www.alzforum.org/therapeutics>). Unfortunately, no clinical proof has been provided yet that validates the amyloid cascade hypothesis (Mullane and Williams 2013). This makes preventive therapies a desirable aim in future clinical trials. No doubt, the Alzheimer's prevention (API) Colombia initiative, a plan to accelerate the evaluation of pre-symptomatic treatments in the "Paisa" community, might provide critical information for the advancing of AD treatments (Reiman et al. 2011; Ayutyanont et al. 2014; Kosik et al. 2015).

A β has naturally been at the center of therapeutic strategies (Aisen 2009; Bayer and Wirths 2014), and tau has recently been recognized as another important player in the neurodegenerative AD process. It is thus reasonable to assume that therapeutic

approaches should be jointly directed against both amyloidogenic proteins (Boutajangout and Wisniewski 2014; Golde et al. 2011). Furthermore, due to the multifactorial nature of AD, we think that pleiotropic interventions are urgently needed for individuals at genetic risk of AD, as elegantly proposed by Frautschy and Cole (Frautschy and Cole 2010), especially for asymptomatic *PSEN 1* p.E280A mutation carriers. It is hoped that present and future development efforts may release not only the "Paisas" from Dr. Alzheimer's visit, but help the scientific community to further understand this fatidic *maladie*.

At present, there is as yet no definitive cure for AD. It is therefore imperative to answer two basic questions. First, what is (are) the molecular mechanism(s) responsible by which A β induces selective neural damage and death leading to brain dysfunction? Second, what are the potential therapeutic approaches that could be implemented to retard, prevent or interrupt the pathophysiological process in AD? Our approach to clarify these uncertainties involves the use of lymphocytes as a cell model, and *Drosophila melanogaster*, as *in vivo* model of AD.

2.3 β Adly Treated Lymphocytes: A Model of Oxidative Stress in Alzheimer's Neurons

Oxidative stress (OS) refers to an imbalance in which free radicals (defined as any atom or molecule that has one or more unpaired electrons in its outer shells) and their products (e.g. superoxide anion radical, O $_2^{\bullet-}$; hydroxyl radical, \bullet HO; sulfhydryl radical, $-S\bullet$; nitric oxide, NO \bullet ; lipid peroxy radical, LOO \bullet) are in excess of enzymatic (e.g., catalase, CAT; glutathione peroxidase, GPx) and non-enzymatic defense (e.g. vitamin E & C) mechanisms. Chemically, free radicals and products, collectively known as reactive oxygen species (ROS), are predominantly generated by the transfer of one electron to oxygen (O $_2$) into superoxide ion (O $_2^{\bullet-}$). Addition of one more electron to O $_2^{\bullet-}$ will generate the peroxide ion (O $_2^{2-}$). The addition of 2H $^+$ to

O_2^{2-} ion results in hydrogen peroxide (H_2O_2 , a non-radical oxygen species) which decomposes rapidly into the more reactive hydroxyl radical, $HO\bullet$ in the presence of $O_2^{\bullet-}$ (Haber-Weiss reaction: $O_2^{\bullet-} + H_2O_2 + H^+ \rightarrow HO\bullet + O_2 + H_2O$) or metals (e.g. Fenton reaction: $Fe_2^+ + H_2O_2 + H^+ \rightarrow Fe_3^+ + HO\bullet + H_2O$). Because those ROS are capable of reacting with lipids, proteins, nucleic acids and other molecules, OS can lead to alterations in cells with an accumulation of oxidized products such as aldehydes and isoprostanes from lipid peroxidation, protein carbonyls from protein oxidation, and base adducts from DNA oxidation.

All of these chemical imprints of OS have been amply found in AD brain (Praticò 2008) and support the hypothesis that genetic alterations (e.g. over-production of $A\beta(1-42)$ by *PSEN-1* mutations), environmental factors (e.g., Fe, Cu or Mn metal ions (Gaeta and Hider 2005)), diminished cerebral energy metabolism and excitotoxic events, working in a convergent or complimentary manner, induce OS to damage and kill AD neurons (Markesbery 1997). Indeed, recent evidence indicates that OS phenomenon is an early event and might have a pathological role in the pathogenesis of this disease (Nunomura et al. 2001).

Not surprisingly, $A\beta$ has been suggested to play a central role in the OS damage in AD brain (Butterfield et al. 2001; Khan et al. 2006; Smith et al. 2007). Although the mechanism by which $A\beta(1-42)$ generates ROS is not yet fully established, methionine (met) at position 35 (Met³⁵) of $A\beta(1-42)$ (Fig. 2.2c) has been implicated as a critical amino acid for the OS and neurotoxic properties in AD brain (Butterfield and Sultana 2011; Butterfield et al. 2001). Most probably, the reduction of molecular O_2 by the sulfur-containing element of met³⁵ yields $O_2^{\bullet-}$, which then undergoes spontaneous or enzyme-catalysed dismutation to H_2O_2 (Fig. 2.2d). Indeed, H_2O_2 is generated as a short burst early on in the aggregation process of amyloid peptides and British dementia peptide (Tabner et al. 2005). Furthermore, Behl et al. (1994) demonstrated that $A\beta(1-42)$ causes increased levels of H_2O_2 to

accumulate in cells and lipid peroxidation. Interestingly, they also found that $A\beta(1-42)$ induces the activity of the nuclear factor-kappaB (NF- κ B), a transcription factor thought to be regulated by OS. This last observation implicates $A\beta(1-42)$ as being able to trigger oxidative signaling mechanisms involved in cell death.

Together, these data prompted us to test whether $A\beta(1-42)$ might induce H_2O_2 generation, NF- κ B activation and cytotoxicity in lymphocyte cells. Lymphocytes were selected as a model system based on their biochemistry, metabolic and physiological resemblance to neuronal cells (Jimenez-Del-Rio and Velez-Pardo 2012). In fact, lymphocytes not only express homologous neurochemical systems (e.g. catecholaminergic, serotonergic, cholinergic, glutamatergic, adrenergic systems) pertaining to neurons, but also express β APP751 and β APP770 mRNA isoforms (Ebstein et al. 1996), PSEN1 and PSEN 2 proteins (Mirinics et al. 2002; Takahashi et al. 2003), glucose transporters Glut 1 and 3 (Fu et al. 2004), and IGF-1 (Tapsom et al. 1988; Kooijman et al. 1992), among others. Most importantly, lymphocytes and neurons are post-mitotic cells, i.e. in the G_0 phase of the cell cycle, and both cells express similar molecular death machinery. It is worth mentioning that lymphocytes have even been postulated as potential diagnostic and/or progression biomarkers of AD (Sultana et al. 2013; Song et al. 2012). Hence, peripheral blood lymphocytes are valid and important experimental tools for understanding the molecular signalling machinery and metabolic processes of cell death/survival in AD neurons.

Effectively, we have demonstrated that lymphocytes exposed to (10 μ M) $A\beta(25-35)$ (i.e. the cytotoxic functional sequence of the amyloid peptide, for a review see (Kaminsky et al. 2010)) induces apoptosis either alone or in the presence of (25 μ M) iron in a concentration-dependent fashion by an OS mechanism (Velez-Pardo et al. 2002). Interestingly, $A\beta(25-35)$ was able not only to generate H_2O_2 and induce NF- κ B activation, but also induced p53 and c-Jun transcription factors, c-Jun N-terminal kinase

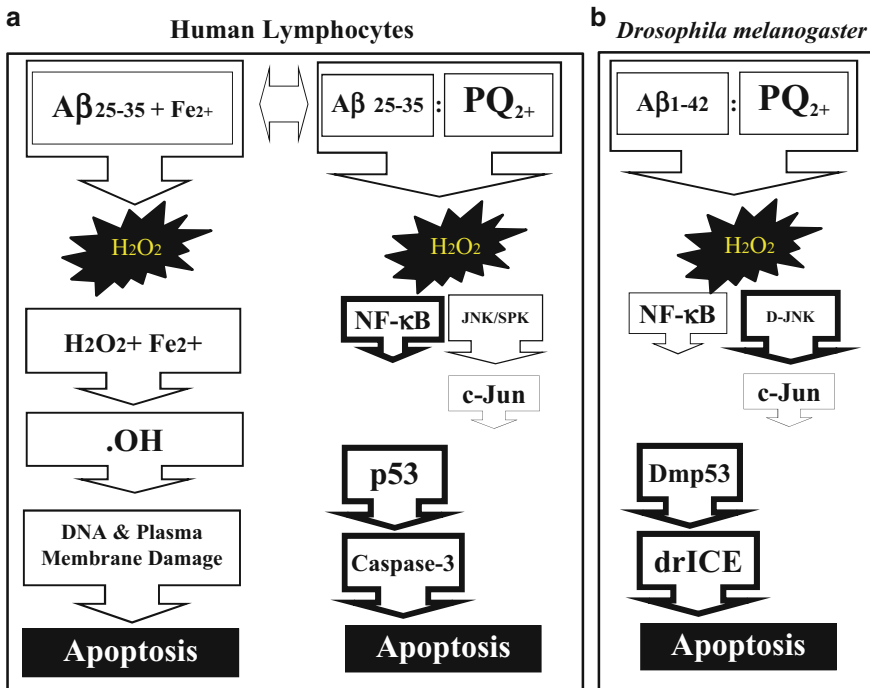


Fig. 2.3 Schematic model of Aβ(25–35) and Fe²⁺-induced apoptosis by oxidative stress mechanism in peripheral blood lymphocytes (PBL) and *Drosophila melanogaster* neurons. (a) The Aβ(25–35) fragment alone or in combination with Fe²⁺ generates H₂O₂. The latter might activate in parallel NF-κB and JNK/SAPK pathways, which in turn activate the p53 and c-Jun transcription factors, respectively. P53 in turn activates the pro-apoptotic Bax protein, which induces cytochrome c release from mitochondria to activate the apoptosome com-

plex. This complex activates caspase-3, leading PBL cells to cell death by apoptosis. Alternatively, H₂O₂ reacts with Fe²⁺ to produce highly reactive oxygen species (probably hydroxyl radicals by Fenton reaction), which directly damage nuclear DNA and provoke plasma membrane rupture leading to apoptosis. Noticeably, paraquat (PQ²⁺) induced a similar mechanism in PBL. (b) PQ²⁺ induces a similar oxidative stress mechanism in *Drosophila melanogaster* neuronal cells (a is modified and reproduced with permission from Velez-Pardo et al. (2002))

(JNK) and caspase-3 activation, mitochondrial damage, chromatin condensation and nuclei fragmentation – all features typical of apoptosis (Fig. 2.3a). Not surprisingly, specific inhibitors (e.g. pifithrin-α and Ac-DEVD-cho, inhibitors of p53 and caspase-3, respectively), antioxidant agents (e.g. *N*-acetyl-cysteine, the cannabinoids CP55940 and JHW-015) and growth factors (e.g. insulin growth factor-1) protected lymphocytes against Aβ(25–35)/H₂O₂-induced apoptosis (Jimenez Del Rio and Velez-Pardo 2006; Velez-Pardo and Del Rio 2006). These results imply that H₂O₂ generation and OS precede the apoptotic process, and that once it is generated, H₂O₂ is able to trigger apoptotic signaling (Marinho et al. 2014).

But then how does H₂O₂ induce apoptosis in lymphocytes? Although H₂O₂ itself is a mild oxidant and is relatively inert with most biomolecules, it is able to oxidize cysteine residues in proteins to either cysteine sulfenic acid (Cys-SOH) or disulfide bond (Cys-S-S-Cys). This oxidative chemical process might change the quaternary structure of the target protein, thereby activating/inactivating it. In fact, it has been demonstrated that H₂O₂ is able to indirectly activate the NF-κB by at least three different mechanisms (Jung et al. 2008; Schoonbroodt et al. 2000). Once NF-κB is freed from inhibition, the p50/p65 active dimeric protein translocates into the nucleus and transcribes either anti-apoptotic (e.g. *Bcl-2*, *cIAP-1* & 2, *Bcl-xL*) or

pro-apoptotic (e.g. *p53*) genes. At this point, a vicious cycle is established wherein *p53* plays a critical role by balancing the cell to a death decision because of its many actions. First, *p53* transcribes pro-apoptotic genes such as *Bax*, which in turn contribute to the permeabilization of the outer mitochondrial membrane by antagonizing anti-apoptotic proteins (e.g., Bcl-2, cIAP-1-2, Bcl-xL). Second, *p53* not only induces pro-oxidant genes (e.g., *p53-induced gene-3*, *proline oxidase*), which generate more H_2O_2 , but also represses the transcription of antioxidant genes (e.g., *phosphoglycerate mutase*, *NAD(P) H: quinone oxidoreductase-1*) (Olovnikov et al. 2009). Elevated stress stimuli by H_2O_2 production and further activation of NF- κ B induce up-regulation of pro-apoptotic genes (e.g., *p53*), which in turn amplify the initial H_2O_2 -induced cell death signal. Formation of the mitochondrial permeabilization transition pore allows the release of apoptogenic proteins, by a not fully established mechanism, such as the apoptosis-inducing factor (AIF, (Norberg et al. 2010)) responsible for DNA fragmentation and chromatin condensation (i.e., stage I nuclei morphology) and cytochrome *c*, which together with Apaf-1, dATP, and procaspase-9 (i.e., the apoptosome) elicits caspase-3 protease activation (Zou et al. 1999). This protease is essential for the fragmentation and morphological changes associated with apoptosis (i.e., stage II nuclei morphology (Jänicke et al. 1998)). H_2O_2 is also able to induce the apoptosis signal-regulating kinase (ASK1) activation by a mechanism that involves oxidation and release of thioredoxin from ASK-1 complex (Saitoh et al. 1998). Once activated, ASK-1 phosphorylates MKK4/MAPK kinase, which in turn phosphorylates c-Jun N-terminal kinase (JNK)/stress apoptotic protein kinase (SAPK) (Yang et al. 1997). JNK/SAPK signals cell death either via c-Jun activation (Gupta et al. 1996) or via stabilization and/or activation of *p53* (Oleinik et al. 2007; Fuchs et al. 1998). Importantly, JNK/SAPK can also phosphorylate tau protein *in vitro* (Goedert et al. 1997; Reynolds et al. 1997). In agreement with others, this last observation suggests that tangle pathology is a downstream process related to

A β signalization, consistent with the amyloid cascade hypothesis (Götz et al. 2001; Oddo et al. 2003; LaFerla 2010).

Is there a link between A β -induced apoptosis in lymphocytes and AD brain pathology? It would be possible to answer this question affirmatively, if one considers that A β -induced OS and most probably apoptosis in (*in vivo*) model AD brain (LaFerla et al. 1995; Selznick et al. 1999) and (*in situ*) human post-mortem AD brains (Selznick et al. 1999; Su et al. 2001; Zhao et al. 2003) occurs by similar mechanisms as those shown in A β -induced toxicity (*in vitro*) in lymphocytes. Of note, Garcia-Ospina et al. (2003) showed a correlation between DNA damage, detected by TdT-mediated dUTP Nick End Labeling (TUNEL) technique, with a positive detection of c-Jun, NF- κ B, *p53* and Par-4 transcription factors in post-mortem *PSEN-1* E280A AD brains. Further, it has been directly demonstrated that a subset of amyloid plaques produce ROS in living, Alzheimer's models (e.g., Tg2576 APP overexpressing transgenic mice) and in human post-mortem Alzheimer tissue (McLellan et al. 2003). Huang and co-workers found NF- κ B precursor p105 and I κ B γ inhibitor protein elevated in AD brain (Huang et al. 2005). Likewise, activation of *p53* (Cenini et al. 2008), caspase-3 (Su et al. 2001), c-Jun and JNK (Thakur et al. 2007; Shoji et al. 2000), AIF (Yu et al. 2010; Lee et al. 2012) and DJ-1, an antioxidant and OS sensor protein (Choi et al. 2006; Baulac et al. 2009), have been shown in AD brains. Collectively, these findings further support the notion that in AD brain either β -plaques or A β oligomers may induce cell death by a molecular signalling process similar to that found in our non-neuronal model exposed to A β . Indeed, increased OS, elevated apoptosis and decreased basal mitochondrial membrane potential levels, as well as enhanced sensitivity to different complex inhibitors of the respiratory chain, have all been reported in lymphocytes from AD patients (reviewed by Leuner et al. (2012)). Although other investigators have proposed complex disease pathways in AD pathogenesis (e.g. Crouch et al. 2008; Bettens et al. 2010), the A β -induced toxicity in lymphocytes should be interpreted as

a proposal for a minimal mechanism of cell death signaling induced by OS to explain neuronal demise.

Interestingly, we found that A β (25–35)-induced cytotoxicity can be recapitulated by paraquat (PQ), a mitochondrial redox cycling compound. Indeed, PQ induced a similar OS cell death mechanism in lymphocytes (Fig. 2.3a, Rio and Velez-Pardo 2008). Cannabinoids and growth factor protect and rescue lymphocytes against both A β and PQ exposure (Rio and Velez-Pardo 2008). Hence, it may be concluded that although the initiator cytotoxic molecules are clearly different in nature, they evidently converge on a common activator compound and pathway, namely the H₂O₂ signaling molecule that triggers an OS-mediated apoptotic mechanism (Jimenez-Del-Rio and Velez-Pardo 2004, 2012).

In conclusion, lymphocytes have been useful in deciphering molecular signaling events in AD. Nonetheless, they have to be viewed as a limited model for the study of other important aspects of the disease. For instance, lymphocytes do not express tau protein (Herrera-Rivero et al. 2013). Therefore, they provide no clues as to how A β interacts with such a protein. Furthermore, as a model system, lymphocytes are limited to answer questions in relation to A β -induced alterations of higher functions such as locomotion, memory and learning process. To address this problem, we changed our focus to the fruitfly *Drosophila melanogaster*. In the next section, we discuss the advantages of using the fly to model AD, and the fact that PQ might replace A β to further advance our understanding of the molecular mechanism of neuronal death in AD.

2.4 Next Patient: *Drosophila melanogaster*

In 1933, Thomas Hunt Morgan was awarded the Nobel Prize “for his discoveries concerning the role played by the chromosome in heredity” using *Drosophila melanogaster* as a genetics model. Since then, several other scientists have been awarded this prominent prize for discoveries in the fields of genetics (Hermann Muller, 1946),

development (Edward B. Lewis, Christiane Nusslein-Volhard and Eric F. Wieschaus, 1995), odour/olfactory systems (Richard Axel and Linda Buck, 2004) and immunity (Jules Hoffman, Bruce Beutler and Ralph Steinmann, 2011). What is it about flies that make them particularly useful to study Alzheimer's disease? From an evolutionary standpoint, *Drosophila* conserves several AD-related pathophysiological systems. To start with, basic local alignment search (BLAST) comparison between human and *Drosophila* proteins involved in memory processes showed high identity and similarity homology (Table 2.1). Strikingly, the fundamental molecular mechanisms implicated in implicit and explicit memory storage work similarly in *Drosophila* as well as in *Aplysia* and mammals (see Frank and Greenberg 1994; Silva and Murphy 1999; Skoulakis and Grammenoudi 2006; Barco et al. 2006; Kandel 2012; Kandel et al. 2014, among others). The fly has a complex nervous system that consists of 100,000 neurons, which includes a subset of 200 neurons that contain dopamine and serotonin neurotransmitters involved in memory processes (Waddell 2010; Berry et al. 2012; Sitaraman et al. 2012). Moreover, the fly possesses 2,500 Kenyon neuron cells and axons (i.e. α/β , α'/β' , γ lobes) constituting the mushroom bodies (MB), which are anatomical structures analogous to the human hippocampus (Campbell and Turner 2010), and central complex ellipsoid bodies involved in olfactory memory (Perisse et al. 2013a) and visual place learning and memory. Consequently, the fly can be trained to learn and perform complex behaviours requiring short- and long-term memory tasks (Ali et al. 2011; Ofstad et al. 2011; Perisse et al. 2013b; Pitman et al. 2009; Foucaud et al. 2010). Conclusively, research with *Drosophila* has provided essential information to enhance our understanding of human memory (Chakraborty et al. 2011) and early-onset cognitive disorders (Oortveld et al. 2013; van der Voet et al. 2014).

Notably, APP orthologs have been identified too in *Drosophila* (Rosen et al. 1989; Luo et al. 1992), and their processing is also evolutionarily preserved (Tharp and Sarkar 2013,

Table 2.1 Comparison of amino acid sequences of proteins involved in learning and memory, amyloid- β processing protein, cell death (apoptosis) and metal metabolism by basic local alignment search tool (BLAST) in *Homo sapiens* and *Drosophila melanogaster*

System	Protein name	<i>Homo sapiens</i>	Accession number	<i>Drosophila melanogaster</i>	Accession number	BLAST result: Identity, %; Similarity, %; (E)
Learning and memory	Ca(2+)/calmodulin-activated adenylyl cyclase	Adenylate cyclase type 1	Q08828.2	Ca(2+)/calmodulin-responsive adenylyl cyclase	P32870.2	46 %; 64 %; (1×10^{-112})
		Adenylate cyclase type 8	P40145.1	Ca(2+)/calmodulin-responsive adenylyl cyclase	P32870.2	58 %; 76 %; (6×10^{-139})
	Protein kinase A	PKA alpha (beta)	P17612.2	PKA c1 Isoform B (C,D)	AAN10703.1	82 %; 89 %; (0.0)
		PKA gamma	P22612.3 P22612.3 P22612.3	PKA c1 Isoform B (C,D)	AAN10703.1	75 %; 85 %; (0.0)
		Protein kinase C	PKC epsilon	CAA46388.1	PKC	P13678.1
		PKC alpha	EAW89014.1	PKC	P13678.1	63 %; 79 %; (8×10^{-154})
		PKC gamma	EAW72161.1	PKC	P13678.1	44 %; 57 %; (8×10^{-144})
		PKC eta	AAH37268.1	PKC	P13678.1	55 %; 68 %; (0.0)
		PKC beta	AAH36472.1	PKC	P13678.1	63 %; 70 %; (1×10^{-153})
		PKC iota	AAB17011.1	PKC	P13678.1	43 %; 57 %; (3×10^{-136})
		PKC zeta	AAA36488.1	PKC	P13678.1	43 %; 60 %; (6×10^{-135})
		PKC theta	AAI13360.1	PKC	P13678.1	46 %; 62 %; (0.0)
		PKC mu	CAA53384.1	PKC	P13678.1	30 %; 49 %; (6×10^{-31})
	Protein phosphatase 1	Protein phosphatase-1	AAA36508.1	Protein phosphatase1	CAA39820.	88 %; 92 %; (0.0)
	Ubiquitin hydroxylase	Ubiquitin carboxyl-terminal hydrolase isozyme L1	P09936.2	Ubiquitin carboxyl-terminal hydrolase 36	Q9VRP5.3	50 %; 70 %; (8.9)
Phospholipase C		PLC beta1	AAF86613.1	Phospholipase C	AAA28820.1	50 %; 68 %; (1×10^{-140})
		PLC beta2	AAP35551.1	Phospholipase C	AAA28820.1	41 %; 60 %; (2×10^{-48})
		PLC beta3	CAA85776.1	phospholipase C	AAA28820.1	46 %; 63 %; (0.0)
		PLC beta4	AAI43869.1	Phospholipase C	AAA28820.1	46 %; 64 %; (3×10^{-128})
		PLC delta1	AAA73567.1	Phospholipase C	AAA28820.1	46 %; 59 %; (1×10^{-58})

(continued)

Table 2.1 (continued)

System	Protein name	<i>Homo sapiens</i>	Accession number	<i>Drosophila melanogaster</i>	Accession number	BLAST result: Identity, %; Similarity, %; (E)
		PLC delta3	AAH72384.1	phospholipase C	AAA28820.1	35 %; 52 %; (2×10^{-55})
		PLC delta4	AAH06355.1	Phospholipase C	AAA28820.1	45 %; 61 %; (4×10^{-63})
		PLC epsilon1	AAI51855.1	Phospholipase C	AAA28820.1	42 %; 58 %; (6×10^{-43})
		PLC gamma1	AAI44137.1	Phospholipase C	AAA28820.1	44 %; 58 %; (6×10^{-47})
		PLC gamma2	AAQ76815.1	Phospholipase C	AAA28820.1	34 %; 50 %; (2×10^{-49})
		PLC eta	AAI13951.1	Phospholipase C	AAA28820.1	29 %; 46 %; (5×10^{-88})
		PLC zeta	AAN71895.1	Phospholipase C	AAA28820.1	38 %; 56 %; (4×10^{-53})
	CCAAT-box-enhanced binding protein	C/EBP gamma	AAC50201.1	C/EBP	AAA28415.1	40 %; 73 %; (4×10^{-15})
		C/EBP delta	EAW86679.1	C/EBP	AAA28415.1	40 %; 70 %; (2×10^{-16})
		C/EBP beta	EAW75629.1	C/EBP	AAA28415.1	44 %; 68 %; (1×10^{-16})
		C/EBP epsilon	EAW66183.1	C/EBP	AAA28415.1	43 %; 64 %; (1×10^{-15})
		C/EBP zeta	AAH34475.1	C/EBP	AAA28415.1	34 %; 48 %; (0.078)
	Ca ²⁺ /Calmodulin protein kinase II- α	CaMKII- α	AAH40457.1	CaM-kinase II alpha	Q00168.1	71 %; 81 %; (0.0)
	Calcineurin	Calcineurin	AAC37581.1	Calcineurin A1, isoform A (D)	AAF57105.3	24 %; 48 %; (1.8)
		Calcineurin	AAC37581.1	Calcineurin B, isoform A (B)	AAF46026.1	36 %; 50 %; (4.6)
	Cytoplasmic polyadenylation element binding protein	CPEB1	Q9BZB8.1	orb2, isoform A (B,C,D, H)	AAF50352.1	42 %; 59 %; (2×10^{-63})
		CPEB2,3,4	Q7Z5Q1.3	orb2, isoform A (B,C,D,H)	AAF50352.1	88 %; 93 %; (2×10^{-174})
	NMDA receptor	NMDA	AAA21180.1	dNR 1	AAF52016.1	48 %; 67 %; (0.0)
		NMDA	AAA21180.1	dNR 2, isoform A (B,C,D,E,F,G)	AAN09051.2	29 %; 47 %; (3×10^{-91})

(continued)

Table 2.1 (continued)

System	Protein name	<i>Homo sapiens</i>	Accession number	<i>Drosophila melanogaster</i>	Accession number	BLAST result: Identity, %; Similarity, %; (E)
AMPA receptor	AMPA1	AMPA1	P42261.2	Glutamate receptor IA	AAF50652.2	40 %; 57 %; (0.0)
	AMPA1	AMPA1	P42261.2	Glutamate receptor IB, isoform A (B,C)	AAF50306.2	36 %; 53 %; (2×10^{-114})
	AMPA2	AMPA2	P42262.3	Glutamate receptor IA	AAF50652.2	42 %; 58 %; (0.0)
	AMPA2	AMPA2	P42262.3	Glutamate receptor IB, isoform A (B,C)	AGB94317.1	36 %; 53 %; (2×10^{-114})
	AMPA3	AMPA3	P42263.2	Glutamate receptor IA	AAF50652.2	42 %; 58 %; (0.0)
	AMPA3	AMPA3	P42263.2	Glutamate receptor IB, isoform A (B,C)	AAF50306.2	36 %; 52 %; (2×10^{-120})
	AMPA4	AMPA4	P48058.2	Glutamate receptor IA	AAF50652.2	43 %; 60 %; (0.0)
	AMPA4	AMPA4	P48058.2	Glutamate receptor IB, isoform A (B,C)	AAF50306.2	42 %; 56 %; (0.0)
Serotonin receptor	Serotonin receptor	Serotonin receptor	CAA05851.1	Serotonin receptor 1	P20905.1	83 %; 100 %; (1.5)
	Serotonin receptor	Serotonin receptor	AAA66493.1	Serotonin receptor 2A	P28285.2	48 %; 69 %; (3×10^{-61})
	Serotonin receptor	Serotonin receptor	AAA66493.1	Serotonin receptor 2B	P28286.3	48 %; 68 %; (3×10^{-62})
Dopamine receptor	Dopamine receptor D1A	Dopamine receptor D1A	P21728.1	Dopamine receptor	AAC47161.1	35 %; 48 %; (6×10^{-59})
	Dopamine receptor D1B(5)	Dopamine receptor D1B(5)	CAA41360.1	Dopamine receptor	AAC47161.1	42 %; 57 %; (4×10^{-40})
	Dopamine receptor D2	Dopamine receptor D2	P14416.2	Dopamine receptor	AAC47161.1	34 %; 49 %; (1×10^{-62})
	Dopamine receptor D3	Dopamine receptor D3	P35462.2	Dopamine receptor	AAC47161.1	36 %; 50 %; (3×10^{-65})
	Dopamine receptor D4	Dopamine receptor D4	P21917.2	Dopamine receptor	AAC47161.1	40 %; 54 %; (8×10^{-26})
	Dopamine receptor D1A	Dopamine receptor D1A	P21728.1	Dopamine receptor	CAA54451.1	40 %; 55 %; (6×10^{-71})
	Dopamine receptor D1B(5)	Dopamine receptor D1B(5)	CAA41360.1	Dopamine receptor	CAA54451.1	36 %; 52 %; (3×10^{-70})

(continued)

Table 2.1 (continued)

System	Protein name	<i>Homo sapiens</i>	Accession number	<i>Drosophila melanogaster</i>	Accession number	BLAST result: Identity, %; Similarity, %; (E)
		Dopamine receptor D2	P14416.2	Dopamine receptor	CAA54451.1	36 %; 56 %; (2×10^{-35})
		Dopamine receptor D3	P35462.2	Dopamine receptor	CAA54451.1	33 %; 46 %; (3×10^{-52})
		Dopamine receptor D4	P21917.2	Dopamine receptor	CAA54451.1	38 %; 57 %; (6×10^{-24})
	Tyrosine hydroxylase	Tyrosine hydroxylase	AAI43612.1	Tyrosine hydroxylase	CAA53802.1	54 %; 74 %; (5×10^{-168})
APP	APP	APP ₇₇₀	P05067.3	Appl	AAD55414.1	(36 %) (51 %) (8×10^{-37})
		APP ₇₅₁	NP_958816.1	Appl	AAD55414.1	(36 %) (51 %) (9×10^{-37})
		APP ₆₉₅	NP_958817.1	Appl	AAD55414.1	(36 %) (51 %) (2×10^{-37})
	Aβ42	Amyloid beta-peptide (1–42)	IITYT_A P05067.3	–	–	–
	PSEN	PS1	P49768.1	PS-D A (B-E)	AAF51598.1	53 %; 66 %; (5×10^{-145})
		PS2	AAP35630.1	PS-D A (B-E)	AAF51598.1	53 %; 67 %; (8×10^{-110})
	Nicastrin	Nicastrin-1	NP_056146.1	Nicastrin-A (C-E)	AAF56349.2	31 %; 48 %; (8×10^{-90})
		Nicastrin-2	NP_001277113.1	Nicastrin-A (C-E)	AAF56349.2	30 %; 48 %; (9×10^{-90})
		Nicastrin-3	NP_001277115.1	Nicastrin-A (C-E)	AAF56349.2	30 %; 46 %; (1×10^{-38})
	APH-1	APH-1A	Q96BI3.1	dAPH-1A	AAF51212.1	47 %; 66 %; (2×10^{-69})
		APH-1B	Q8WW43.3	dAPH-1B	AGB92500.1	44 %; 60 %; (5×10^{-59})
	PEN-2	PEN-2	Q9NZ42.1	pen-2	Q86BE9.3	61 %; 74 %; (3×10^{-45})
	BACE	BACE-1	P56817.2	dBACE	NP_609235.1	26 %; 42 %; (5×10^{-30})
		BACE-2	Q9Y5Z0.1	dBACE	NP_609235.1	29 %; 44 %; (2×10^{-30})
	α-Secretase	ADAM10	O14672.1	Kuzbanian	P07207.3	28 %; 41 %; (1.1)
	MAPT	MAPT	P10636.5	dTau	AAK54456.1	38 %; 56 %; (6×10^{-33})

(continued)

Table 2.1 (continued)

System	Protein name	<i>Homo sapiens</i>	Accession number	<i>Drosophila melanogaster</i>	Accession number	BLAST result: Identity, %; Similarity, %; (E)
Apoptosis	ASK-1	ASK-1	Q99683.1	DASK-1	BAC16514.1	42 %; 58 %; (0.0)
	MEKK-1	MEKK-1	Q13233.4	D-MEKK-1	AAF55592.3	34 %; 54 %; (1×10^{-45})
MEKK-1		Q13233.4	B-MEKK-1	AAN13787.1	34 %; 54 %; (1×10^{-45})	
MKK4	MKK4	P45985.1	MKK4-A (B,C)	AAF54258.1	67 %; 78 %; (7×10^{-153})	
IKK	IKK α	p25963	Cactus	AAA85908	37 %; 51 %; (1×10^{-28})	
	IKK β	Q15653	Cactus	AAA85908	30 %; 45 %; (5×10^{-19})	
NF- κ B	NF- κ B 1 (p50)	P19838	RELISH	AAF20138	34 %; 54 %; (2×10^{-44})	
	NF- κ B 1 (p50)	P19838	Dorsal	AAQ65068	39 %; 53 %; (1×10^{-47})	
	NF- κ B 1 (p50)	P19838	Dif	EDW90181	33 %; 48 %; (3×10^{-38})	
	NF- κ B 1 (p65)	Q04206.2	RELISH	AAF20138	36 %; 53 %; (1×10^{-41})	
	NF- κ B 1 (p65)	Q04206.2	Dorsal	AAQ65068	49 %; 61 %; (1×10^{-75})	
	NF- κ B 1 (p65)	Q04206.2	dif	EDW90181	36 %; 56 %; (6×10^{-62})	
JNK	JNK-1/MAPK8	AAI30571.1	dJNK (Basket)	P92208.1	80 %; 88 %; (0.0)	
c-JUN	c-JUN	CAG46552	d-Jun	CAA73154	55 %; 76 %; (1×10^{-22})	
P-53	p-53	AAC12971	Dmp-53	AAF75270	24 %; 42 %; (5×10^{-6})	
BAX	BAX	AAB35593.1	Drob-1	BAA89603.2	67 %; 66 %; (3.0)	
Cyto C	Cyto-C	AAA35732	Cyto c-A (B)	AAF55946.1	40 %; 40 %; (6.4)	
Caspase-9	Caspase-9	BAA87905.1	Dronc	AAF50180.1	33 %; 54 %; (1×10^{-13})	
Caspase-3	Caspase-3	CAC88866	drICE	CAA72937	42 %; 60 %; (1×10^{-46})	
DFF45	DFF45	O00273.1	dICAD-A (B)	AAF58589.1	35 %; 48 %; (2×10^{-9})	
DFF40	DFF40	BAA32250.1	dCAD	BAA97120	28 %; 43 %; (4×10^{-30})	
APAF-1	APAF-1	O14727.2	Dark: Apaf1-A (B)	AAM68488.1	23 %; 44 %; (3×10^{-9})	
AIF	AIF	AAD16436.1	AIF-A (B,C)	AAF51299.2	52 %; 68 %; (8×10^{-175})	

(continued)

Table 2.1 (continued)

System	Protein name	<i>Homo sapiens</i>	Accession number	<i>Drosophila melanogaster</i>	Accession number	BLAST result: Identity, %; Similarity, %; (E)
Metal metabolism	Ferritin	Ferritin heavy chain	NP_002023.2	Ferritin 1	AAF57034.1	37 %; 52 %; (5×10^{-28})
		Ferritin light chain	NP_000137.2	Ferritin 2	AAF57038.1	24 %; 43 %; (1×10^{-11})
	Transferrin	Transferrin	ABI97197.1	Transferrin-1	NP_523401.2	26 %; 41 %; (2×10^{-28})
		Transferrin	ABI97197.1	Transferrin-2	AAF49900.1	34 %; 49 %; (2×10^{-53})
		Transferrin	ABI97197.1	Transferrin-3	AAF58039.1	23 %; 38 %; (1×10^{-27})
	Divalent metal ion transportin	DMT-1	P49281.2	Malvolio	NP_732584.1	61 %; 76 %; (0.0)
	CTR-1	hCTR-1	O15431.1	Ctr1A	AAF46182.3	49 %; 61 %; (9×10^{-48})
		hCTR-1	O15431.1	Ctr1B	AAF54223.1	35 %; 51 %; (1×10^{-30})
	ATP7	ATP7A (B)	Q04656.3	DmATP7	AAF48104.3	47 %; 62 %; (0.0)

Abbreviations: AMPA α -amino-3-hydroxy-5-methyl-4-isoxazolepropionic acid (receptor), APP amyloid- β precursor protein, APH-1 anterior pharynx-defective 1, AIF apoptosis inducing factor, APAF apoptosis protease-activating factor-1, ASK-1 apoptosis signal-regulating kinase 1, BAX BCL2-associated X protein, BACE beta-site amyloid precursor protein cleaving enzyme 1, CAD Caspase Activated DNase, JNK c-Jun N-terminal kinases, CTR-1 Copper transporter 1, DIF differentiation-inducing factor-1, DFF DNA fragmentation factor, DRONC *Drosophila melanogaster* NEDD2-like caspase, DARK *Drosophila* Apaf-1-related killer, drICE *Drosophila* ICE (Interleukin-1 β -converting enzyme), ICAD Inhibitor of Caspase Activated DNase, IKK I κ B kinase, MEKK MAPK/ERK kinase, MAPT microtubule-associated protein tau, MKK Mitogen-activated protein kinase kinase, NMDA N-methyl-D-aspartate (receptor), NF- κ B nuclear factor-kappaB, PEN-2 presenilin enhancer 2

Fig. 2.2b). In fact, several reports have shown that *in vivo* reconstitution of the γ -secretase complex (i.e. fly PSEN, NCT, APH-1 and PEN-2 proteins) has biochemical properties and a subcellular distribution resembling those of endogenous γ -secretase (Hu and Fortini 2003; Stempfle et al. 2010). Additionally, *Drosophila* express α -secretase-like protein encoded by the *kusbanian* gene (Rooke et al. 1996), and BACE-1 like secretase activity (Greeve et al. 2004). Although there is a lack of amino acid sequence conservation between the human β -peptide and that of the fly (Fig. 2.2b), Carmine-Simmen et al. (2009) have shown that A β fragments derived from the *Drosophila* orthologue APPL aggregate into intracellular fibrils, amyloid deposits, and cause age-dependent behavioral deficits and neurodegeneration. Most importantly, it was shown that the behavioral phenotypes

precede extracellular amyloid deposit formation, supporting results that intracellular A β plays a key role in AD (Crowther et al. 2005). Therefore, *Drosophila* provides a comparable neurotoxic peptide effect from fly APPL. Additionally, several *Drosophila* AD models that express the toxic human A β 42 have shown amyloid deposits, learning and memory deficiencies, locomotor impairment, and premature death (Jahn et al. 2011) (for review, see refs. Iijima-Ando and Iijima 2010; Prüßing et al. 2013). Lately, Mhatre et al. (2014) have characterized a novel model of AD using *Drosophila melanogaster*, where they expressed the human AD-associated proteins APP and BACE in the central nervous system of the fly and showed synaptic defects in the larval neuromuscular junction (NMJ) leading to defective larval locomotion behavior, decreased presynaptic connections, altered mitochondrial

localization in presynaptic motor neurons and decreased postsynaptic protein levels. Most importantly, treatment of this model with the γ -secretase inhibitor L-685,458 suppressed the behavioral defects as well as the pre- and postsynaptic defects. These data comply with the notion that A β provokes early neuronal damage/alteration prior to A β aggregation and/or deposition. Based on such an assumption, one may wonder whether AD can be considered as a neurodevelopmental disorder rather than a neurodegenerative disease. Further investigation is needed to fully answer this question.

Another relevant point is that the apoptosis machinery is conserved in *Drosophila* (Abdelwahid et al. 2011, and Table 2.1). Despite the fact that A β induces neuronal cell death by activation of JNK (Troy et al. 2001; Tare et al. 2011), and causes mitochondrial dysfunction through OS (Abramov et al. 2004; Pagani and Eckert 2011; Spuch et al. 2012), no information is available concerning the role of p53 and other ortholog genes. Since PQ induces a similar molecular cascade of cell death as A β (Rio and Velez-Pardo 2008; Velez-Pardo et al. 2002) and *Drosophila* is amenable to ectopic gene knockdown or constitutive gene overexpression, we made use of the bipartite GAL4/Upstream Activation Sequence (UAS) system (Phelps and Brand 1998; Elliott and Brand 2008) to demonstrate that *Dmp53* (i.e. p53), *basket* (JNK) and *drICE* (caspase-3) gene knockdown in dopaminergic neurons (Fig. 2.4a–j) prolong life span and locomotor activity in *Drosophila melanogaster* lines chronically exposed to PQ, compared to untreated transgenic fly lines (Fig. 2.4h–j, Ortega-Arellano et al. 2013). In contrast, overexpression of *basket* and *Dmp53* flies sensitizes the transgenic flies to PQ treatment (Fig. 2.4k–m). It is anticipated that overexpression of A β might activate both *Dmp53* and *drICE*, leading to mitochondrial damage and contributing to neuronal death in *Drosophila* brain (Fig. 2.3c). It is therefore reasonable to suppose that A β might be able to trigger activation of *Dmp53*, *basket* and *drICE* via O $_2^{\bullet-}$ /H $_2$ O $_2$ signaling and cell death in a similar fashion as proposed in mammalian cells (Fig. 2.3a, b and Jimenez-Del-Rio and Velez-

Pardo 2012). Finally, *Drosophila* expresses similar protein sequences involved in iron and copper metabolism such as ferritin, transferrin, human high-affinity copper importer Ctr1, and divalent metal ion transporter (DMT-1) that may modify A β -induced neurotoxicity (Table 2.1). Effectively, overexpressing the subunits of the iron-binding protein ferritin can rescue the toxicity of the A β peptide in *Drosophila* AD model system (Rival et al. 2009). It has also been found that inhibition of Ctr1 or overexpression of the copper exporter DmATP7 reduce copper accumulation in the brains of the flies, ameliorate neurodegeneration, enhance climbing ability, and prolong lifespan (Lang et al. 2013). Therefore, iron and copper might play a critical role in AD pathophysiology (Greenough et al. 2013).

Interestingly, Iijima et al. (2010) have shown that co-expression of A β 1–42 with tau using the pan-neuronal retinal *glass multiple reporter* (gmr) promoter-GAL4 significantly enhanced the reduction in the external size of eyes and internal retina thickness compared to A β (1–42) expression alone, which did not significantly affect eye structures at this level of expression. Similarly, co-expression of A β (1–42) and tau using the pan-neuronal *embryonic lethal, abnormal vision* reporter (elav) promoter-GAL4 caused a complete loss of calyx (a dendritic region in the mushroom body) structures, whereas a normal calyx structure was observed in flies expressing A β (1–42) alone. Furthermore, phosphorylation of tau Ser²⁶² was found critical for A β (1–42)-induced tau toxicity. In agreement with these findings, Folwell and co-workers (2010) report that co-expression of A β (1–42) with tau (wild-type) increases tau phosphorylation and exacerbates tau-mediated phenotypes (i.e. neuronal dysfunction by disrupting axonal transport and synaptic structure resulting in behavioural impairments and reduced lifespan). Altogether, these results indicate that tau enhances A β -induced toxicity in *Drosophila* brain neurons, and reinforce the notion that tau – pathology is a downstream process related to A β signaling (Ploia et al. 2011; Chabrier et al. 2012). Based on the above observations, *Drosophila* seems to be an extremely useful and valid *in vivo* model

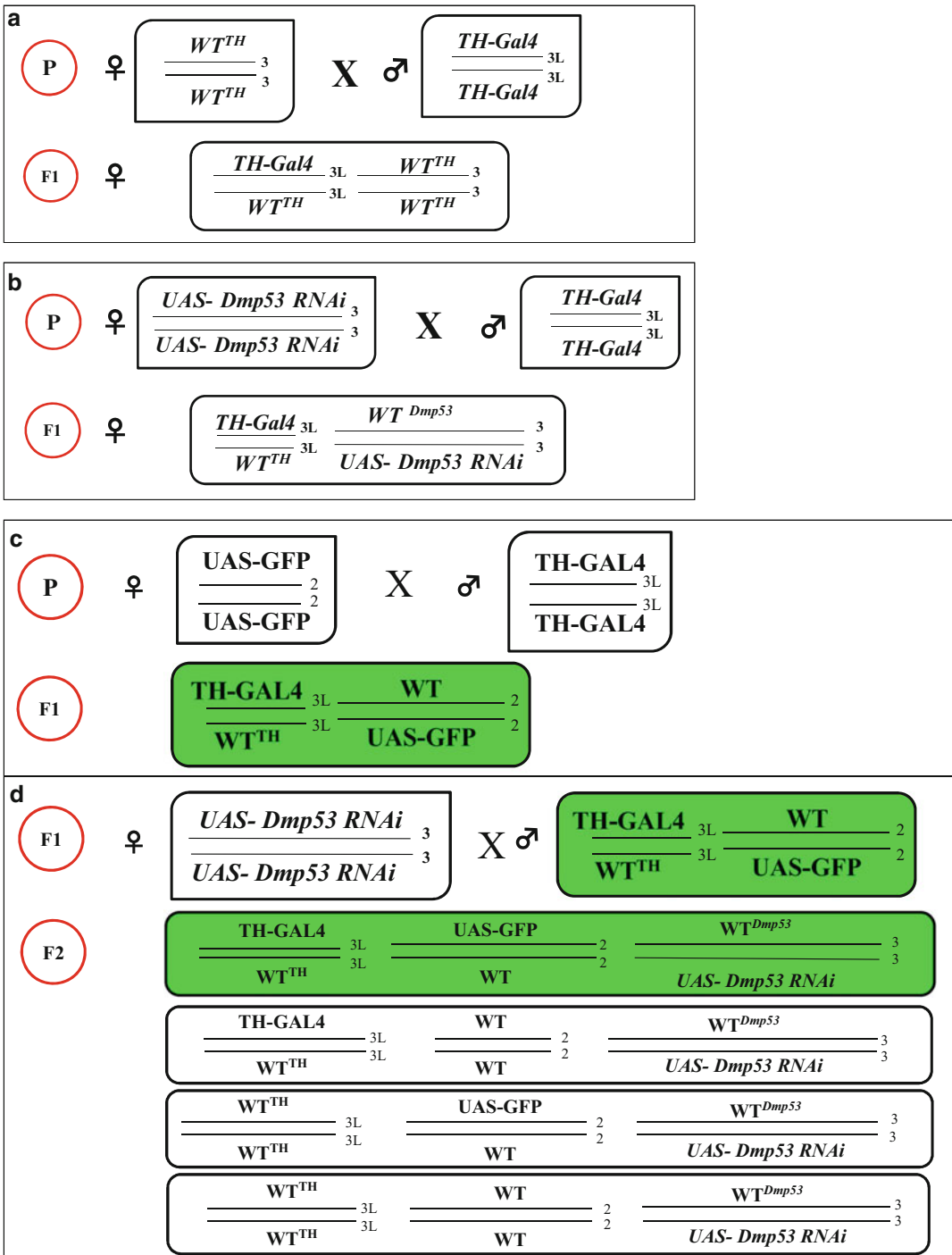


Fig. 2.4 (a) Scheme for basic fly cross and selection. *TH-Gal4*^{+/-} flies were obtained by crossing *TH-Gal4*^{+/+} (males, *n* = 10) and wild type Canton S *Drosophila melanogaster* (females, *n* = 10). After five days of husbandry, parental flies were discarded from the mating tubes. F1 flies were then reared according to stan-

dard procedures. (b) *TH-Gal4*^{+/-}, *UAS-Dmp53RNAi*^{+/-} flies were obtained by crossing *TH-Gal4*^{+/+} (males, *n* = 10) and *UAS-Dmp53RNAi*^{+/+} (females, *n* = 10). (c) F1 *TH-Gal4*^{+/-}; *UAS-GFP*^{+/-} flies were obtained by crossing *TH-Gal4*^{+/+} (males, *n* = 10) and *UAS-GFP*^{+/+} *Drosophila melanogaster* (females, *n* = 10).

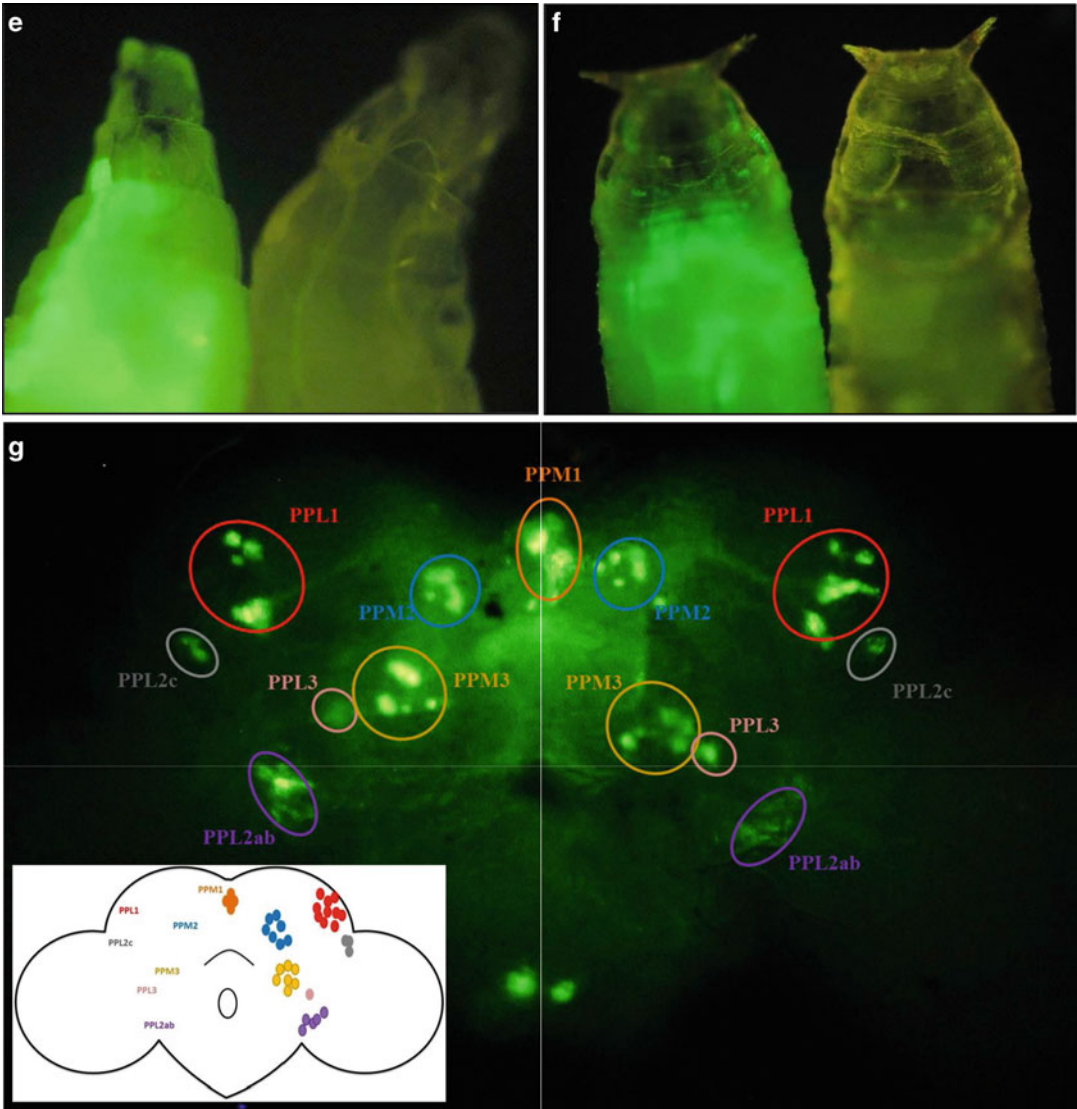


Fig. 2.4 (continued) (d) Males F1 *TH-Gal4^{+/-}; UAS-GFP^{+/-}* were crossed with females *UAS-Dmp53 RNAi^{+/+}* to obtain F2, where 1/4 represent *TH-Gal4^{+/-}; UAS-GFP^{+/-}; UAS-Dmp53 RNAi^{+/-}*. (e, f) Negative (translucent) and positive (green fluorescent) *TH-Gal4^{+/-}; UAS-GFP^{+/-}; UAS-Dmp53 RNAi^{+/-}* larvae (e) and pupae (f). (g) Illustrates six clusters on the posterior side in whole mount of dissected brain from untreated *TH-GAL4^{+/-}; UAS-GFP^{+/-}* flies: PPM1 (unpaired), PPM2 (paired), PPM3 (paired) (protocerebral posterior medial); PPL1, PPL2ab, PPL2c and PPL3 (paired) (protocerebral posterolateral). (h–m) Effect of paraquat (PQ) on the survival (h, k), locomotor activity (i, l) and dopaminergic neurons (j, m) knockdown (h–j, *TH-GAL4^{+/-}; UAS-Dmp53 RNAi^{+/-}*) or overexpression (k–m, *TH-GAL4^{+/-}; UAS-Dmp53*) in *Drosophila melanogaster*. *TH-GAL4^{+/-}; UAS-Dmp53 RNAi^{+/-}* (j)

and *TH-GAL4^{+/-}; UAS-Dmp53* (m) flies were fed with either 1 or 5 mM PQ alone or in combination with (0,1 mM) GA. Whole mounts of dissected brains were examined for DAergic neuron morphology and number count of neurons per brain hemisphere was recorded using a fluorescent microscopy (Axiostar plus 50) in each treatment according to ref. Bonilla-Ramirez et al. (2011). There were six clusters on the posterior side observed according to ref. Mao and Davis (2009): PPM1 (unpaired), PPM2 (paired), PPM3 (paired) (protocerebral posterior medial); PPL1, PPL2ab, PPL2c and PPL3 (paired) (protocerebral posterolateral). Average number of neurons of the medial and lateral dopaminergic neurons was scored for untreated and treated flies. The nonparametric Kruskal–Wallis test was performed to compare more than two independent groups. Differences were considered statistically significant at $P < 0.001$. The Mann–Whitney

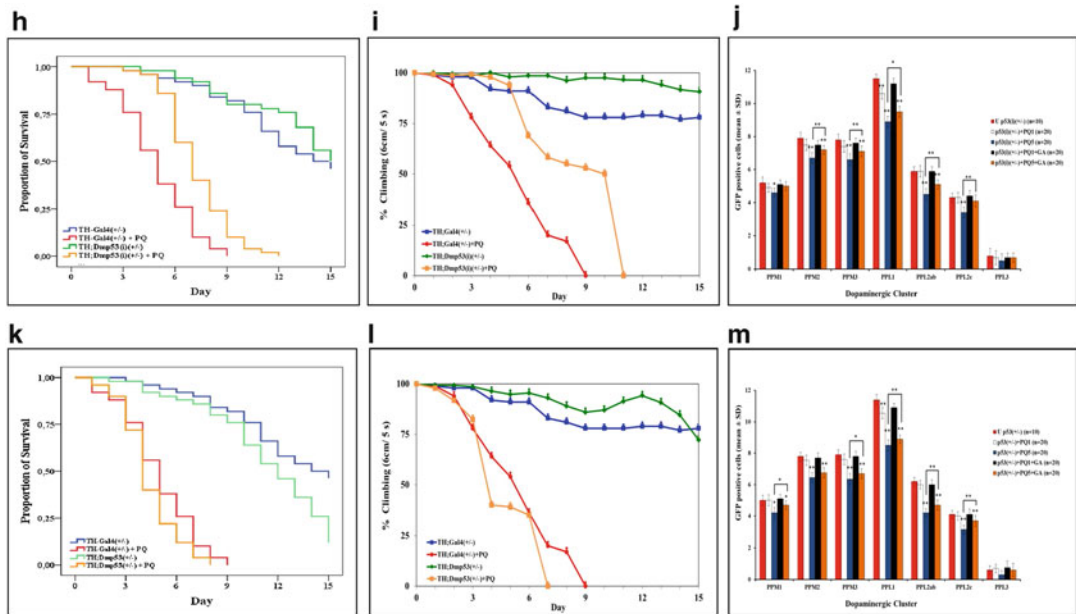


Fig. 2.4 (continued) *U* Test was used to compare differences between two independent groups. Differences were considered statistically significant at $P < 0.05$. ($n=$) represents the number of brain hemispheres examined (except PPM1) per treatment. Error bars indicate SEM. By day 7, PQ (1 mM) significantly eroded only PPL1 DAergic cluster in gene overexpressed/knocked down fly lines compared to untreated flies (**j**, **m**). When flies were fed

with PQ (5 mM), except PPL3, all other DAergic clusters in both overexpressed/knocked down fly lines showed significant differences in the number of DAergic neurons compared with untreated flies. However, overexpressed fly lines showed a decrease in DAergic clusters compared to knocked down fly lines ($p < 0.05$). (**a**, **b**, **h** and **i** are reproduced from ref. Ortega-Arellano et al. (2013) under a Creative Commons attribution-type BY-NC)

for β -neurotoxicity currently available, making it a powerful tool for gene and therapy discovery (Pandey and Nichols 2011).

2.5 *Drosophila's* $\text{A}\beta$ Brain Is Urgently Seeking a Cure: Polyphenols Enter the Stage

Since AD is a complex and multifactorial neurodegenerative condition, an ideal therapeutic agent should have multiple targets, including, antioxidant, metal chelation, anti-aggregation, anti-inflammatory, modulate cell and neurotransmitter signaling, and ultimately have an effect on learning, memory and neurocognitive performance. Polyphenols may comply with such requirements (e.g. Soobrattee et al. 2005; Ramassamy 2006; Weinreb et al. 2004; Rivière et al. 2007; Mandel et al. 2008; Perron and Brumaghim 2009; Kim et al. 2010; Viña et al.

2011; Sharoar et al. 2012; Hirohata et al. 2012; Kostomoiri et al. 2013; Vauzour 2014).

Polyphenols are a group of chemical substances present in plants, fruits, and vegetables, characterized by the presence of one or more than one phenol unit per molecule with several hydroxyl groups on aromatic rings. Based on differences in molecular backbone structure (Fig. 2.5) polyphenols may generally be classified into phenolic acids, flavonoids, stilbenes, and lignans. Flavonoids, the largest group of polyphenols, in turn may be classified into 2 different classes: anthocyanins and anthoxantins. This last group may be classified into flavonols, flavanones, flavones, isoflavones, and flavanols, which give origin to the catechins. In this review, we focused our attention on the effect of polyphenols as antioxidant, metal chelator, anti-amyloidogenic and anti-tau aggregator in *Drosophila*. Given that more than 8,000 polyphenolic compounds

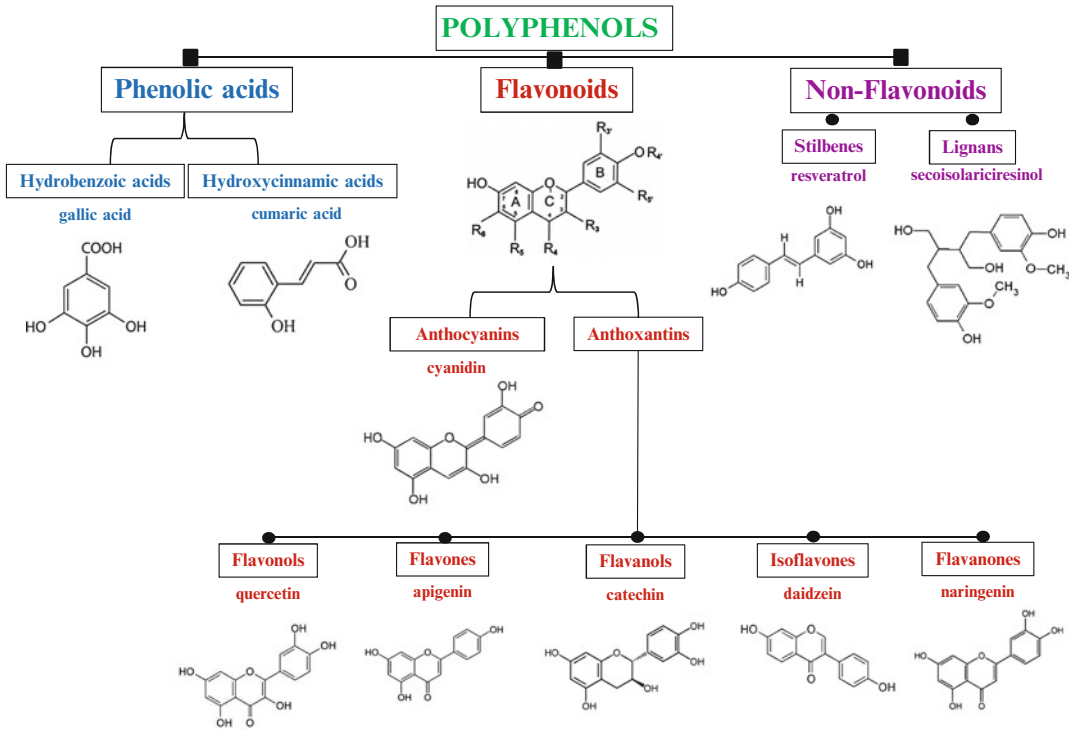


Fig. 2.5 Schematic classification of polyphenols

have been identified in various plant species, we selected a handful of known polyphenols (Fig. 2.5) to be assumed modifiers of AD. Surprisingly, no data is available to establish whether flavonoids (e.g. gallic acid, ferulic acid, caffeic acid, coumaric acid, propyl gallate, epicatechin, epigallocatechin, epigallocatechin gallate) are effective against A β -induced AD-like pathophysiology in *Drosophila*. This despite the fact that they are well-known to be effective as *in vitro* free radical scavengers and antioxidants (Rice-Evans et al. 1997; Sroka and Cisowski 2003; Tsimogiannis and Oreopoulou 2004, 2006; Villaño et al. 2007), *in vitro* inhibitors of A β -aggregation (Porat et al. 2006; Ono et al. 2012; Sato et al. 2013), *in vitro* and *in vivo* metal chelators (Hider et al. 2001; Perron et al. 2010; He et al. 2012; Bonilla-Ramirez et al. 2011), and *in vivo* gene regulatory molecules (Li et al. 2007).

Interestingly, curcumin has been shown to promote amyloid fibril conversion by reducing the pre-fibrillar/oligomeric species of A β , resulting in a reduced neurotoxicity in

Drosophila (Caesar et al. 2012). Recently, it has been shown that structural characteristics, such as degrees of saturation, types of carbon skeleton and functional groups, and hydrophobicity appear to play a role in determining the inhibitory potency of curcuminoids (e.g. diarylalkyls curcumin (CCN), demethoxycurcumin (DMCCN), bisdemethoxycurcumin (BDMCCN)) on BACE-1 activity (Wang et al. 2014). Effectively, it has been found that BDMCCN has the strongest inhibitory activity toward BACE-1. Furthermore, BDMCCN (and CNN) proved not only to be effective in rescuing morphological defects observed in flies expressing APP and BACE-1 in compound eyes (GMR > APP/BACE-1), but also to improve movement in *elav*-BACE-1 and *elav*-APP/BACE-1 female flies. These data suggest that either curcumin (Yanagisawa et al. 2011), curcumin-derived analogs (Orlando et al. 2012) or curcuminoids from rhizomes of *Curcuma longa* might be potential pharmacological agents against AD (Mourtas et al. 2014). It is worth mentioning that ferulic acid, tannic acid and

icariin (a flavonol) have been shown to improve behavioral impairment and Alzheimer-like pathology in transgenic mice (Urano and Tohda 2010; Mori et al. 2012, 2013). We therefore expect that flavonoids, other polyphenols and non-polyphenol molecules might also be operative in transgenic AD flies (Rivière et al. 2007, 2010). This line of research is currently being undertaken in our laboratory (Bonilla-Ramirez et al. 2014).

Since ancient times, traditional medicine has been the only source of drugs to treat diseases including AD (Liu et al. 2014). Today, herbal medicines have the potential to be developed into optimal pharmaceuticals and nutraceuticals for AD because of their multi-function, multi-target characteristics (Kim and Oh 2012; Gao et al. 2013). Recently, it has been shown that a modified version of SuHeXiang Wan (SHXW), a Chinese traditional medicinal extract of 15 crude herbs, improved the developmental defects and motor activity of flies expressing A β 42 in neurons (Hong et al. 2011). Further studies have demonstrated that inhibition of JNK or EGFR/ERK signaling pathways ameliorated the A β -induced lethality and locomotor deficit (Park et al. 2013). Similar to the SHXW, *Gardenia jasminoides Ellis* components and *Gastrodia elata blume* extract have been proved effective in improving short-term-memory in A β transgenic AD flies (Yu et al. 2009, 2012) and in reducing the A β -induced neurodegeneration in *Drosophila* (Ng et al. 2013) (for a review, see Lee et al. 2014). Concomitantly, it has been found that the *Gardenia jasminoides Ellis* fruit extracts displayed antioxidant and superoxide radical-scavenging activities, reducing power, nitrite scavenging activity, inhibition of linoleic acid oxidation activity, superoxide dismutase-like activity and catalase activity (Debnath et al. 2011). Most importantly, the antioxidant and superoxide radical-scavenging activities were highly correlated with the phenolic and polyphenol content. These findings may explain the *Gardenia* extract effect on memory in A β -*Drosophila* model. Another polyphenol-enriched extract, grape-seed polyphenolic extract, improves the eye phenotype in a *Drosophila* model of tauopathy (Pfleger et al. 2010). In

conclusion, there is little doubt that polyphenols (Fraga et al. 2010) and/or natural products (Ansari and Khodaghali 2013) exert several beneficial effects, not least potent antioxidant activity (Galleano et al. 2010) on transgenic A β *Drosophila melanogaster* or PQ-exposed flies (Ortega-Arellano et al. 2011; Jimenez-Del-Rio et al. 2008, 2010).

Notwithstanding, several key questions remain unanswered. Why have antioxidant treatments largely failed in AD? (Persson et al. 2014). Can polyphenol/natural product in vitro actions be translated to benefit animal models and ultimately the human condition? Can polyphenols/natural products retard memory loss in early-onset AD *PSEN 1* E280A presymptomatic patients? Last but not least, are we dealing with an intractable neurological disorder? Further investigations are warranted to address these issues.

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Biflavonoids as Potential Small Molecule Therapeutics for Alzheimer's Disease

3

Arjun Thapa and Eva Y. Chi

Abstract

Flavonoids are naturally occurring phytochemicals found in a variety of fruits and vegetables and offer color, flavor, aroma, nutritional and health benefits. Flavonoids have been found to play a neuroprotective role by inhibiting and/or modifying the self-assembly of the amyloid- β ($A\beta$) peptide into oligomers and fibrils, which are linked to the pathogenesis of Alzheimer's disease. The neuroprotective efficacy of flavonoids has been found to strongly depend on their structure and functional groups. Flavonoids may exist in monomeric, as well as di-, tri-, tetra- or polymeric form through C-C or C-O-C linkages. It has been shown that flavonoids containing two or more units, e.g., biflavonoids, exert greater biological activity than their respective monoflavonoids. For instance, biflavonoids have the ability to distinctly alter $A\beta$ aggregation and more effectively reduce the toxicity of $A\beta$ oligomers compared to the monoflavonoid moieties. Although the molecular mechanisms remain to be elucidated, flavonoids have been shown to alter the $A\beta$ aggregation pathway to yield non-toxic, unstructured $A\beta$ aggregates, as well as directly exerting a neuroprotective effect to cells. In this chapter, we review biflavonoid-mediated $A\beta$ aggregation and toxicity, and highlight the beneficial roles biflavonoids can potentially play in the prevention and treatment of Alzheimer's disease.

Keywords

Alzheimer's disease • Amyloid- β peptide ($A\beta$) • Protein aggregation • Neurotoxicity • Polyphenols • Biflavonoids

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3.1 Alzheimer's Disease

Alzheimer's disease (AD) is the most common neurodegenerative disorder that causes memory loss and cognitive impairment, especially among

the elderly (Selkoe 1991; Hardy and Selkoe 2002; Blennow et al. 2006; Pimplikar 2009). AD is chronic and complex. It is the fourth leading cause of death and accounts for 50–80 % of all dementia cases. Nearly 10 % of individuals above 65 years of age are afflicted with AD and it is believed that about 30 million people world wide are living with this devastating illness (Hamley 2012; Thies and Bleiler 2013). Genetic mutations in three genes that encode the amyloid precursor protein (*APP*), presenilin-1 (*PSEN-1*), and presenilin-2 (*PSEN-2*) proteins, as well as polymorphisms in apolipoprotein E (*APOE*) have been shown to be linked to early-onset (or familial) AD (Hamley 2012; Blennow et al. 2006; Yu et al. 2014). Close to 25 other genes have also been documented and recognized to increase the risk for AD. In addition to genetics, aging, environmental factors, diet that promotes obesity, and an inactive life style are also known risk factors for late-onset (or sporadic) AD, which accounts for over 95 % of all AD cases (Hamley 2012; Blennow et al. 2006). The mechanisms by which these risk factors contribute to AD pathogenesis, however, are largely unknown. In fact, many aspects of AD etiology and disease progression remain unclear. Accumulating evidence indicate that the misfolding and aggregation of the amyloid- β ($A\beta$) peptide, which is a fragment of the APP protein, is a central pathogenic event associated with AD development (Suh 1997; Hardy and Selkoe 2002; Haass et al. 1993). Several reports have also shown that reactive oxygen and nitrogen species, metal ions, and inflammation can be triggers for AD (Nuzzo et al. 2014; Cuajungco et al. 2000; Chauhan and Chauhan 2006; Gu et al. 2008).

The $A\beta$ peptides are generated from the abnormal processing of APP by α -, β - and γ - secretase enzymes (Hamley 2012; Suh 1997; Haass et al. 1993). Although the biological function of APP is still unclear, the protein has been found to be involved in multiple cellular processes, including neuronal growth and survival (Nie et al. 2011; Haass et al. 1993). APP cleavage by β -secretase at methionine 671 position produces a soluble APP fragment and a C-terminal fragment

with 99 amino acids. The C-terminal fragment is further cleaved by γ -secretase at valine 711 and alanine 713 positions to generate $A\beta$ isoforms of 40 or 42 amino acid residues ($A\beta_{40}$ or $A\beta_{42}$) depending upon the cleavage site. These two $A\beta$ isoforms are the major disease-causing peptides (Suh 1997; Hamley 2012; Blennow et al. 2006). Alternatively, α -secretase cleaves APP at lysine 687 and generates a soluble APP fragment and a C-terminal fragment of 83 amino acid residues. The latter fragment is further cleaved by γ -secretase to produce an $A\beta$ (17–42) fragment, which is known as P3 (Suh 1997; Blennow et al. 2006). The P3 fragment does not undergo spontaneous aggregation like the other $A\beta_{40}$ or $A\beta_{42}$ peptides generated by the sequential cleavage of β - and γ - secretases (Suh 1997; Hamley 2012; Blennow et al. 2006). Moreover, the soluble fragment has been described to be neurotrophic and neuroprotective (Nie et al. 2011).

Genetic mutations that affect APP, as in familial AD, have been shown to increase the protein's susceptibility to be cleaved by secretases, resulting in higher rates of $A\beta$ production. The increased $A\beta$ production and subsequent aggregation and accumulation of the peptides into insoluble deposits (or amyloid plaques) over time is believed to cause neurodegeneration that leads to AD (Nie et al. 2011; Hamley 2012; Haass et al. 1993). It is important to note that no mutations in APP are found in sporadic AD, which represents over 95 % of all AD cases, that enhances $A\beta$ production (Hamley 2012; Nie et al. 2011). Regardless of mutation-associated variations in APP processing and $A\beta$ peptide production, the $A\beta$ peptides can self-associate following their release from APP and over time gain pathogenicity. Non-genetic factors that influence AD development include age, oxidative stress, inflammation, and increased levels of metal ions. In fact, aging is recognized as the primary risk factor. Reactive oxygen species (ROS) and metal ions accumulated during aging greatly influence AD progression. ROS has been shown to induce $A\beta$ peptide misfolding and aggregation and enhance $A\beta$ toxicity (Gu et al. 2008; Chauhan and Chauhan 2006; Cuajungco et al. 2000). Accumulating knowledge

suggests that the aggregation and accumulation of A β , as well as the factors that influence A β aggregation, contribute toward AD pathogenesis.

The progressive self-association and deposition of A β peptides that leads to amyloid plaque formation comprise a histopathological hallmark in AD brains (Hardy and Selkoe 2002; Roychaudhuri et al. 2009; Blennow et al. 2006). It is also generally believed that the aggregation process itself is directly involved in the development of AD. However, the mechanism by which the aggregation process occurs and the pathway by which the peptide aggregates cause widespread neuronal death in the brain are still under investigation. Aggregation of A β has been shown to trigger oxidative stress, inflammation, and neurotoxicity resulting in cognitive dysfunction (Hardy and Selkoe 2002). Currently available therapeutics for AD provide moderate and temporary relief of symptoms (Blennow et al. 2006). A significant number of clinical trials for AD have been unsuccessful. The lack of a clear understanding of the molecular mechanisms of A β misfolding and aggregation is one of the major challenges and an obstacle in AD drug discovery (Uversky 2010; Hardy and Selkoe 2002).

3.2 A β -Peptide Toxicity in Alzheimer's Disease

It is now well accepted that A β peptide production and subsequent aggregation is a major causative factor for AD pathogenesis (Hardy and Selkoe 2002; Pike et al. 1991; Roychaudhuri et al. 2009). Although various N- or C-terminus truncated A β peptide fragments have been found in the brains of AD patients, A β 40 and A β 42 are the major isoforms present (Roychaudhuri et al. 2009; Suh 1997). The most abundant isoform of the A β peptide found in AD patients is A β 40, which is less toxic and forms fibrils more slowly compared to A β 42. The presence of the two additional amino acids in A β 42, isoleucine and alanine, increases the peptide's aggregation propensity and neuronal toxicity. The production of A β 42 has been found to be enhanced by

genetic mutations linked to the familial form of AD (Findeis 2007; Blennow et al. 2006).

The native A β peptides are unstructured monomers with a molecular mass of 4 kDa. A β peptide is rich in hydrophobic amino acid residues and is amphiphilic. The C-terminus and particularly the central part of the peptide contain highly hydrophobic residues, whereas the N-terminus contains hydrophilic and charged residues. More than half of the amino acid residues in A β peptides are hydrophobic and are prone to aggregation (Walsh et al. 1999; Dasilva et al. 2010). A β peptides spontaneously undergo conformational changes from unstructured monomers to form misfolded intermediates. The misfolded A β species are highly unstable and undergo further aggregation to form higher-order oligomers, protofibrils, and fibrils that are rich in β -sheet structures (Walsh et al. 1999; Dasilva et al. 2010). A β fibrils consist of 5–6 strands of protofibrils bundled in parallel. The aggregation of A β peptides is fundamentally associated with its neurotoxic effects and has been found to trigger oxidative stress and inflammation (Walsh et al. 1999; Chauhan and Chauhan 2006; Pimplikar 2009; Nuzzo et al. 2014). *In vitro* and *in vivo* studies have shown that soluble globular A β oligomers consisting of 3 to 24 monomers (A β derived diffusion ligand (ADDL)) and curvilinear, beaded oligomers measuring 3–6 nm in diameter and 200 nm in length (protofibrils) formed during the early phases of the A β aggregation process are the potentially toxic A β species, rather than unstructured monomers or the aggregated end-product, fibrils (Walsh et al. 1999; Klein 2002). In fact, the A β aggregation process is a highly complex and heterogeneous one (Uversky 2010). Distinctly different aggregation behaviors of the two A β isoforms have been found (Chen and Glabe 2006). Although the structures of A β aggregates and mechanism of A β aggregation remain unclear, the steric packing of the hydrophobic residues and/or π -stacking of aromatic amino acid side chains are believed to be the underlying driving forces for A β aggregation (Armstrong et al. 2011; Gazit 2002).

3.3 Therapeutic Strategies for Alzheimer's Disease

As discussed above, increasing evidence suggests that A β aggregation is a primary cause of AD pathogenesis (Walsh et al. 1999; Chromy et al. 2003). Additionally, excessive generation of reactive oxygen and nitrogen species and inflammation are found in the AD brain (Gu et al. 2008; Butterfield et al. 2001). Therapeutic targets for AD remain elusive. A number of clinical trials for AD have been launched over the years and so far, results have been discouraging. It is clear that a lack of understanding of the molecular mechanisms of AD, in particular, a poor understanding of A β aggregation, is a major obstacle in AD drug development. There is also a lack of suitable animal models that recapitulate all of the pathological events seen in human AD brains, as well as a lack of biomarkers to detect and monitor the progression of AD (Ghosh et al. 2012; Jay et al. 2011; LaFerla and Green 2012). However, significant progress has been made in advancing our understanding of AD mechanisms. *In vitro* and *in vivo* studies have indicated that the excessive generation of A β peptides and subsequent A β aggregation primarily induce A β neurotoxicity. Several studies indicate that lowering A β production and stabilizing its native form, inhibiting A β aggregation, and clearing A β pool are promising therapeutic strategies for AD (Soto et al. 2000; Blennow et al. 2006; Permann et al. 2002). Drugs that can alleviate oxidative stress and inflammation are also potential therapeutics for AD. Below, we briefly summarize the range of therapeutic strategies that have been investigated.

3.3.1 α -, β -, and γ - Secretase Modulators

Generation of A β peptides is at the beginning of a cascade that leads to AD. α -, β -, and γ -secretases are aspartic acid proteases that preferentially cleave APP (Haass et al. 1993; Ghosh et al. 2012). As discussed previously, β - and γ -secretases are the principal players involved in

A β peptide production, while α -secretase cleavage of APP has been found to prevent A β deposition. Inhibitors that target β - and γ -secretases as well as activators of α -secretase have therefore been considered promising candidates for treating AD.

The amyloidogenic processing of APP is initiated by β -secretase cleavage, which results in the release of a large soluble fragment and a membrane-tethered C-terminal fragment (Haass et al. 1993; Hardy and Selkoe 2002). The second cleavage in the production of A β peptide is mediated by γ -secretase, which cuts the membrane fragment within the transmembrane domain to release A β into the extracellular milieu (Pimplikar 2009; Hardy and Selkoe 2002). The cleavage mechanism and inhibition pathways of β -secretase that can reduce the rate of A β production are largely known. A study that characterized the structure and catalytic properties of β -secretase suggests that a large inhibitor molecule is required to fit into the relatively large-sized active site of the enzyme (Nie et al. 2011; Ghosh et al. 2012). For drug development, small molecule therapeutics with high molecular weights are expected to have poor blood-brain barrier (BBB) permeability, thus making β -secretase inhibitors generally unsuitable as drugs. The orally available drug LY2886721 has been shown to inhibit β -secretase activity, reduce brain A β production, and rescue the cognitive decline in transgenic AD mice (Lahiri et al. 2014; Nie et al. 2011). Unfortunately, the drug failed phase II clinical trials due to liver dysfunctions in patients (Lahiri et al. 2014).

γ -secretase is the second enzyme involved in the amyloidogenic processing of APP to generate A β peptides. The enzyme is a membrane-embedded protease complex that cleaves the transmembrane region of APP at multiple sites due to its heterogeneous site preference. This second cleavage of APP generates A β peptides of variable length (38–43 amino acids long). As discussed previously, the A β 42 isoform is prone to oligomerization, aggregation, and fibril formation, all of which are considered critical in the development of AD pathology. A number of peptides and small molecules inhibitors of the

γ -secretase complex that block the active site or interact with the initial substrate docking site have been studied (Ghosh et al. 2012; Nie et al. 2011). The γ -secretase inhibitor semagacestat was discovered to reduce A β production in cell culture and also *in vivo*. However, the inhibitor was not successful in a phase III clinical trial (Nie et al. 2011; Doody et al. 2013). Semagacestat was effective at reducing A β peptide levels in the plasma but not in the cerebrospinal fluid. Also, the compound did not improve cognitive functions in patients (Doody et al. 2013). γ -secretase has several physiological substrates, including the Notch protein. Because Notch proteins perform a variety of cellular functions, the inhibition of γ -secretase activity may also lead to other deleterious physiological consequences (Nie et al. 2011; Ghosh et al. 2012; Blennow et al. 2006). Recent studies have shown that small molecules that modulate β - and γ -secretase activities and selectively alter A β production without altering signal transduction may be a suitable strategy for AD intervention (Ghosh et al. 2012; Nie et al. 2011).

In contrast to the amyloidogenic processing of APP by β - and γ - secretases that generate A β peptides, the non-amyloidogenic processing of APP is initiated by α -secretase, which cuts APP 16 amino acids downstream of the β -secretase cleavage site. The cleavage releases a soluble fragment and a truncated transmembrane fragment that is then cleaved by γ -secretase. The truncated A β peptides produced are non-amyloidogenic and have been shown to be neurotrophic and neuroprotective. Thus, stimulating α -secretase activity appears to be a direct gain-of-function strategy and may be a safe route to prevent AD. However, there is also evidence that α -secretase cleaves other membrane proteins aside from APP (Nie et al. 2011; Blennow et al. 2006).

It is clear that modulating α -, β -, and γ -secretase activities may be a promising strategy for preventing and treating AD. However, efforts so far in finding β - and γ - secretase inhibitors and α -secretase activators have not been successful. A more detailed understanding of the underlying mechanisms and the complex interplay with other biochemical processes are needed for identifying

novel and specific inhibitors or activators necessary for AD therapeutic development.

3.3.2 Immunotherapy

The accumulation of A β peptides in the brain is dependent upon a balance between its deposition (production and aggregation) and degradation (phagocytosis and proteasomal degradation) (Blennow et al. 2006; Hardy and Selkoe 2002). Antibodies that target A β peptides and activate the immune system can result in the induction or suppression of immune responses that enhance A β clearance. The injection of A β peptides as antigens (active immunization) or the injection of preformed antibodies against A β (passive immunization) have been considered as viable therapeutic strategies for treating AD (Delrieu et al. 2012; Blennow et al. 2006). Antibodies can clear A β by binding to the peptide in the plasma and sequestering them or by binding to A β in the brain for phagocytosis. Immunization with A β or A β antibodies have been shown to successfully reduce AD pathology and improve cognitive functions in AD mouse model (Rasool et al. 2013). However, although the active immunization with a mixture of A β 42 and the adjuvant QS-21 (AN1792 A β vaccine) has been shown to be effective at clearing amyloid senile plaques in the human brain, the immunization did not improve cognition. Rather, AN1792 was found to induce autoimmune encephalitis (Nie et al. 2011; Delrieu et al. 2012). The failure of phase II clinical trials due to the development of meningoencephalitis in patients has raised serious concerns about the therapeutic strategy of active immunization for the treatment of AD.

3.3.3 Anti-oxidative Therapy

A number of studies have shown that the AD brain shows increased oxidative stress (Butterfield et al. 2001; Chauhan and Chauhan 2006). Reactive oxygen species (e.g., superoxide anions, hydrogen peroxides, hydroxyl and peroxy radicals, and singlet oxygen) and reactive ni-

trogen species (e.g., nitric oxide) produced during oxidative stress are capable of damaging proteins, DNA, and lipids that subsequently result in neuronal death. A number of factors can cause neurons to be particularly vulnerable to free radical attacks. First, oxygen is abundant in the brain as the brain requires a large amount of oxygen for metabolic activities. Second, the neuronal membrane consists of high proportions of easily oxidizable polyunsaturated fatty acids. Third, neurons contain low levels of the natural anti-oxidant glutathione. Fourth, increased levels of metal ions are found in the AD brain that can potentially generate free radicals and enhance protein and DNA oxidation. Moreover, the presence of heme oxygenase-1 and superoxide dismutase-1 (anti-oxidative enzymes) in senile plaques highlight the possible role of oxidative stress in AD (Chauhan and Chauhan 2006; Butterfield et al. 2001; Zhou et al. 2014). The fact that aging is a key risk factor for AD and that reactive oxygen species accumulate over time provides additional support for the free radical hypothesis of AD pathogenesis. Therefore, anti-oxidants taken as supplements, such as vitamin E, vitamin C, coenzyme Q10, and β -carotene, have been tested to prevent and treat AD. However, the supplements have been found to be ineffective at reducing AD symptoms (Yang et al. 2010). Some studies have shown that A β peptides can induce the formation of free radicals while others have shown that A β peptides may serve as anti-oxidants and compensate for the increased oxidative stress in cells. Tissue injury by itself can generate reactive oxygen species. Thus, although there is increasing evidence that oxidative stress is involved in AD, the underlying link between the two is still not properly understood.

3.3.4 Anti-inflammatory Therapy

Aside from the two major known neuropathological AD hallmarks, senile plaques and neurofibrillary tangles, a number of pathological abnormalities, including microglial activation and inflammatory processes, are found in AD brains (Nuzzo et al. 2014; Zhou et al. 2014;

Kim et al. 2008). Chronic inflammatory reactions are also common symptoms among age-related neurodegenerative disorders. Inflammation is a natural immune response linked to the excessive activation of the glial cells during fighting against harmful substances or injuries. An abundant number of activated glial cells such as microglia and astrocytes are characteristically found around neurons and amyloid plaques. Inflammation mediator molecules such as cytokines, chemokines, macrophages, prostaglandins, coagulation factors, and reactive oxygen species have been detected in the AD brain (Kim et al. 2008). Genetic studies have shown that polymorphisms in some genes that encode for inflammatory interleukins 1 α and 1 β enhance the risk of AD (Kim et al. 2008; Nuzzo et al. 2014). These findings suggest chronic inflammation may play an important role in AD pathogenesis. However, anti-inflammatory drugs (NSAIDs) such as naproxen and celecoxib have been shown to be ineffective in reducing AD pathology and symptoms (Lyketsos et al. 2007). Recent evidence suggests that NSAIDs may lower A β load by inhibiting the small GTP binding protein Rho and its effector, Rho-associated kinase (Rock), instead (Gasparini et al. 2004; Blennow et al. 2006).

3.3.5 Metal Chelators

Studies have shown that APP and A β peptides can interact with and bind to heavy metal ions, including copper, iron, aluminum, and zinc ions. The binding can subsequently induce the misfolding and aggregation of A β peptides (Hane et al. 2013; Cuajungco et al. 2000). In fact, elevated levels of such metal ions have been found in the brains of AD patients, as well as in elderly individuals (Cuajungco et al. 2000). Monomeric and oligomeric A β have been shown to bind to metal ions, which give rise to ordered A β structures. Therefore, metal chelation has been explored as one of the therapeutic strategies for reducing A β -metal ion interactions by removing metal ions in the AD brains. Clioquinol, a copper-zinc chelator, is one of the FDA approved metal chelators for

AD (Schimmer et al. 2012). However, its use has been discontinued due to its adverse effects, including neurotoxic effects in some patients. It remains to be understood which form, in terms of conformational and aggregation state, the A β peptide adopts after the metal ions are removed from A β aggregates. Moreover, the delivery of metal chelating compounds to the A β enriched areas of the brain remains a challenging task (Blennow et al. 2006; Nie et al. 2011).

3.3.6 Inhibitors and Modulators of Neurotransmitters

Neurotransmitters (neurochemical messengers such as acetylcholine) are signaling molecules involved in several aspects of normal brain functioning including cognition, memory, and learning. The dysregulation of the biosynthesis of acetylcholine neurotransmitter has been found in AD (Brogi et al. 2014; Rosini et al. 2014). Anti-acetylcholinesterases, which inhibit the action of the acetylcholinesterase enzyme that breaks down acetylcholine and increase the level, duration, and action of acetylcholine, have been suggested to improve learning and memory. Examples of these inhibitors include galantamine, remynyl, razadyne, rivastigmine, and donepezil (Hamley 2012; Blennow et al. 2006). Anti-acetylcholinesterases have also been found to protect cells from free radical toxicity and A β -induced toxicity. In fact, cholinesterase inhibitors have been used as a first line medication for AD as they have been shown to be effective at delaying cognitive impairment. Elevated levels of another neurotransmitter, glutamate, that activates *N*-methyl-D-aspartate (NMDA) receptors and mediate excitatory signals, have also been found in AD (Johnston et al. 2006). Additionally, high concentrations of the neurotransmitter γ -aminobutyric acid (GABA), which mediates inhibitory signaling in the reactive astrocytes of the dentate gyrus, have been found. A correlation between high concentrations of GABA and deteriorations in learning and memory capacity has been shown (Wu et al. 2014). Memantine is one of the

available AD prophylactic medications that block NMDA receptors. Like other inhibitors that reduce AD symptoms, the inhibitors or modulators of neurotransmitter release are not free from side effects (Blennow et al. 2006). Dizziness, headaches, nausea, vomiting, diarrhea, loss of appetite, and weight are some of the common side effects.

3.3.7 A β -Based Peptide Inhibitors

Given that A β aggregation is critical in AD development, the inhibition of A β aggregation is another AD therapeutic strategy that has been explored (Estrada and Soto 2007; Funke and Willbold 2012). Mutational and computational studies have revealed that residues in the hydrophobic C-terminal half of A β , leucine, isoleucine, phenylalanine, and aromatic amino acids, are critical for β -sheet formation in A β aggregates (Gazit 2002; Porat et al. 2006). Moreover, not only do the full-length A β peptides aggregate, truncated fragments containing the hydrophobic segment can also self-associate. Hence, A β peptides segments that can bind to the hydrophobic region of the full-length A β peptide and prevent A β aggregation have been considered and tested as alternative therapeutics to prevent AD (Estrada and Soto 2007; Mason et al. 2003). Specifically, a pentameric core segment of A β , A β (16–20) with the amino acid sequence of KLVFF, has been tested as a drug lead because it is able to bind to the full-length A β and prevent the peptide's assembly into fibrils (Pallitto et al. 1999). Soto and coworkers introduced a proline-containing A β (17–21) segment, which has an amino acid sequence of LVFFA, and showed that the segment reduced the propensity of full length A β peptides to form β -sheets (Poduslo et al. 1999; Estrada and Soto 2007). The peptide fragment LVPFF has also been found to inhibit A β aggregation and disrupt preformed fibrils (Mason et al. 2003). These β -sheet breaker peptides are believed to be small and hydrophobic enough to cross the BBB. Based on the A β (15–25) sequence, Murphy and co-workers designed a novel peptide series with a 'recognition element' homologous to A β and a

'disrupting element' fused to the C terminus of the peptide (Pallitto et al. 1999). It was shown that a minimum of three lysines is required to disrupt A β aggregation. The peptide KLVFFK β reduced A β toxicity by accelerating A β aggregation kinetics and altering fibril morphology. The study thus demonstrated that A β aggregation does not need to be blocked in order to prevent A β toxicity. Compounds that increase the aggregation kinetics to form A β fibrils at faster rates can be protective as A β fibrils are less toxic than A β oligomers (Blennow et al. 2006). However, proteolysis and short *in vivo* half-life of peptide are the major obstacles associated with this class of A β aggregation and toxicity inhibitors.

3.3.8 Aromatic Small Molecules

Based on the observation that aromatic amino acids are abundant in aggregation-prone proteins and peptides, including A β peptides, Gazit and co-workers proposed that the π - π stacking of the aromatic amino acid side chains stabilizes the fibrillar conformation of A β and is a driving force for amyloid fibril formation (Armstrong et al. 2011; Reinke and Gestwicki 2007; Gazit 2002). In their study, a series of synthetic as well as naturally occurring compounds, including curcumin, resveratrol, apigenin, apomorphine and Congo Red, that are capable of disrupting π - π interactions were tested for their ability to inhibit A β fibril formation and toxicity (Gazit 2002; Porat et al. 2006). Results from the study show that compounds containing at least two aromatic structures (phenyl rings) and three hydroxyl groups linked with two to six atom linkers were effective anti-amyloid compounds (Porat et al. 2006). Two other studies have also shown that the numbers of phenyl rings and hydroxyl groups, as well as their positions, on the molecules are important characteristics that determine the effectiveness of the aromatic compounds in their inhibition of A β aggregation (Lakey-Beitia et al. 2014; Begum et al. 2008).

One class of aromatic compounds, flavonoids, has garnered increasing interest in the prevention and treatment of AD. Flavonoids are plant

secondary metabolites and are a subclass of the naturally-occurring polyphenols. Flavonoids have long been known for their multi-faceted pharmacology, including anti-inflammatory, anti-oxidant, anti-tumorigenic, anti-cancer, anti-microbial, and anti-vascular properties (Ringman et al. 2005; Lakey-Beitia et al. (2014); Kang et al. 2005). The diverse range of biological properties appears to stem from the ability of this class of compounds to bind to a multitude of cellular targets, and thereby regulate multiple cellular pathways (Ringman et al. 2005; Ramassamy 2006). Moreover, the beneficial effects of flavonoids in neurodegenerative diseases have been documented (Ganguli et al. 2000; Gao et al. 2012), although understanding the molecular mechanisms by which flavonoids exert their biological effects remains an active research area. We summarize below the known beneficial roles of flavonoids in AD, with an emphasis on dimeric flavonoids, the biflavonoids. As flavonoids are naturally occurring and an integral part of a diet rich in fruits and vegetables, they are safe, with little or no adverse side effects, and abundant. Additionally, flavonoids are able to target multiple risk factors of AD, including inflammation, oxidation, A β aggregation, and A β toxicity, these compounds are particularly promising therapeutics for both the prevention as well as treatment of AD.

3.4 Beneficial Roles of Naturally Occurring Polyphenolic Compounds in AD

Polyphenols comprise naturally occurring phenyl ring-containing small molecules found in fruits and vegetables and have been shown to offer beneficial effects in AD and other neurodegenerative disorders (Ringman et al. 2005; Lakey-Beitia et al. 2014; Kang et al. 2005). Although their mode of action and targets in these diseases are not yet clear, the phenyl ring structure is believed to be central to polyphenol's beneficial effects (Yang et al. 2005; Gazit 2002). Polyphenols perform a myriad of biological functions to improve the health of the organisms and ac-

tivate a multitude of pharmacological pathways (Gauci et al. 2011; Wang et al. 2001; Ramassamy 2006; Hamaguchi et al. 2009). More than 8,000 naturally occurring polyphenols, ranging from simple small molecules to highly polymerized compounds, have been identified and characterized. The compounds are efficiently absorbed in the intestine and do not have any adverse side effects (Ringman et al. 2005).

A number of pathways have been identified by which polyphenols may exert their protective role in AD. The compounds are capable of efficiently chelating heavy metal ions (Leopoldini et al. 2006; Yang et al. 2005), thereby reducing a primary source of free radicals in neurons that contribute to oxidative stress (Cuajungco et al. 2000). In addition to being chelating agents, polyphenols are also antioxidants as the hydroxyl groups are capable of consuming free radicals and terminating redox reactions that cause cellular damages. The polyphenols myricetin and quercetin have been found to alleviate inflammation, which is also one of the features of AD (Lee and Choi 2010; Boots et al. 2011). The polyphenols curcumin, epigallocatechin gallate, and resveratrol have been found to effectively reduce A β toxicity *in vitro* and *in vivo* by multiple mechanisms (Yang et al. 2005; Feng et al. 2009). Strikingly, the beneficial effects of these compounds have been found to correlate with the delay in the onset of AD and other aging related diseases in the elderly. In addition to their protective role in AD, polyphenols such as quercetin have also been reported to be anti-microbial, exerting toxicity towards many different strains of pathogenic bacteria (Su et al. 2014). Polyphenols from black tea, e.g. catechins and theaflavin, also possess anti-fungal activities (Betts et al. 2013). Furthermore, the polyphenols apigenin, kaemferol, and catechin have been found to exhibit anti-viral and anti-cancer properties (Zhang et al. 2004; Pang et al. 2014).

Aside from the beneficial pharmacological effects described above, polyphenols also interfere with the π - π stacking between aromatic residue side chains of A β peptides, suggesting that these compounds may affect A β aggregation as well (Re et al. 2010; Gazit 2002). The presence of phenyl groups and internal double bonds, the

ability to form hydrogen bonds, and molecular planarity are some of the other features of the polyphenols that may be involved in modulating A β aggregation. A study based on the polyphenol curcumin's structure-function relationship have indicated that the two terminal phenyl groups, the substituted aromatic end groups that participate in hydrogen bonding, and an optimal rigid linker length of 6–19 Å, are critical structural features that inhibit A β aggregation (Yang et al. 2005). It is also suggested that a flat and planar molecule with two aromatic end groups and substituted aromatic groups are suitable ligands for A β peptides and may be neuroprotective (Yang et al. 2005; Reinke and Gestwicki 2007).

Flavonoids are a class of polyphenolic compounds with a C6-C3-C6 backbone (Fig. 3.1a) that adopt a flat and planar structure (Thapa et al. 2011; Re et al. 2010). Flavonoids consist of two aromatic rings (α and β) linked with a pyran ring (γ) and are the largest group of polyphenols available from dietary fruits and vegetables (Lakey-Beitia et al. 2014; Re et al. 2010). The chemical structures of flavonoids are quite diverse. Based on the oxidation state of the pyran ring, flavonoids are classified into flavones, flavonols, flavanols, isoflavonoids, flavanones, and anthocyanins (Lakey-Beitia et al. 2014; Re et al. 2010). Hydroxylation of flavonoids occurs mainly at the C5, C7 and C4' positions (Fig. 3.1a). Flavones such as apigenin and luteolin are rich in parsley and celery. Flavonols possess a keto-hydroxypyrene group, which is hydroxylated at the C3, C5 and C7 positions (Fig. 3.1a). Myricetin and quercetin are examples of flavonols that are found in onions and broccoli. Flavanols have a hydroxylated pyran ring mostly at the C3 position and are present in green tea and red wine. Soy product flavonoids such as genistein and daidzein are categorized as isoflavonoids, which have a substituted keto group on a pyran ring structure. The flavanone group consisting of hesperetin and naringenin has substituted keto-pyran groups on ring α at C3 and C5 positions and are abundant in citrus and tomatoes. Anthocyanin found in berries offers the colors in the plant and fruits. Non-flavonoid types of polyphenols include: (i) phenolic acids (e.g. caffeic acid and gallic acid

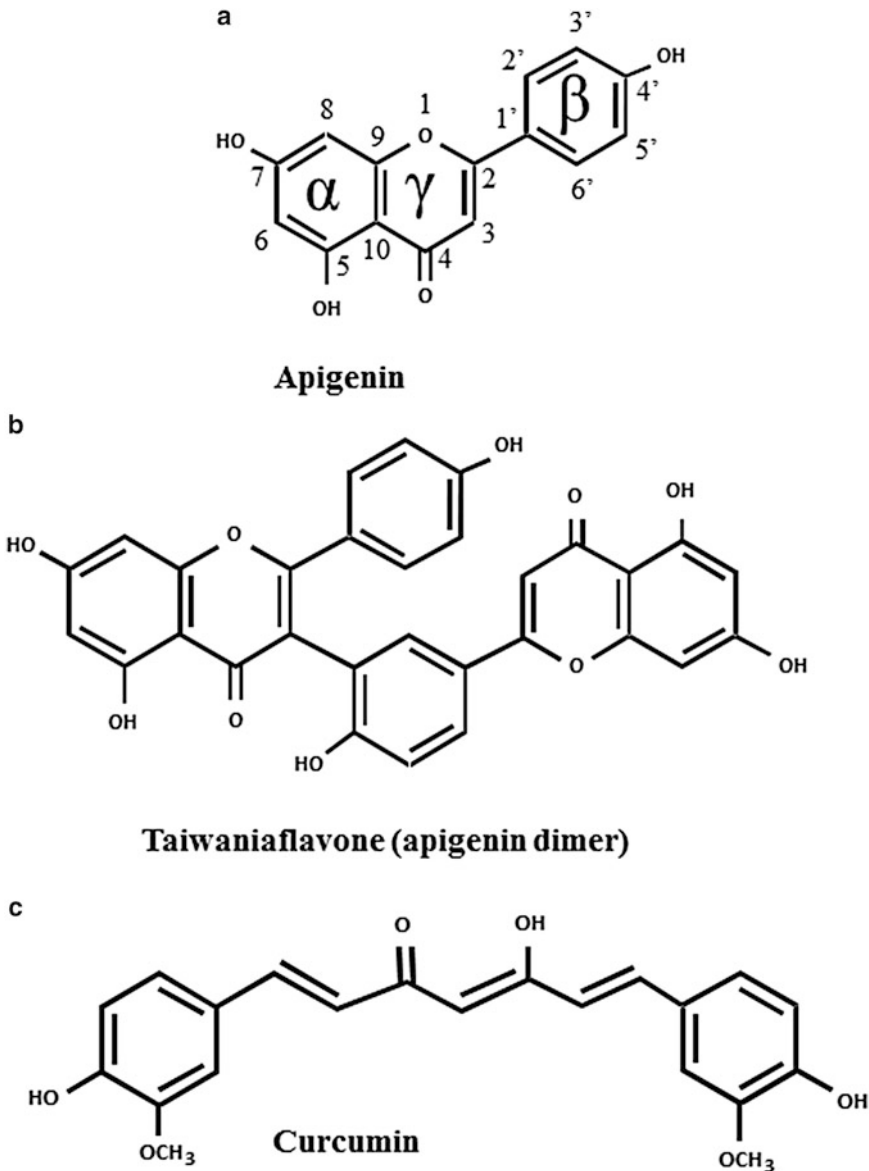


Fig. 3.1 Schematics of monoflavonoid apigenin (a), biflavonoid taiwaniaflavone (apigenin dimer, apigenin-3 and 3'''-apigenin) (b), and curcumin (c). Phenol rings are

denoted as α -, β -, and γ (Reprinted with permission from Thapa et al. 2011)

etc.), (ii) lignans (e.g. secoisolariciresinol), (iii) stilbenes (e.g. resveratrol), and (iv) curcuminoids (e.g. curcumin, Fig. 3.1c) (Lakey-Beitia et al. 2014).

As a subclass of polyphenols, flavonoids exhibit the same wide range of biological functions, including anti-oxidative, anti-inflammatory, anti-tumorigenic, and anti-

amyloidogenic activities (Ramassamy 2006; Lakey-Beitia et al. 2014). Recent studies have shown that larger flavonoid compounds that contain multiple chemical moieties, for example, dimers of single flavonoid structures or biflavonoids, show enhanced biological activities compared to their monomeric flavonoid counterparts (Zhang et al. 2004; Wong et al.

2007; Dhawan et al. 2002). For example, the flavonoid taiwaniaflavone (Fig. 3.1b) is a dimer of the flavonoid apigenin (Fig. 3.1a). Apigenin is a known anti-cancer agent with moderate activity in multidrug resistance breast cancer (Zhang et al. 2004). Dimers of apigenin linked with ethylene glycol linker showed enhanced anti-cancer effect compared to apigenin in multidrug resistance breast cancer cells as well as in leukemia cell lines (Zhang et al. 2004). It was suggested that dimeric molecules could bind to two distinct sites on a single molecule or a defined site on two separate molecules (Hadden and Blagg 2008). With respect to A β peptide aggregation and toxicity, a dimeric polyphenol, curcumin (Fig. 3.1c), has been shown to potently inhibit A β aggregation and toxicity and improve cognitive function (Yang et al. 2005). Studies on curcumin also indicated that a flat and planar shaped polyphenol molecule with two aromatic end groups and substituted aromatic groups may be necessary for inhibiting A β aggregation and toxicity. This conclusion was further supported by the observation that ferulic acid, which is structurally similar to one-half of curcumin, does not inhibit A β aggregation. However, a dimeric form of ferulic acid effectively inhibits A β aggregation (Byeon et al. 2007). The polyphenol that contains both ferulic acid and styryl benzene has been shown to be more effective than either of the two individual compounds (Lee et al. 2005). It has been further suggested that the dimers of A β binding monoflavonoids, separated by an appropriate size linker, may serve as a general structure for inhibitors of A β aggregation and toxicity (Thapa et al. 2011). Such biflavonoid compounds may be promising therapeutic candidates for AD.

3.5 Beneficial Roles of Biflavonoids in Alzheimer's Disease

As discussed earlier, biflavonoids are flavonoid-flavonoid dimers connected by a C-C or a C-O-C linkage (Fig. 3.1b). Naturally occurring biflavonoids contain different combinations

of monoflavonoids, such as flavone-flavone, flavone-flavonol, or flavanone-flavone (Lakey-Beitia et al. 2014; Kim et al. 2008). Biflavonoids are abundant in the *Ginkgo biloba*, *Selaginella*, and *Garcinia* plant species. This subclass of polyphenols exhibits multiple biological and pharmacological properties stemming from the different types of chemical substitutions, including hydroxyl and methoxyl groups. The anti-microbial action of the biflavonoid amentoflavone and its synergistic effect with antibiotics have been documented (Hwang et al. 2013). Amentoflavone stimulates mitochondrial dysfunction and induces apoptotic cell death in yeast (Hwang et al. 2012). The biflavonoid ginkgetin is cytotoxic to human ovarian adenocarcinoma and is also effective against the flu virus (Kim et al. 2008; Miki et al. 2007). We review below the potent anti-oxidant, anti-inflammatory, anti-acetylcholinesterase, β -secretase inhibiting, and anti-amyloidogenic properties of biflavonoids that have been reported to contribute to their beneficial effects in AD.

3.5.1 Biflavonoids as Anti-oxidants

Free radicals are among the normal cellular metabolic byproducts but their accumulation over time is undesirable as they contribute to oxidative stress. Free radicals attack proteins, lipids, and nucleic acids and disrupt their structures and functions. One of the best-known properties of flavonoids is their anti-oxidative capability. Virtually all flavonoids are able to scavenge reactive oxygen species and terminate the detrimental redox reactions. The biflavonoid morelloflavone is known as a potent lipid-peroxidation inhibitor (Gil et al. 1997). It is believed that biflavonoids could more efficiently neutralize free radicals compared to monoflavonoids due to the presence of the additional hydroxyl groups. Further, flavonoids are capable of removing metal ions bound to proteins and lipids; for example, the flavonoid quercetin has been found to chelate heavy metals (Leopoldini et al. 2006). The use of naturally occurring flavonoids, particularly biflavonoids,

for chelating metal ions and scavenging free radicals appears to be an attractive approach to prevent AD.

3.5.2 Biflavonoids as Anti-inflammatory Agents

Brain inflammation is another common feature of AD. The stimulation of pro-inflammatory cytokines, such as interleukin-1 and interferon- γ , results in the production of large amounts of reactive nitric oxides by inducing nitric oxide synthase (iNOS), which could lead to neuronal death (Nuzzo et al. 2014; Pokharel et al. 2006). Inflammation mediators such as prostaglandins that are involved in chronic inflammation are metabolic products of cyclooxygenase (COX) and the pharmacological inhibition of COX and iNOS results in the relief of pain and inflammation. Biflavonoids such as amentoflavone, taiwaniaflavone, bilobetin, and ginkgetin have been found to down regulate iNOS and COX-2 expression, and are known to regulate pro-inflammatory molecules at the transcription level (Pokharel et al. 2006; Kim et al. 1998, 2008).

3.5.3 Biflavonoids as Secretase Inhibitors

Flavonoids have also been viewed as an attractive class of molecules to limit A β production by altering the activities of secretases. The monoflavonoid myricetin has been shown to reduce A β production in cell-free systems as well as in neuronal cells (Shimmyo et al. 2008a). They appear to reduce A β production by upregulating α -secretase activity and/or down regulating β -secretase activity (Shimmyo et al. 2008a, b). For reasons that are still unclear, other monoflavonoids that are structurally similar to myricetin such as apigenin and morin were found to be less effective at reducing β -secretase activity. On the other hand, a study on β -secretase inhibition has revealed that amentoflavone-type biflavonoids (molar mass 600 Da) potently inhibited β -secretase activity (Sasaki et al.

2010). It is believed that biflavonoids, being roughly twice the size of monoflavonoids, better fit the large sized β -secretase catalytic cleft to inhibit β -secretase activity compared to smaller sized monoflavonoids (Ghosh et al. 2012). More recently, the polyphenol tannic acid, a naturally occurring non-flavonoid polymer (molar mass 1,700 Da) was found to inhibit β -secretase activity, prevent cognitive impairment, and reduce AD-like pathology in transgenic mice (Mori et al. 2012). However, due to its large size, tannic acid is impermeable to the BBB (molecular weight limit for BBB is believed to be 600 Da). From these studies, it appears that intermediate-sized biflavonoid molecules may be effective and BBB-permeable candidates for modulating secretase activity.

Although the flavonoids that have exhibited beneficial effects in various aspects of AD are multifunctional and have been shown to reduce oxidative-stress, inflammation, and A β production, the roles they play in the disease process and the interplay between these effects in AD remain to be explored. For instance, some studies have shown that oxidative stress and inflammatory processes are secondary events in AD. As the vast majority of AD cases are sporadic rather than familial, it is reasonable to conclude that A β aggregation rather than A β production is the primary pathogenic event in AD (Pimplikar 2009; Walsh and Selkoe 2007; Chromy et al. 2003). Modulating secretase activities that reduce A β production, therefore, is expected to have only a limited effect on A β aggregation. Removal of the triggers of A β aggregation, e.g., oxidative stress, or direct disruption of the aggregation process itself, are also needed in order to achieve complete inhibition of A β aggregation, and thereby AD pathogenesis.

3.5.4 Biflavonoids as A β Aggregation Inhibitors

Studies that have investigated small molecule inhibitors of A β aggregation suggest that the inhibitor should strongly interact with A β (Nie et al. 2011). Additionally, the inhibitor should

have sufficient length and flexibility, which would enable them to capture the entire A β peptide and accommodate conformational changes in the target (Nie et al. 2011). Thus, an effective A β -aggregation inhibitor should contain moieties that strongly interact with A β , be of appropriate size, and exhibit suitable flexibility. In this regard, naturally occurring biflavonoids containing two or more flavonoid units may be a better option to target A β aggregation as compared to simpler polyphenols and flavonoids. It is possible that small polyphenols, including monoflavonoids, may not produce sufficient steric hindrance as necessary to interact with and inhibit A β aggregation as compared to biflavonoids. Thus, flavonoids with multiple phenol groups that are sufficiently long and flexible, such as biflavonoids, hold greater potential for inhibiting A β aggregation and toxicity, and thereby prevent AD (Reinke and Gestwicki 2007). However, there are substantial challenges associated with utilizing biflavonoids as aggregation inhibitors since their effects vary greatly even with minor chemical substitutions (Ono et al. 2003, 2006; Shimmyo et al. 2008a). A clear understanding of their aggregation inhibitory effects at the molecular level has yet to emerge.

A β peptides spontaneously aggregate *in vitro* at concentrations above its critical micelle concentration. A β fibril formation can be easily detected with the Thioflavin-T fluorescence assay, as the dye binds to β -sheet rich A β fibrils and upon binding, the fluorescence of the dye at 480 nm is greatly enhanced upon excitation at 445 nm. When incubated alone in solution, the formation of A β fibrils follows a sigmoidal curve, which represents a nucleation-dependent A β polymerization mechanism. A lag phase during which fibril nuclei are formed followed by a rapid fiber extension phase are typical features of this aggregation process (Naiki et al. 1998). The final plateaued level of fluorescence in this model is indicative of the complete conversion of monomeric A β into fibrils. A study that compared the effects of the monoflavonoid apigenin

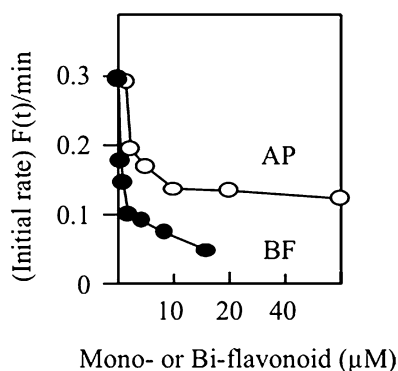


Fig. 3.2 The initial rate of A β 42 fiber extension in the presence or absence of biflavonoid taiwaniaflavone (•, BF) or monoflavonoid apigenin (○, AP) obtained by extrapolating Thioflavin-T fluorescence data (Reprinted with permission from Thapa et al. 2011)

and biflavonoid taiwaniaflavone on A β fibril formation showed that the biflavonoid reduced the final level of Thioflavin-T fluorescence to a greater extent compared to the monoflavonoid (Thapa et al. 2011). The study also showed that neither the monoflavonoids nor the biflavonoids significantly affected the nucleation step. Seeding experiments, where predominantly monomeric A β samples were seeded with preformed fibrils to eliminate the nucleation phase, were also carried out to explore the effects of the flavonoids on A β fibril extension kinetics. Initial rates of fibril extension in the presence of different concentrations of a monoflavonoid and a biflavonoid are shown in Fig. 3.2. As shown, both flavonoids decreased the rate of fibril extension with increasing flavonoid concentration. A saturation effect was apparent for both flavonoids as the decrease in rate plateaued at high flavonoid concentrations, indicating binding saturation of the flavonoids to A β species. Strikingly, the biflavonoid reduced the rate of fibril extension to a much greater extent compared to the monoflavonoid and its saturation effect is also less pronounced than that of the monoflavonoid (Fig. 3.2). Thus, the biflavonoid was found to be more effective at reducing A β fibril extension compared to the monoflavonoid (Thapa et al. 2011).

3.5.5 Biflavonoids as Inhibitors of β -Sheet Formation

The structural transition of the intrinsically unstructured A β peptides into β -sheet-enriched fibrils is a characteristic feature of the A β aggregation process. Disruption of β -sheet formation can potentially inhibit A β aggregation and fibril formation. The far UV-circular dichroism (CD) spectrum, which gives information regarding protein secondary structural content, of monomeric A β peptides in solution exhibited negative ellipticity at 190 nm, indicating a random coil structure (Fig. 3.3). Upon aggregation, the CD spectrum of A β changed and showed negative ellipticity at 217 nm, indicating the presence of β -sheet structures (Fig. 3.3) (Thapa et al. 2008). A β aggregates formed in the presence of biflavonoids contained predominantly random coil structures, whereas aggregates formed in the presence of monoflavonoids show a similar level of β -sheet structure compared to A β aggregates formed in the absence of flavonoids (Fig. 3.3). These findings suggest that while smaller flavonoid molecules such as monoflavonoids had little effect on the β -sheet formation of A β , larger flavonoid molecules such as biflavonoids can perturb β -sheet formation by A β peptides. The β -sheet structure blocking/disrupting capability of the biflavonoids also provides an explanation to its observed inhibitory effect on A β fibril extension as described in the previous section (Thapa et al. 2011).

3.5.6 Biflavonoids as Specific Inhibitors of A β -Induced Toxicity

The broad-spectrum biological activities of polyphenols and flavonoids, including anti-oxidant, anti-microbial, anti-cancer, anti-inflammatory, and metal-chelating activities, are now well known. How some of these biological activities, such as anti-oxidant, metal chelating, and anti-inflammatory effects, contribute towards the beneficial roles polyphenols and flavonoids

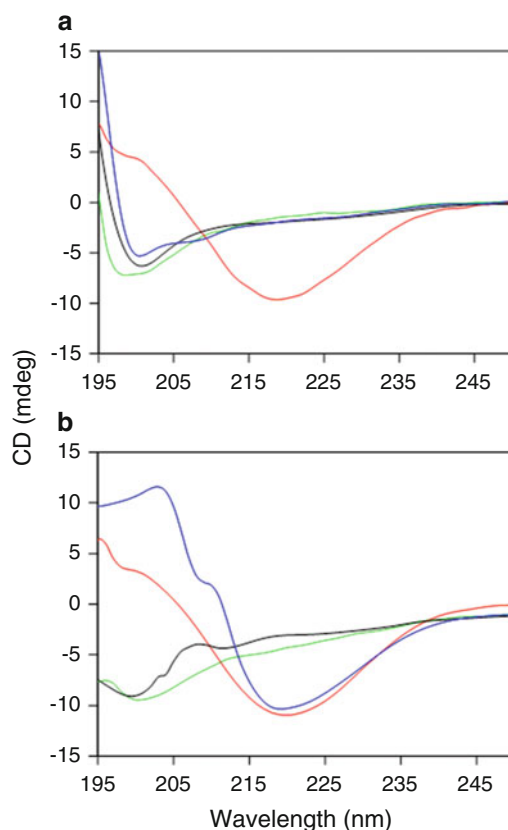


Fig. 3.3 Circular dichroism (CD) spectra of 20 μ M A β 42 incubated under different conditions. (a): A β incubated alone for 0 h (green) or 12 h (red) and A β incubated with 20 μ M biflavonoid taiwaniallavone for 0 h (black) or 12 h (blue). (b): A β incubated alone for 0 h (green) or 12 h (red) and A β incubated with 20 μ M monoflavonoid apigenin for 0 h (black) or 12 h (blue) (Reprinted with permission from Thapa et al. 2011)

play in AD and other aging related illnesses are also becoming more clear (Shin et al. 2006; Kim et al. 2008; You et al. 2013). In addition to these known effects, many reports have documented that polyphenols can directly mediate A β aggregation as well as ameliorate A β -induced toxicity. However, these effects vary and a consensus regarding their molecular mechanisms has yet to emerge.

A number of A β aggregation inhibitors that have been identified (see Sect. 3.3.8) are polyphenols and some are flavonoids. However, despite their anti-amyloidogenic activity, many of the compounds do not exhibit a neuroprotective

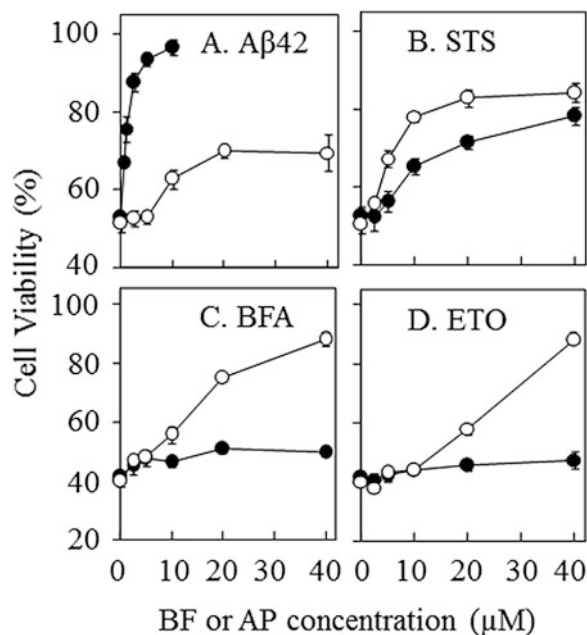


Fig. 3.4 Percentage cell viability of human neuroblastoma SH-SY5Y cells treated with 0.5 μM predominantly monomeric A β 42 (A: A β 42), staurosporine (B: STS), brefeldin A (C: BFA), and etoposide (D: ETO) in the presence or absence of varying concentrations of biflavonoid taiwaniaflavone (●, BF) or monoflavonoid api-

genin (○, AP). Cells were treated for 12 h in DMEM/F12 containing media without phenol red and fetal bovine serum. Cell viability was determined by MTT assay. Data are presented as the mean \pm standard deviation from 3 to 4 independent experimental measurements (Reprinted with permission from Thapa et al. 2011)

effect (Thapa et al. 2011; Ramassamy 2006; Ono et al. 2003, 2006). The flavonoid quercetin inhibits A β fibril formation to a similar extent compared to myricetin, but it is less effective in ameliorating A β toxicity compared to myricetin, which possesses an additional -OH group (Ono et al. 2003). Some polyphenols, including flavonoids, have been reported to inhibit A β aggregation but not A β -induced toxicity, while others have been found to be neuroprotective without significantly reducing A β aggregation (Ono et al. 2003; Wang et al. 2001; Feng et al. 2009; Sharoar et al. 2012). While it is difficult to distinguish and establish the specific roles these multifunctional polyphenols, flavonoids, and biflavonoid molecules play in a complex disease such as AD, some studies have shown that biflavonoids are more effective at inhibiting A β aggregation and reducing A β -induced toxicity and may be a better class of naturally-occurring compounds to target AD. We review below

recent publications on the effects and proposed neuroprotective mechanisms of biflavonoids on A β -induced toxicity.

A study comparing the effects of monoflavonoids and biflavonoids on A β -induced toxicity in cultured neuronal cells has revealed some insights into the neuroprotective properties of these two types of flavonoids (Thapa et al. 2011). In agreement with other published reports, flavonoids were shown to rescue cells exposed to monomeric A β 42 peptides (Thapa et al. 2011). However, the effects of the biflavonoids and monoflavonoids are distinctly different. The study showed that while the presence of 10 μM monoflavonoid apigenin increased the viability of cultured SH-SY5Y neuroblastoma cell by 50 % when exposed to 0.5 μM of monomeric A β 42 for 12 h, only approximately 1 μM of the biflavonoid was required to achieve the same level of neuroprotection (Fig. 3.4) (Thapa et al. 2011).

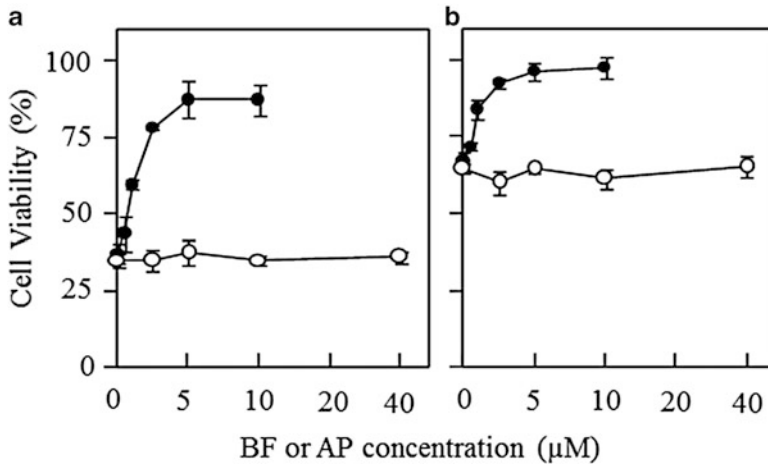


Fig. 3.5 Percentage cell viability of human neuroblastoma SH-SY5Y cells treated with 0.5 μM preformed A β 42 oligomers (a) or fibrils (b) in the presence or absence of biflavonoid taiwaniflavone (•, BF) or monoflavonoid apigenin (○, AP). Cells were treated for

12 h in DMEM/F12 containing media without phenol red and fetal bovine serum. Cell viability was determined by MTT assay. Data are presented as the mean \pm standard deviation from 3 to 4 independent experimental measurements (Reprinted with permission from Thapa et al. 2011)

To test the specificity of the rescuing effects of the flavonoids towards A β -induced toxicity, the effects of the two types of flavonoids on the viability of cultured SH-SY5Y cells exposed to a number of known toxic compounds including staurosporine (an inhibitor of protein kinase), brefeldin A (a protein transport inhibitor to Golgi complex), and etoposide (an inhibitor of DNA topoisomerase) were also studied (Karki et al. 2007; Shin et al. 2006). The monoflavonoid apigenin was found to be moderately effective in increasing the viability of cells exposed to A β and was found to be equally effective at rescuing cells exposed to the other toxic compounds. Previously, it was shown that apigenin abrogates the activation of the executioner molecule caspase-3 in the apoptotic cell death cascade (Wang et al. 2001). Such a cytoprotective mechanism explains the moderate and non-specific rescuing effect that has been observed for the monoflavonoid. In contrast to the nonspecific rescuing effect of monoflavonoids, biflavonoids rescued cells to a greater extent towards toxicity induced by monomeric A β 42 compared to toxicity induced by the other compounds (Fig. 3.4). These data suggest that biflavonoids more specifically and more effectively reduce toxicity induced

by A β peptide compared to monoflavonoids (Fig. 3.4). The enhanced rescuing effects can arise from biflavonoid's ability to either more potently inhibit A β peptide aggregation during the incubation period with the cells (Figs. 3.2 and 3.3) or that biflavonoids are more effective at directly blocking monomeric A β 42 toxicity.

In a separate study, the neuroprotective effects of mono- and bi-flavonoids against oligomeric and fibrillar A β 42 were examined. In contrast to monomeric A β 42, the soluble A β 42 oligomers are known to be significantly more toxic and are believed to be the primary neurotoxic species in the A β peptide aggregation cascade (Deshpande et al. 2006; Walsh and Selkoe 2007). As shown in Fig. 3.5, preformed 0.5 μM A β 42 oligomers induced nearly 16% more cell death compared to monomeric A β 42. The biflavonoid taiwaniflavone significantly increased the viability of cells exposed to preformed A β oligomers and fibrils whereas the monoflavonoid apigenin did not offer neuroprotection against A β oligomers and fibrils (Fig. 3.5). These findings suggest that the biflavonoids are uniquely effective at inhibiting the toxicity of soluble A β oligomers (Fig. 3.5) and are therefore promising small molecule therapeutics for the prevention of A β toxicity and neurodegeneration in AD.

3.5.7 Biflavonoids Accumulate Nontoxic A β Oligomers

Since soluble A β oligomers are believed to be the primary toxic species of A β causing neurodegeneration, small molecules, including flavonoids, that selectively bind to A β oligomers over monomeric or fibrillar A β are considered as potential therapeutics for targeting and neutralizing A β toxicity in AD (Walsh and Selkoe 2007; Klein 2002; Yang et al. 2005). Several other small molecules that inhibit A β aggregation have also been shown as inhibitors of A β oligomerization and/or A β fibril formation (Necula et al. 2007). A study on biflavonoids has revealed that these compounds efficiently capture small A β oligomers and reduce subsequent A β aggregation (Thapa et al. 2011). Immunoblotting analysis of glutaraldehyde cross-linked A β 42 aggregates formed by incubating A β alone showed the presence of high molecular weight A β species along with the disappearance of monomeric A β . Analysis of soluble A β aggregates formed in the presence of the biflavonoid taiwaniaflavone showed that significantly more low-molecular-weight (LMW) A β 42 aggregates (including monomeric, dimeric, and trimeric A β 42) were present, along with small amounts of high-molecular-weight (HMW) soluble oligomers spanning a broad size range. Moreover, the levels of LMW A β aggregates and HMW A β oligomers formed in the presence of the biflavonoid increased in a dose-dependent manner (Fig. 3.6). In contrast, A β aggregates formed in the presence of the monoflavonoid apigenin did not contain LMW A β aggregates; the oligomers that were present were larger than 30 kDa (Fig. 3.6). Similar results were also obtained from studies that used other biflavonoids, including 2',8''-biapigenin, amentoflavone, sumafavone (Thapa et al. 2011). These results suggest that biflavonoids uniquely and selectively bind to soluble LMW A β oligomers. As a result, these A β oligomers are thermodynamically favored, leading to their increased levels. Critically, these small and soluble A β oligomers bound to biflavonoids have been shown to be nontoxic and unstructured. The A β aggregates formed in the

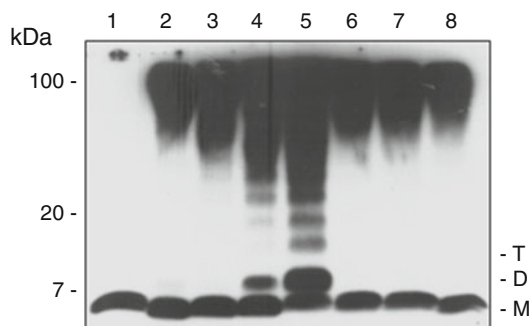


Fig. 3.6 Western blot analysis of 0.5 μ M A β incubated alone or in the presence of biflavonoid taiwaniaflavone or monoflavonoid apigenin. Lane-1: A β incubated alone; lane-2: A β incubated for 3 h; lanes 3–5: A β incubated in the presence of taiwaniaflavone (2.5, 5 and 10 μ M) for 3 h; lanes 6–8: A β incubated in the presence of apigenin (2.5, 5 and 10 μ M) for 3 h. A β incubated in PBS at 37 $^{\circ}$ C was cross-linked with 0.01 % (v/v) glutaraldehyde before separating on 16 % acrylamide gel. The samples were then transferred onto PVDF membrane and probed with A β specific 6E10 monoclonal antibody at 1:10,000 dilution. Monomer, dimer, and trimer bands are indicated by M, D and T, respectively (Reprinted with permission from Thapa et al. 2011)

presence of biflavonoids were thus structurally and functionally different than on-pathway toxic A β aggregates (Thapa et al. 2011).

3.5.8 Mechanisms of Biflavonoid-Mediated Neuroprotection Against A β -Induced Toxicity

Elucidating the molecular mechanisms by which polyphenols affect A β aggregation and toxicity remains an active research area. Several recently published reports gave useful insights to the mechanisms by which polyphenols and flavonoids antagonize A β -induced toxicity. In one study, Necula and co-workers investigated the effects of a large number of small molecules on the oligomerization and fibrillization of A β 42. Results from the study showed that the small molecules could be categorized into three groups based on their inhibitory effects on A β 42 aggregation. One group was found to be effective at inhibiting the oligomerization

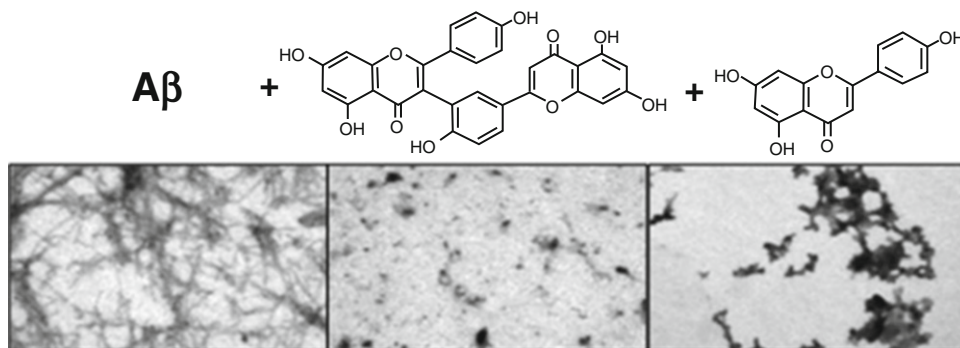


Fig. 3.7 Schematics of mono- and bi-flavonoids (top) and TEM images (bottom) of A β incubated alone (left image) or in the presence of biflavonoid taiwaniaflavone (middle image) or monoflavonoid apigenin (right image) (Reprinted with permission from Thapa et al. 2011)

of A β while another group was found to be effective at inhibiting the fibrillization of A β . The third group of small molecules were effective at inhibiting both fibrillization and oligomerization of A β (Necula et al. 2007). In a separate study, Ladiwala and co-workers investigated the effects of polyphenolic compounds on A β 42 aggregation and classified the aromatic molecules according to their abilities to (i) remodel soluble oligomers into non-toxic off-pathway aggregates, (ii) convert soluble A β oligomers into fibrils, and (iii) disaggregate soluble oligomers or fibrils into non-toxic, LMW species (Ladiwala et al. 2011).

In a study that specifically examined the effects of biflavonoids on A β 42 aggregation kinetics, Thapa and co-workers showed that the compounds primarily affected the fibril extension phase of A β aggregation. Biochemical analysis further revealed that biflavonoids captured soluble A β aggregates more efficiently than monoflavonoids (Fig. 3.6) (Thapa et al. 2011). Moreover, transmission electron microscopy (TEM) imaging of A β aggregates showed that A β incubated alone contained predominantly fibrillar aggregates whereas the A β sample incubated with biflavonoids contained small globular oligomeric aggregates (Fig. 3.7). In contrast, beaded protofibril-like A β aggregates were found in the A β sample incubated with monoflavonoids. Because the biflavonoids, but not the monoflavonoids, were found to be neuroprotective against preformed A β oligomers and fibrils in this study, it was concluded that

the oligomers accumulated by the biflavonoids are non-toxic (Fig. 3.6). This study shows that while the biflavonoids are effective at reducing A β fibril extension, they are also able to direct the A β aggregation pathway towards a non-toxic pathway, as has been proposed for the neuroprotective mechanism of tea polyphenols (Ehrnhoefer et al. 2008).

3.6 Conclusion

Sporadic AD represents over 95 % of all AD cases (Hamley 2012; Thies and Bleiler 2013). The dominant prevalence of sporadic over familial AD indicates that reduction of A β production by manipulating secretases that cleave APP may only have a limited effect in preventing and treating AD (Nie et al. 2011). It also highlights the fact that A β aggregation is critical in AD development; hence, inhibiting or modifying A β aggregation is a highly promising AD therapeutic strategy. Because the soluble oligomers and protofibrils formed during A β fibril formation are believed to be the primary toxic species and their toxicity correlates well with the progression of AD, the inhibition of A β aggregation, especially at an early stage that inhibits the formation of toxic oligomers, appears to be a promising preventive strategy for AD (Ehrnhoefer et al. 2008; Ladiwala et al. 2010; Tomic et al. 2009).

Unfortunately, a number of clinical trials that have been conducted in the past few years, including heavy metal chelators, oxidative

stress reducers, and anti-inflammatory inhibitors, have yielded largely unsatisfactory outcomes (Blennow et al. 2006; Nie et al. 2011). Perhaps an important lesson that emerges from these trials is that compounds that have multiple modes of action against AD development, especially those that inhibit A β aggregation, may be promising drug candidates for the prevention and treatment of AD.

Polyphenols are a class of naturally occurring compounds with well-documented anti-oxidant and anti-inflammatory properties. There is increasing interest in utilizing these compounds as AD therapeutics as they have also been shown to be capable of disrupting hydrophobic and π - π interactions of A β peptides and prevent A β aggregation and toxicity (Armstrong et al. 2011; Gazit 2002). Studies on the structure-function relationship of polyphenols revealed that a minimum of two phenyl rings connected by a suitably long and flexible linker are essential features that contribute to aggregation inhibition (Reinke and Gestwicki 2007). In this regard, the flat and planar biflavonoid compounds that are endowed with multiple phenyl ring moieties connected by appropriately sized and flexible linkers may target multiple and specific regions on the A β peptide in a concerted manner. Biflavonoids have indeed been shown to exert a stronger aggregation inhibition effect compared to the smaller-sized monoflavonoids, possibly due to their higher affinity to A β and the greater steric hindrance for fibril growth when bound to A β peptides. These properties are believed to contribute to the enhanced anti-amyloidogenic activity of biflavonoids, resulting in reduced A β toxicity. More importantly, biflavonoids have been shown to reduce A β oligomers toxicity more specifically while capturing soluble A β oligomers and reducing A β aggregation.

These unique and potent biochemical and biological activities of biflavonoids suggest that they may be better candidate small molecules in preventing AD as compared to polyphenols, including monomeric flavonoid, in AD interventions. Further studies in an *in vivo* setting using transgenic mouse models of AD are therefore urgently warranted.

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Natural Phenolic Compounds as Therapeutic and Preventive Agents for Cerebral Amyloidosis

4

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Abstract

Epidemiological studies have suggested that diets rich in phenolic compounds may have preventive effects on the development of dementia or Alzheimer's disease (AD). We investigated the effects of natural phenolic compounds, such as myricetin (Myr), rosmarinic acid (RA), ferulic acid (FA), curcumin (Cur) and nordihydroguaiaretic acid (NDGA) on the aggregation of amyloid β -protein ($A\beta$), using *in vitro* and *in vivo* models of cerebral $A\beta$ amyloidosis. The *in vitro* studies revealed that these phenolic compounds efficiently inhibit oligomerization as well as fibril formation of $A\beta$ through differential binding, whilst reducing $A\beta$ oligomer-induced synaptic and neuronal toxicity. Furthermore, a transgenic mouse model fed orally with such phenolic compounds showed significant reduction of soluble $A\beta$ oligomers as well as of insoluble $A\beta$ deposition in the brain. These data, together with an updated review of the literature, indicate that natural phenolic compounds have anti-amyloidogenic effects on $A\beta$ in addition to well-known anti-oxidative and anti-inflammatory effects, hence suggesting their potential as therapeutic and/or preventive agents for cerebral $A\beta$ amyloidosis, including AD and cerebral amyloid angiopathy (CAA). Well-designed clinical trials or preventive interventions with natural phenolic compounds are necessary to establish their efficacy as disease-modifying agents.

Keywords

Alzheimer's disease • Amyloid β -protein • Amyloidosis • Polyphenols • Therapy

Abbreviations

$A\beta$	amyloid β -protein
AD	Alzheimer's disease
ADAS-Cog	Alzheimer's Disease Assessment Scale – Cognitive Subscale

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α S	α -synuclein
ApoE	apolipoprotein E
APP	amyloid- β precursor protein
CAA	cerebral amyloid angiopathy
CD	circular dichroism
CSF	cerebrospinal fluid
Cur	curcumin
DLB	dementia with Lewy bodies
EGCG	(-)-epigallocatechin-3-galate
FA	ferulic acid
fA β	A β fibrils
GSPE	grape seed polyphenolic extract
LTD	long-term depression
LTP	long-term potentiation
MCI	mild cognitive impairment
MMSE	Mini-Mental State Examination
MPTP	1-methyl-4-phenyl-1,2,3,6-tetrahydro-pyridine
MTT	3-[4,5-dimethylthiazol-2-yl]-2,5-diphenyltetrazolium bromide
Myr	myricetin
NDGA	nordihydroguaiaretic acid
NMR	nuclear magnetic resonance
PD	Parkinson's disease
PHF	paired helical filament
PICUP	photo-induced cross-linking of unmodified proteins
RA	rosmarinic acid

4.1 Epidemiological Studies Suggesting Preventive Effects of Phenol Compound-Rich Diets on Dementia or Alzheimer's Disease

Epidemiological studies have reported that diets rich in phenolic compounds or polyphenols may be associated with a reduced risk of dementia or Alzheimer's disease (AD). These include vegetables, fruits, spice, and derived products such as wine and non-alcoholic beverages.

The Mediterranean diet, characterized by a high intake of vegetables, fruits, cereals, olive oil, fish, in combination with a low intake of meat and poultry, was reported to be associated with

a reduction in risk of dementia, mild cognitive impairment (MCI), and AD in prospective longitudinal studies (Scarmeas et al. 2006, 2009; Féart et al. 2009). In a randomized trial with nutritional intervention comparing two Mediterranean diets supplemented with either extra-virgin olive oil or nuts versus a low-fat control diet for 6.5 years, cognitive performance examined by Mini-Mental State Examination (MMSE) and Clock Drawing Test was significantly better in the group of Mediterranean diets than in a low-fat control group, after adjusting multiple confounding factors (Martínez-Lapiscina et al. 2013). Recent systematic reviews with meta-analysis indicate that a higher adherence to Mediterranean diet is associated with a reduced risk of MCI, dementia, and AD; nevertheless, further prospective cohort studies with longer follow-up and randomized controlled trials are necessary to unequivocally establish the effects of this type of diet on cognitive decline and AD (Psaltopoulou et al. 2013; Singh et al. 2014).

Traditional Indian diets and medicines contain spices such as yellow curry spice turmeric, curcumin. Frequency of AD in India is about one-quarter of that in the US (aged 70–79 years, 0.7 % vs 3.1 %; aged 80 years or older, 4.0 % vs 15.7 %), suggesting influence of ethnic differences in environmental, including dietary, apart from genetic factors (Ganguli et al. 2000). In a population-based cohort of non-demented elderly Asian subjects, more curry consumption was associated with better cognitive performance suggesting possible preventive effects of curry spice curcumin on cognitive decline, although further prospective cohort studies with long follow-up are required (Ng et al. 2006).

Several prospective cohort studies also reported that moderate intake of wine was associated with a reduced risk of dementia, AD, or cognitive decline (Orgogozo et al. 1997; Truelsen et al. 2002; Luchsinger et al. 2004; Solfrizzi et al. 2007; Mehlig et al. 2008; Arntzen et al. 2010). As this protective effect was not seen for alcoholic beverages other than wine (Truelsen et al. 2002; Luchsinger et al. 2004; Mehlig et al. 2008; Arntzen et al. 2010), it is suggested that

the association for wine may be attributable to components of wine other than ethanol itself. In a population-based prospective study, consumption of fruit and vegetable juices, containing a high concentration of polyphenols, decreased a risk of AD (Dai et al. 2006).

Coffee, black tea, and green tea are enriched in polyphenols, and may be protective against onset of dementia including AD. Several longitudinal studies (Lindsay et al. 2002; van Gelder et al. 2007; Ritchie et al. 2007; Eskelinen et al. 2009) have investigated the relationship between coffee consumption and dementia, AD, or cognitive decline, but findings from these studies are inconsistent. Longitudinal studies of black tea consumption have not found any association with reduced risks for dementia, AD, or cognitive decline (Laurin et al. 2004; Dai et al. 2006). One cross-sectional study has shown that higher green tea consumption is associated with lower prevalence of cognitive impairment (Kuriyama et al. 2006). To determine whether the consumption of green tea, coffee, or black tea influences the incidence of dementia and MCI in older people, we recently conducted a population-based prospective study with Japanese residents aged >60 years from Nakajima, Japan (the Nakajima Project) (Noguchi-Shinohara et al. 2014). Participants received an evaluation of cognitive function and blood tests. The consumption of green tea, coffee, and black tea was also evaluated at baseline. Of 723 participants with normal cognitive function at a baseline survey (2007–2008), 490 completed the follow up survey in 2011–2013. The incidence of dementia during the follow-up period (mean \pm SD: 4.9 \pm 0.9 years) was 5.3 %, and that of MCI was 13.1 %. To analyze the independent effects of green tea, coffee, and black tea consumption on the risk of developing dementia or MCI, multivariate logistic regression analysis was performed with adjustment for sex, age, history of hypertension, diabetes mellitus, and hyperlipidemia, formal education, apolipoprotein E (*ApoE*) phenotype status (*ApoE* E4+ or E4-), smoking status, alcohol consumption, green tea, coffee, and/or black tea consumption, physical activities and/or hobbies. The multiple-adjusted odds ratio for the incidence of overall cogni-

tive decline (dementia or MCI) was 0.32 (95 % CI: 0.16–0.64) among individuals who consumed green tea every day and 0.47 (95 % CI: 0.25–0.86) among those who consumed green tea 1–6 days per week compared with individuals who did not consume green tea at all (Fig. 4.1). No association was found between coffee or black tea consumption and the incidence of dementia or MCI. Our results indicate that green tea consumption is significantly associated with reduced risk of cognitive decline, even after adjustment for possible confounding factors. This was the first prospective longitudinal study that examined the association between green tea consumption and incidence of dementia or cognitive decline.

Figure 4.2 shows components of green tea, black tea, and coffee with possible effects on cognitive decline. The major tea-related polyphenols present in green tea are catechins, especially (–)-epigallocatechin-3-gallate (EGCG), whereas black tea mainly contains theaflavins (Peterson et al. 2005). In addition, green tea contains greater amounts of myricetin (Myr) compared with black tea (Peterson et al. 2005). Other tea-related polyphenols such as quercetin, kaempferol, apigenin, and luteolin, are also present in both green and black tea, but the amounts of these polyphenols are not significantly different between tea types (Peterson et al. 2005). The caffeine content is 40–57 mg/100 mL in coffee (Barone and Roberts 1996), 25.5 mg/100 mL in black tea, and only 15.3 mg/100 mL in green tea (Khokhar and Magnusdottir 2002). High intake of ascorbic acid was reported to be associated with lower risk of AD (Engelhart et al. 2002). The content of ascorbic acid is 6 mg/100 mL in green tea, which is the most common source of ascorbic acid in Japan (Ogawa et al. 2002); on the other hand, coffee and black tea do not contain ascorbic acid. As the serum levels of ascorbic acid were associated with the frequency of coffee consumption, but not green tea consumption in our study (Noguchi-Shinohara et al. 2014), it is unlikely that the effects of green tea on cognitive function could be explained as those of ascorbic acid. Taken together, phenolic compounds enriched in green

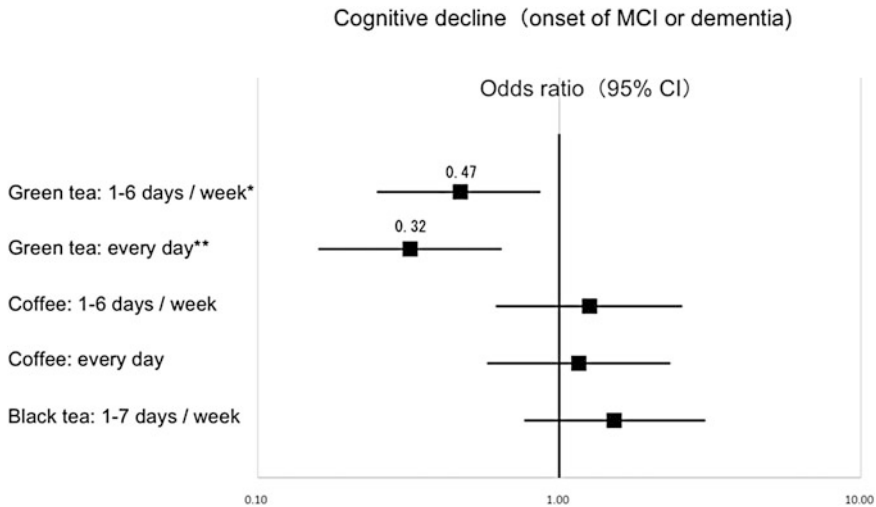


Fig. 4.1 Association between green tea, coffee, or black tea consumption and the incidence of cognitive decline [mild cognitive impairment (MCI) or dementia] in cognitively normal subjects (age > 60 years) ($n = 490$) during the follow-up period (mean \pm SD: 4.9 ± 0.9 years). The multiple-adjusted odds ratio[#] for the incidence of cognitive decline (MCI or dementia) is shown compared with individuals who did not consume green tea, coffee, or black tea at all. Details were reported in the reference (Noguchi-Shinohara et al. 2014)
[#]Multivariate logistic regression models were used to

analyze the independent effects of green tea, coffee, and black tea consumption on the risk of developing dementia or MCI so that the lowest category (none) served as the reference group. Model was adjusted for sex, age, history of hypertension, diabetes mellitus, and hyperlipidemia, formal education, apolipoprotein E (*ApoE*) phenotype status (*ApoE* E4+ or E4-), smoking status, alcohol consumption, green tea, coffee and/or black tea consumption, physical activities and/or hobbies* P -value < 0.05, ** P -value < 0.01

tea, such as EGCG and myricetin, would be candidates to exert preventive effects on cognitive decline (Fig. 4.2).

4.2 Effects of Natural Phenolic Compounds in Cerebral Amyloidosis Models

4.2.1 Studies Using *In Vitro* Models of Cerebral Amyloidosis

Natural phenolic compounds have been commonly reported as having anti-oxidant, anti-inflammatory, and other activities that may exert neuroprotective effects on AD and other dementias. However, the remarkable effects of such compounds on cognitive decline observed in epidemiological studies with older people suggest that they may have more specific effects on pathways involved in the pathophysiology of cerebral amyloidosis and

other neurodegenerative disorders such as dementia with Lewy bodies (DLB).

Cerebral parenchymal deposition of the amyloid β -peptide ($A\beta$) is a central feature of AD. In addition, $A\beta$ deposits in the cerebral vasculature of older subjects and AD, called cerebral amyloid angiopathy (CAA), cause cerebral hemorrhages and other cerebrovascular disorders. As amyloid deposition is considered to be the most upstream event in AD pathogenesis (amyloid cascade hypothesis), the process of $A\beta$ deposition is the main target of drug development in AD. Although α -cleavage of amyloid- β precursor protein (APP) by α -secretase prevents production of $A\beta$, β - and γ -cleavages of APP by β - and γ -secretases produce $A\beta$; $A\beta$ peptides subsequently aggregate from monomers to oligomers, protofibrils, and fibrils (Fig. 4.3). Moreover, tau protein is phosphorylated and aggregates forming intracellular neurofibrillary tangles composed of paired helical filaments. Finally, synaptic dysfunction and neuronal death occur. In DLB as well as Parkinson's

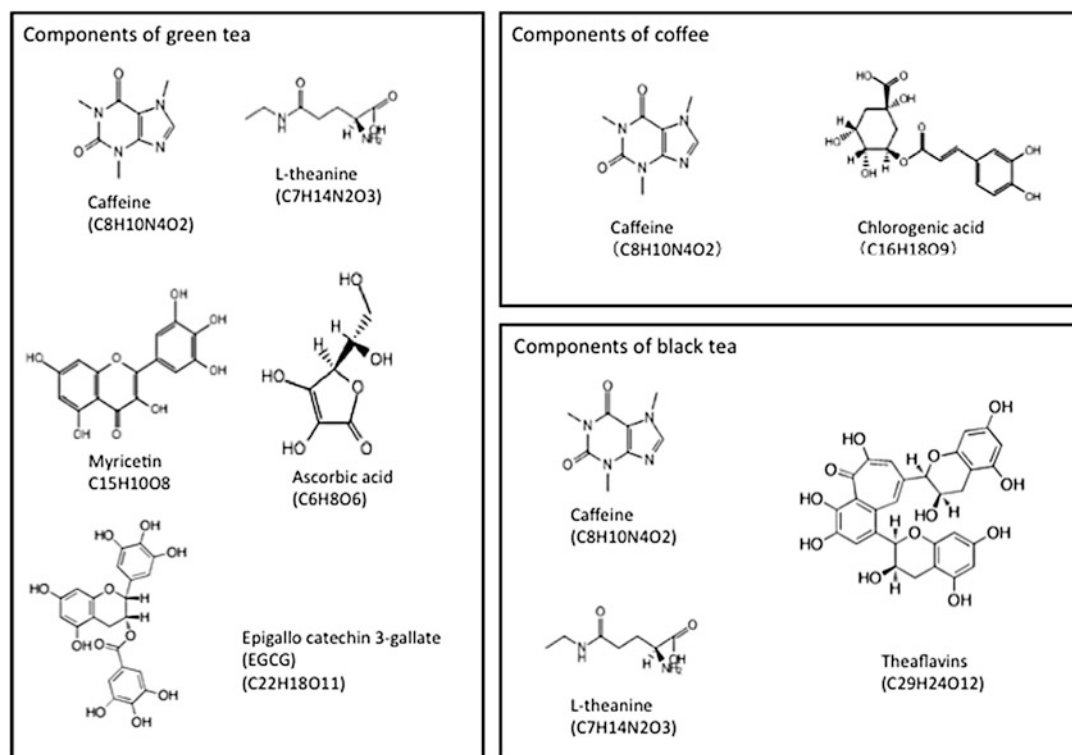


Fig. 4.2 Components of green tea, coffee, and black tea that may be implicated in preventive effects on cognitive decline

disease (PD), α -synuclein (α S) is aggregated in neuronal cell bodies and neurites (Lewy bodies and neurites) in the brain (α -synucleinopathies). Widespread tau aggregation is found in other neurodegenerative dementias than AD (non-AD tauopathies), such as Pick's disease, argyrophilic grain disease, and senile dementia of the neurofibrillary tangle type.

Recent studies have reported that natural phenolic compounds have the following specific actions: modulation of the processing of APP (Levites et al. 2003; Rezai-Zadeh et al. 2005; Obregon et al. 2006; Chakraborty et al. 2011; Kostomoiri et al. 2013; Yoshida et al. 2014; Zhang et al. 2013), inhibition of A β aggregation and remodeling and destabilization of aggregates (Ono et al. 2002, 2003, 2004a, b, 2005, 2008, 2012; Yang et al. 2005; Bastianetto et al. 2006; Rivière et al. 2007, 2009; Ehrnhoefer et al. 2008; Shoval et al. 2008; Wang et al. 2008, 2014; Bieschke et al. 2010; Grelle et al.

2011; Thapa et al. 2011; Rigacci et al. 2011; Hirohata et al. 2012; Ge et al. 2012; Cheng et al. 2013; Sinha et al. 2012; Rushworth et al. 2013; Palhano et al. 2013; Zhang et al. 2013; Ho et al. 2013; Cui et al. 2013; Richard et al. 2013; da Silva Bittencourt et al. 2014), promotion of A β degradation/clearance (Marambaud et al. 2005; Vingtdoux et al. 2010), alleviation of A β -induced oxidative stress/toxicity/synaptic dysfunction (Ono et al. 2003; Savaskan et al. 2003; Sultana et al. 2005; Bastianetto et al. 2006; Joshi et al. 2006; Feng et al. 2009, 2013; Bieschke et al. 2010; Choi et al. 2010; He et al. 2011; Fuentealba et al. 2011, 2012; Grelle et al. 2011; Rushworth et al. 2013; Ho et al. 2013; Wong et al. 2013; Cimini et al. 2013; Camilleri et al. 2013; da Silva Bittencourt et al. 2014), inhibition of α S aggregation (Ono and Yamada 2006; Masuda et al. 2006, 2009; Ehrnhoefer et al. 2008; Bieschke et al. 2010; Grelle et al. 2011; Marchiani et al. 2013), detoxification

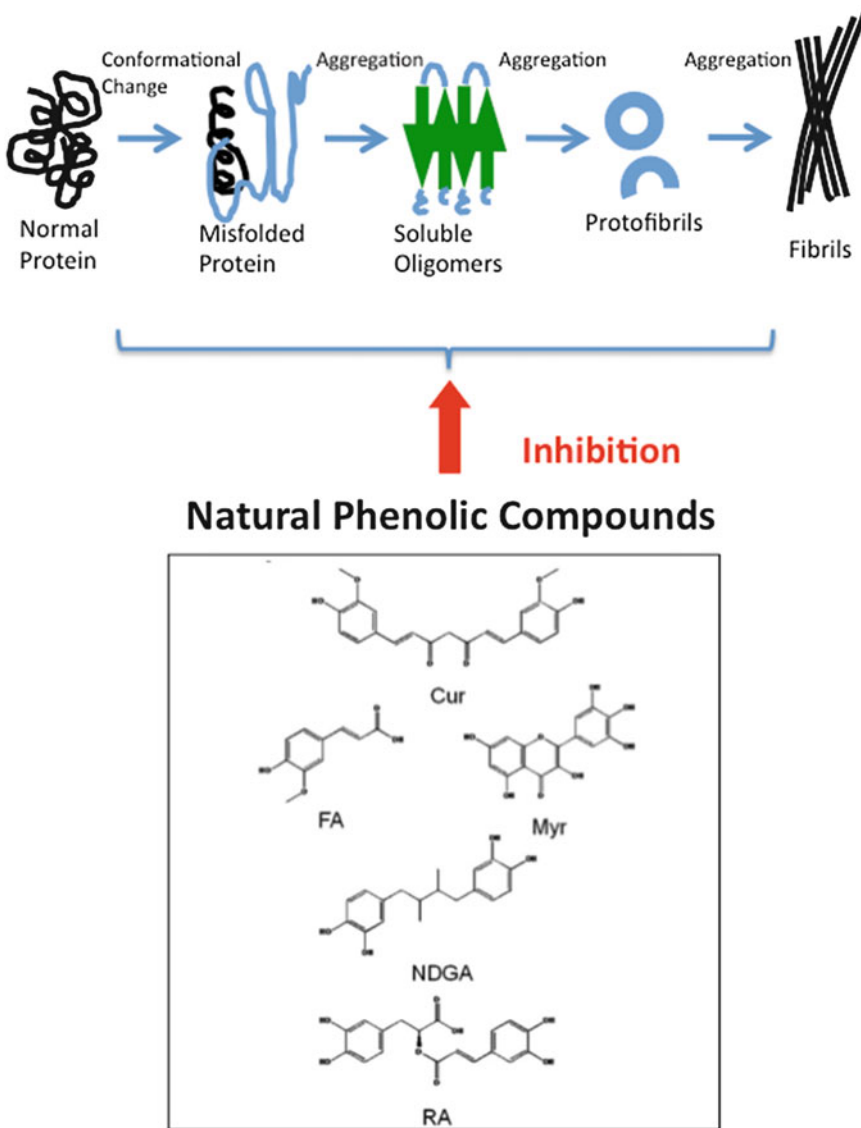


Fig. 4.3 A pathway of protein aggregation for the amyloid- β peptide ($A\beta$) and α -synuclein protein. The same five phenolic compounds with inhibitory effects on $A\beta$ aggregation in our *in vitro* studies were used for our

in vivo studies with an animal model. *Cur* curcumin, *FA* ferulic acid, *Myr* myricetin, *NDGA* nordihydroguaiaretic acid, *RA* rosmarinic acid

of α S aggregates (Bieschke et al. 2010; Grelle et al. 2011; Caruana et al. 2012; Marchiani et al. 2013; Lorenzen et al. 2014), and inhibition of tau phosphorylation and aggregation (Taniguchi et al. 2005; Ho et al. 2009b; Ksiezak-Reding et al. 2012; Patil et al. 2013; Yao et al. 2013). Such effects have been reported in various phenolic compounds: flavones such as baicalein (Caruana

et al. 2012), flavonols such as Myr, quercetin, and morin (Ono et al. 2003, 2006a, 2012; Masuda et al. 2006; Chakraborty et al. 2011; Caruana et al. 2012; Ho et al. 2013), isoflavones such as glycitein and genistein (Hirohata et al. 2012), flavanols such as EGCG and theaflavins (Ono et al. 2003, 2006a; Levites et al. 2003; Rezai-Zadeh et al. 2005; Obregon et al. 2006;

Bastianetto et al. 2006; Ehrnhoefer et al. 2008; Bieschke et al. 2010; He et al. 2011; Grelle et al. 2011; Cheng et al. 2013; Sinha et al. 2012; Rushworth et al. 2013; Palhano et al. 2013; Zhang et al. 2013; Lorenzen et al. 2014), stilbenes such as resveratrol, nordihydroguaiaretic acid (NDGA), piceid, and viniferin (Ono et al. 2002, 2003, 2006a, 2012; Savaskan et al. 2003; Marambaud et al. 2005; Rivière et al. 2007, 2009; Gauci et al. 2011; Capiralla et al. 2012; Ge et al. 2012; Feng et al. 2013; Vingtdeux et al. 2010; Caruana et al. 2012; Rushworth et al. 2013; Richard et al. 2013), phenolic acids such as rosmarinic acid (RA), tannic acids (TA), ferulic acid (FA), ellagic acid, and gallic acid (Ono et al. 2003, 2004b, 2005, 2006a, 2012; Sultana et al. 2005; Joshi et al. 2006; Feng et al. 2009; Cui et al. 2013; Zhang et al. 2013; Yao et al. 2013; Yoshida et al. 2014), curcuminoids such as curcumin (Cur) (Ono et al. 2004b, 2006a, 2012; Yang et al. 2005; Shoal et al. 2008; Marchiani et al. 2013; Patil et al. 2013), secoiridoids such as oleuropein (Rigacci et al. 2011; Kostomoiri et al. 2013), and others (Thapa et al. 2011). In addition, extracts of phenolic compounds of natural products have been used for studies, including extracts of grape seeds, wine, berries, tea, cocoa, guarana, and *Pueraria lobata* (Ono et al. 2008; Wang et al. 2008, 2014; Ho et al. 2009a; Choi et al. 2010; Fuentealba et al. 2011, 2012; Gauci et al. 2011; Caruana et al. 2012; Ksiezak-Reding et al. 2012; Wong et al. 2013; Cimini et al. 2013; da Silva Bittencourt et al. 2014).

To develop therapeutics and preventives for cerebral A β amyloidosis (AD and CAA), we investigated whether such natural phenolic compounds with possible anti-dementia/AD effects suggested in the epidemiological studies have anti-aggregation effects on A β . We first examined the effects of Myr, morin, quercetin, kaempferol (+)-catechin, (-)-epicatechin, NDGA, Cur, RA, and FA on the formation, extension, and destabilization of A β fibrils (fA β) *in vitro*, using fluorescence spectroscopy with thioflavin T and electron microscopy (Ono et al. 2003, 2004b, 2006b). All examined phenolic compounds dose-dependently inhibited formation of fA β from fresh A β (1–40) (A β 40)

and A β (1–42) (A β 42), as well as their extension (Fig. 4.4a–c). Moreover, these polyphenols dose-dependently destabilized preformed fA β 40 and fA β 42. The effective concentrations (EC50) of Myr, morin, quercetin, NDGA, Cur, and RA for the formation, extension and destabilization of fA β 40 and fA β 42 were in the order of 0.1–1 μ M. In cell culture experiments, Myr-treated fA β were less toxic than intact fA β , as demonstrated by 3-[4,5-dimethylthiazol-2-yl]-2,5-diphenyltetrazolium bromide (MTT) assays.

We further investigated the effects of Myr, NDGA, FA, Cur, and RA on A β oligomerization and a mechanistic basis of the anti-aggregation effects of these compounds (Ono et al. 2012). We revealed that, using the method of photo-induced cross-linking of unmodified proteins (PICUP), these five phenolic compounds dose-dependently inhibited oligomerization of A β 40 and A β 42 (Fig. 4.4d). The circular dichroism (CD) spectroscopy studies showed that both Myr and RA stabilized A β populations comprising mostly random coil and inhibited statistical coils to β -sheet conversion. However, at the atomic level, a study with nuclear magnetic resonance (NMR) spectroscopy showed that Myr and RA behave differently, in that Myr shows significant binding to monomeric A β 42 (Fig. 4.5), whereas RA does not bind to the monomer. It is possible that RA could prevent aggregation by binding to non-NMR detectable early-formed oligomers or distinct monomer conformers/structures causing the inhibition of oligomerization (Fig. 4.5).

There has been mounting evidence that A β oligomers rather than mature fibrils are toxic and considered to induce the deleterious cascade(s) involved in the pathophysiology of AD [see review (Larson and Lesne 2012)]. We investigated whether these phenolic compounds with anti-oligomerization effects could attenuate toxicity (Ono et al. 2012). Long-term potentiation (LTP) and depression (LTD) are neurophysiological models of neuronal plasticity for memory and learning; using electrophysiological assays for LTP and LTD in hippocampal slices, we found that Myr and RA decreased A β oligomer-induced synaptic toxicities. We evaluated the effects of these phenolic compounds on A β -oligomer in-

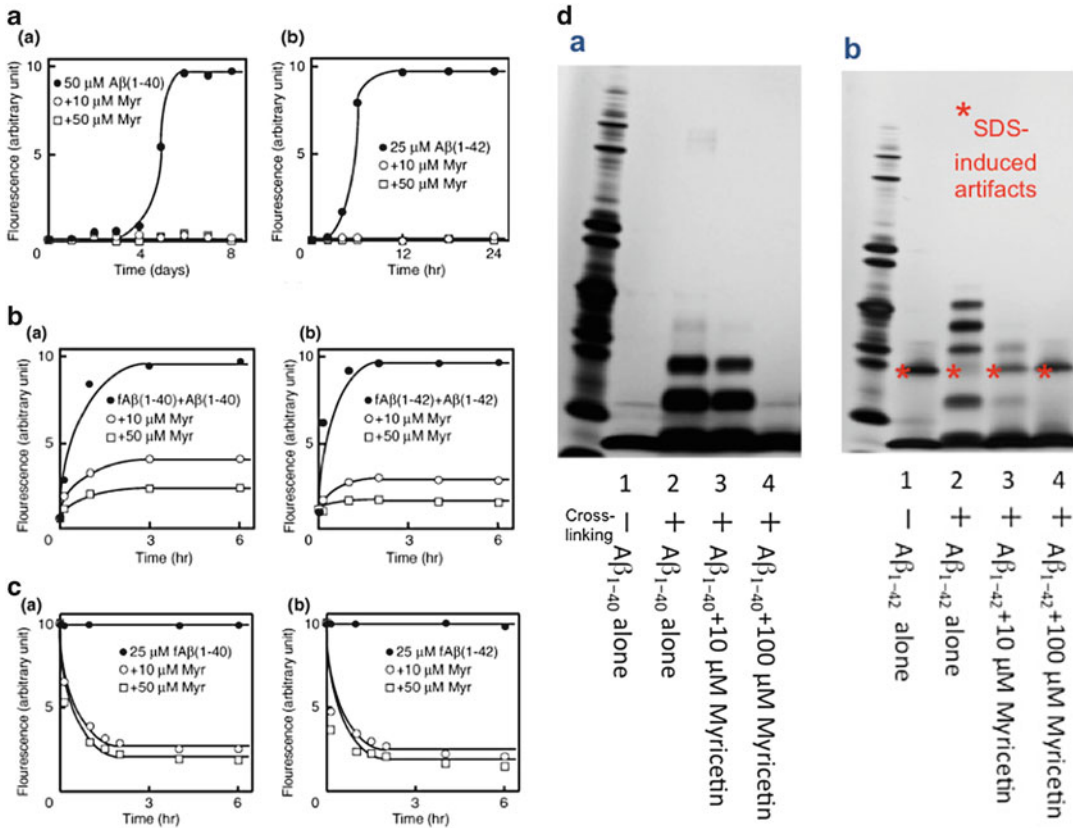


Fig. 4.4 Myricetin inhibits formation of A β fibrils (fA β) from fresh A β (1–40) (A β 40) and A β (1–42) (A β 42) (A), extension of fA β (B), destabilized preformed fA β (C), and inhibit oligomerization of A β 40 and A β 42 (D). The inhibitory and destabilizing effects are also demonstrated

with other methods such as electron microscopy and atomic force microscopy (not shown). Details of the experiments were described in the references (Ono et al. 2003, 2012 [A, B, C: thioflavin T; D: Photo-induced Cross-linking of Unmodified Proteins (PICUP)])

duced cytotoxicity using MTT assays. A β 40 and A β 42 oligomers exhibited cellular toxicity, however, Myr and RA reduced the A β oligomer-induced cytotoxicity.

4.2.2 Studies with *In Vivo* Models of Cerebral Amyloidosis

The natural phenolic compounds with *in vitro* anti-amyloidogenic effects have been tested for the effects in *in vivo* models of cerebral amyloidosis. Reductions of amyloid deposition, A β oligomer levels, inflammation, or oxidative stress in the brain with attenuation of cognitive deterioration have been reported in transgenic

mouse models treated with: Cur (Lim et al. 2001; Yang et al. 2005; Hamaguchi et al. 2009; Ray et al. 2011), EGCG (Rezai-Zadeh et al. 2005, 2008), Myr (Hamaguchi et al. 2009), RA (Hamaguchi et al. 2009), resveratrol (Karuppagounder et al. 2009; Capiralla et al. 2012; Solberg et al. 2014; Porquet et al. 2014), tannic acid (Mori et al. 2012), FA (Mori et al. 2013), rutin (a glycone of quercetin) (Xu et al. 2014), oleuropein (Grossi et al. 2013), hopeahainol A (Zhu et al. 2013), grape seed polyphenolic extract (GSPE) (Wang et al. 2008; Liu et al. 2011), proanthocyanidins of GSPE (Wang et al. 2012), red wine/its polyphenolic contents (Wang et al. 2006; Ho et al. 2009a), anthocyanin-enriched blueberry and blackcurrant

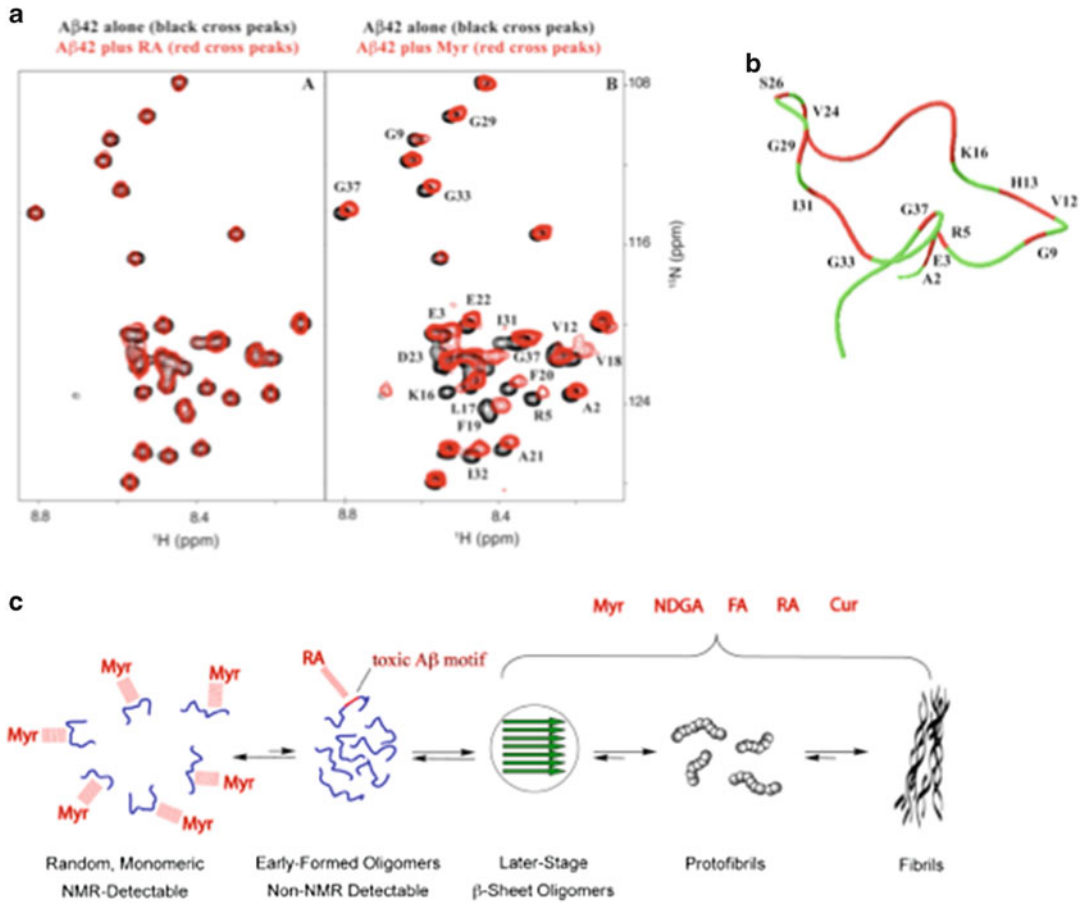


Fig. 4.5 Binding of phenolic compounds to $A\beta$ (a). In nuclear magnetic resonance (NMR) spectroscopy studies, myricetin (Myr) shows NH chemical shift movements indicative of binding (right), but rosmarinic acid (RA) does not (left) (b). A representative structural model of $A\beta_{42}$ that shows binding locations with Myr (indicated by red color) (c). The summary of our studies for

mechanism of polyphenolic inhibition of $A\beta$ aggregation. The phenolic compounds [Myr, RA, nordihydroguaiaretic acid (NDGA), ferulic acid (FA), and curcumin (Cur) (see Fig. 4.3)] exert inhibitory effects through different binding to $A\beta$. Details of the studies were described in the reference (Ono et al. 2012)

extracts (Vepsäläinen et al. 2013), pomegranate juice containing high levels of polyphenols (Hartman et al. 2006), and a natural diet rich in polyphenols and polyunsaturated fatty acids (LMN diet) (Fernández-Fernández et al. 2012). Other models include an $A\beta$ -infused rat AD model treated with Cur (Hoppe et al. 2013), and a transgenic *Caenorhabditis elegans* model of $A\beta$ amyloidosis treated with quercetin (Regitz et al. 2014) and oleuropein (Diomedea et al. 2013; Grossi et al. 2014). Furthermore, attenuation of neuropathology was reported in a tau transgenic mouse model of tauopathy

treated with GSPE (Wang et al. 2010; Santa-Maria et al. 2012). In addition, EGCG prevented the accumulation of αS in 1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine (MPTP)-treated mice, a model of α -synucleinopathy (Mandel et al. 2004).

We focused on the natural phenolic compounds that exerted anti- $A\beta$ aggregation effects in our *in vitro* studies as described above, and systematically investigated whether these five phenolic compounds (Cur, FA, Myr, NDGA, and RA) (Fig. 4.3) also have *in vivo* effects in APP transgenic mice (Tg2576) that show

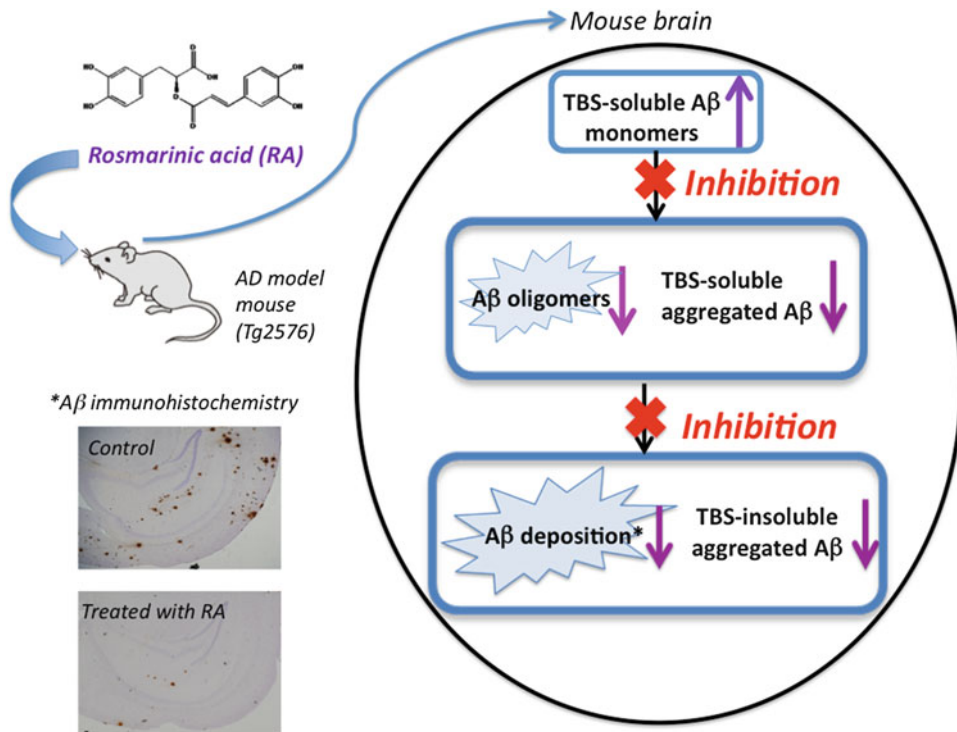


Fig. 4.6 Treatment of Alzheimer's disease (AD) model mice (Tg2576) with rosmarinic acid (RA) showed reductions of both soluble aggregated A β , such as A β oligomers, and insoluble aggregated A β , and an increase of A β monomers in the brain. These findings indicate

that RA inhibits both the steps from A β monomers to oligomers and from oligomers to fibrils. Details of the *in vivo* study on the effects of treatment with diets of the natural phenolic compounds were described in the reference (Hamaguchi et al. 2009)

cerebral A β amyloidosis including parenchymal and vascular amyloid deposition (Hamaguchi et al. 2009). Mice were fed Cur, FA, Myr, NDGA, or RA for 10 months from the age of 5 months. Immunohistochemical analysis, in both the NDGA- and RA-treated groups, revealed that A β deposition was significantly decreased in the brain ($p < 0.05$). In the RA-treated group, the level of soluble A β monomers was increased ($p < 0.01$), while that of oligomers, as probed with the A11 antibody (A11-positive oligomers), was decreased ($p < 0.001$) (Fig. 4.6). However, in the NDGA-treated group, the abundance of A11-positive oligomers was increased ($p < 0.05$) without any change in the levels of soluble or insoluble A β . In the Cur- and Myr-treated groups, changes in the A β profile were similar to those in the RA-treated group, but A β plaque deposition was not significantly decreased. In

the FA-treated group, there was no significant difference in the A β profile. These results showed that oral administration of the natural phenolic compounds influenced AD pathology and A β monomer/oligomer/fibril deposition levels in the brain by differentially modulating A β aggregation pathways *in vivo*. From our results, RA appeared to be the best compound, because it was found to inhibit both steps from monomers to soluble oligomers, and from soluble oligomers to insoluble aggregated A β deposition (Fig. 4.6). Cur and Myr also seemed effective because they significantly decreased soluble oligomer levels, although the reduction of A β deposition did not reach significant levels. FA showed no significant effect. NDGA would be inappropriate, because it significantly increased soluble A β oligomer levels, which would suggest that it inhibited only the step from soluble oligomers to

insoluble aggregated A β deposition, resulting in an increase of potentially toxic soluble oligomers.

4.3 Clinical Trials with Natural Phenolic Compounds for Alzheimer's Disease

For clinical use, several phenolic compounds have been investigated or are under current investigation in clinical trials. Concerning Cur, two clinical trials in AD have been published. In a double-blind, placebo-controlled, randomized, 6-month trial of Cur with 34 AD patients in Hong Kong, 4 g, 1 g (plus 3 g placebo), or 0 g (plus 4 g placebo) of Cur in addition to 120 mg ginkgo leaf extract were orally administered once daily, and showed no significant difference in changes in MMSE or plasma A β 40 levels between 0 and 6 months (Baum et al. 2008). Cur showed no significant side effects in this pilot study (Baum et al. 2008). In another double blind, placebo-controlled, randomized, 24-week trial of Cur in California, 34 patients with AD daily received placebo, 2 g, or 4 g of Curcumin C3 Complex[®] (Ringman et al. 2012). There were no differences between treatment groups in clinical or biomarker efficiency measures including the Alzheimer's Disease Assessment Scale-Cognitive Subscale (ADAS-Cog), levels of A β 40 and A β 42 in plasma, levels of A β 42 and total and phosphorylated tau in cerebrospinal fluid (CSF) (Ringman et al. 2012). For adverse effects, Cur was largely well-tolerated, however, three subjects in the Cur group withdrew due to gastrointestinal symptoms (Ringman et al. 2012). Pharmacokinetic results for Cur and its metabolites suggested limited bioavailability of this compound; levels of native Cur were undetectable in the CSF (Ringman et al. 2012). These published data failed to demonstrate clinical or biomarker evidence of efficacy of a half-year oral Cur intake. Further studies are necessary with a larger number of patients, a longer duration of treatment, and better Cur preparations with higher bioavailability and penetration to the brain.

The website of the U.S. National Institute of Health (NIH) (ClinicalTrials.gov) reports that

clinical studies with Cur for subjects at the stage of MCI are ongoing. In a double blind, randomized interventional study at UCLA, subjects with MCI or age-associated cognitive impairment are recruited and receive Cur (Theracurmin[®] 180 mg/day) or placebo; outcome measures include cognitive testing, amyloid PET, and inflammatory markers. Effects of Cur (800 mg) and yoga in subjects with MCI are investigated in a double blind, randomized trial using interventions by Cur or placebo, and aerobic or non-aerobic yoga.

Regarding resveratrol, to our knowledge, there have been no publications of clinical trials for AD or dementia. The NIH website reports that three clinical trials of resveratrol for AD or MCI are active or completed (ClinicalTrials.gov). A double blind, placebo-controlled, randomized, multi-center study operated by the Alzheimer's Disease Cooperative Study in the U.S. is ongoing; it is scheduled that 120 subjects with AD will be enrolled and receive resveratrol (500 mg to 2 g/day by mouth) or placebo for 52 weeks, and CSF markers and MRI as well as safety and tolerability are primary outcome measures. A single-center, multi-site, randomized, double blind, placebo-controlled 12-month trial of liquid resveratrol with glucose and malate to slow the progression of AD in New York has been completed with enrollment of 27 AD subjects, but no results are posted. To test enhancement of memory functions in subjects with MCI by dietary interventions and in combination with exercise and cognitive training, a double blind, placebo-controlled, randomized trial in Germany is ongoing with multiple arms that include a group of resveratrol supplementation.

Clinical trials with EGCG for AD or Down syndrome are also posted on the NIH website (ClinicalTrials.gov). A double-blind, placebo-controlled, randomized, 18-month trial of EGCG in early or mild AD is ongoing in Germany; it is planned that 50 patients will be recruited and receive EGCG (200–800 mg) or placebo added to donepezil with evaluation of cognitive functions. Older subjects with Down syndrome show progression of AD-like lesions in the brain. To test improvement of cognitive performance and deceleration of AD-like progression in

Down syndrome by EGCG, a double-blind, placebo-controlled, randomized study is ongoing in Spain; the participants receive a daily oral dose containing 9 mg/kg of EGCG or placebo, and changes in cognitive functions and amyloid markers are primary outcome measures.

In addition, two clinical trials of isoflavones (including genistein) in AD and two interventional trials of pomegranate polyphenol extract or juice in non-demented subjects are ongoing (ClinicalTrials.gov).

Our group started clinical studies with RA, based on the results of our *in vitro* and *in vivo* studies with models of A β amyloidosis (Ono et al. 2004b, 2012; Hamaguchi et al. 2009). First, we completed a double-blind, placebo-controlled, randomized trial in healthy individuals to reveal pharmacokinetics, safety, and tolerability of RA. Next, we are conducting a double-blind, placebo-controlled, randomized trial of RA for mild AD with investigations of cognitive functions and biomarkers including amyloid PET and CSF markers.

Targets of future clinical trials with natural phenolic compounds will extend to other cerebral amyloidoses or protein aggregation disorders than AD, including CAA, PD, DLB, and non-AD tauopathies.

4.4 Conclusions

Epidemiological studies suggest an association of diets rich in phenolic compounds or polyphenols (Mediterranean diet, red wine, green tea, etc.) with reduction of risk of dementia or AD. In addition to the general beneficial effects of these compounds such as anti-oxidant and anti-inflammatory properties, experimental studies indicate that natural polyphenols have specific effects on pathways involved in the pathophysiology of cerebral amyloidosis; the effects include modulation of APP processing, inhibition of A β aggregation and destabilization of aggregates, promotion of A β degradation/clearance, alleviation of A β -induced oxidative stress, leading to reductions of amyloid deposition, A β oligomer levels and inflammation in the brain, with at-

tenuation of cognitive deterioration in treated animal models. For clinical use, several phenolic compounds are under investigation by clinical trials for AD or MCI, although no compounds have been yet proved to have certain therapeutic or preventive effects so far. Further clinical trials and preventive interventions of these phenolic compounds with efforts to improve oral bioavailability and brain penetration are necessary to establish their efficacy in AD and other human cerebral amyloidoses.

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Abstract

Healthy brain aging and the problems of dementia and Alzheimer's disease (AD) are a global concern. Beyond 60 years of age, most, if not everyone, will experience a decline in cognitive skills, memory capacity and changes in brain structure. Longevity eventually leads to an accumulation of amyloid plaques and/or tau tangles, including some vascular dementia damage. Therefore, lifestyle choices are paramount to leading either a brain-derived or a brain-deprived life. The focus of this review is to critically examine the evidence, impact, influence and mechanisms of natural products as chemopreventive agents which induce therapeutic outcomes that modulate the aggregation process of beta-amyloid (A β), providing measureable cognitive benefits in the aging process. Plants can be considered as chemical factories that manufacture huge numbers of diverse bioactive substances, many of which have the potential to provide substantial neuroprotective benefits. Medicinal herbs and health food supplements have been widely used in Asia since over 2,000 years. The phytochemicals utilized in traditional Chinese medicine have demonstrated safety profiles for human consumption. Many herbs with anti-amyloidogenic activity, including those containing polyphenolic constituents such as green tea, turmeric, *Salvia miltiorrhiza*, and *Panax ginseng*, are presented. Also covered in this review are extracts from kitchen spices including cinnamon, ginger, rosemary, sage, salvia herbs, Chinese celery and many others some of which are commonly used in herbal combinations and represent highly

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promising therapeutic natural compounds against AD. A number of clinical trials conducted on herbs to counter dementia and AD are discussed.

Keywords

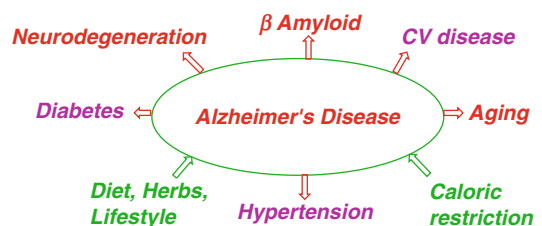
Alzheimer's disease • Dementia • Amyloid-beta • Traditional Chinese medicine (TCM) • Herbal polyphenols

5.1 Beyond the Molecular Frontier – The Threats of Our Age

During the past hundred years, treatments for human diseases have helped raise life expectancy significantly. However, an aging population brings increased burdens and costs to individuals and society from age-related cognitive decline; indeed, the latter has emerged as one of the major health threats and challenges of our age. In another 36 years there will be triple the number of persons 80 years or older, with approximately 50 % of adults over 85 years afflicted with Alzheimer's disease (AD). The total number of new cases of dementia each year worldwide is nearly 7.7 million, which translates to 15 new cases every minute (International 2012). Estimates indicate that between 2 and 10 % of all cases of dementia appear before the age of 65. Advancing age is the highest risk factor for AD, with age-specific prevalence nearly doubling every 5 years beyond the age of 65. The financial estimated worldwide cost of dementia was \$604 billion in 2010 (Wimo et al. 2013). Unless we act now, by 2050 the problem will be unmanageable. Recent advances in the biology of aging in model organisms, together with molecular and multidisciplinary studies of neurodegenerative and aging-related disease risks and personal practices (outlined in Scheme 5.1), are beginning to uncover these mechanisms and their potential roles in cognitive decline (Witte et al. 2009; Villeda et al. 2011)

Interrelationships between aging, apolipoprotein E (*APOE*) ϵ 4 allele, oxidative damage, reactive oxygen species (ROS), amyloid metabolism/toxicity and neurodegenerative dysfunctions leading to dementia and AD

are highly probable. Nevertheless, the precise mechanisms remain unknown. Ideally, the opportunities for making lifestyle, diet and nutritional choices to enhance human brain and body function is available and practiced by many (Gomez-Pinilla and Tyagi 2013). The theme of positive aging is to be proactive in minimizing/preventing cognitive decline and disease. Dementia and AD research priorities have also advanced from simply considering clinical symptoms. The focus is now more on early detection of the pre-symptomatic phase and the prevalence of early dementia signs, as these are considered to be potential windows of opportunity for successful therapeutic interventions and preventions. For instance, recent research supports mounting evidence implicating dysfunctional lipid metabolism in the pathophysiology of AD indicating that lipid biomarkers have the potential to predict memory impairment at a preclinical stage of AD. Changes in the blood profile of a set of ten lipids critical for proper cell membrane structure and function in elderly persons who showed no signs of cognitive problems, predicted they would go on to develop either mild memory impairment or AD within 2–3 years, with greater than 90 % accuracy (Mapstone et al. 2014).



Scheme 5.1 The interventions and disease risks related to Alzheimer's disease

Humans are able to consume a vast range of foodstuffs. However, the ready availability and low cost of food, and the freedom of being able to eat anything, does not mean that we should maximize eating practices to eat everything (Ulijaszek et al. 2012). The diet-related chronic diseases of modern society are now the single largest cause of death encompassing diabetes, cardiovascular disease, hypertension, obesity and cognitive decline (Scheme 5.1). For foods to promote the health of our aging, physical frailty and mental state, we need to reduce the consumption of processed foods and fatty diets, with negative nutritional attributes such as high-energy refined sugars, saturated fats and high sodium content, whilst increasing affinity and tendency to consume those with positive health attributes including phytochemicals and micronutrient rich foods.

5.2 Herbal Polyphenols – Modulation of Oxidative Stress, Dementia and AD

From our previous analysis of well-designed, randomized double-blind controlled trials on Chinese herbal medicines beneficial for the improvement of cognitive function, we found that neuroprotective benefits of suppression of oxidative stress as the most common feature provided by single herbs or herbal mixtures (May et al. 2009, 2012).

5.2.1 Epigallocatechin-3-Gallate

Oxidative stress may directly initiate neurodegeneration, and herbal antioxidant neuroprotection is considered as a preventative and therapeutic approach (Hugel et al. 2012). Crucially, the scientific evidence confirms that the majority of herbal polyphenolic compounds have a good safety profile, are affordable and are globally readily available to significantly reduce the burden of dementia and AD.

It has been known for at least a decade that polyphenols possess anti-amyloidogenic activity. A diverse range of herbal polyphenolic

constituents including tannic acid, quercetin, kaempferol, curcumin, catechin and epicatechin are known to dose-dependently inhibit the formation of amyloid-beta ($A\beta$) fibrils as well as their elongation. Importantly, polyphenols can bind directly to $A\beta$ or mature aggregates and impair their stability. Epigallocatechin-3-gallate (EGCG), a major component of green tea, significantly inhibits $A\beta$ aggregation and has the ability to remodel large $A\beta$ fibrils into smaller aggregates that are non-toxic (Wang et al. 2010). The gallate functionality in EGCG is critical in facilitating the reduction of $A\beta$ and increasing APP α -proteolysis. Evidence has indicated that EGCG reduces $A\beta$ production in both neuronal and mouse AD models in concert with activation of anti-amyloidogenic amyloid precursor protein (APP) α -processing. An extensive screening of the effect of other gallate-containing phenolic compounds on APP anti-amyloidogenic processing found that long chain gallate esters (Zhang et al. 2013b) such as octyl gallate (OG; 10 mM), a commercial food antioxidant, drastically decreased $A\beta$ generation, in concert with increased APP α -proteolysis in murine neuron-like cells transfected with human wild-type APP or “Swedish” mutant APP. OG markedly increased production of the neuroprotective amino-terminal APP cleavage product, soluble APP- α (sAPP α). OG increases anti-amyloidogenic APP α -secretase processing by activation of ER α /PI3k/Akt signaling and ADAM10. Fish oil has been shown to have a synergistic effect in combination with EGCG, with co-treatment leading to a reduction in $A\beta$ plaque formation and levels of $A\beta$ (1–40) and $A\beta$ (1–42) in AD transgenic Tg2576 mice (Giunta et al. 2010). The potential role of polyphenols in neurodegeneration and the pathogenesis of AD has expanded with discoveries that they can modulate a class of proteins called sirtuins that are involved in longevity and cell survival (Jayasena et al. 2013) (Table 5.1).

EGCG has numerous health-promoting effects (Hugel and Jackson 2012) including anti-cancer, antioxidant, anti-inflammatory, anti-diabetic, anti-aging and in particular its $A\beta$ -sheet disruption (Palhano et al. 2013) capacity

Table 5.1 Anti-amyloidogenic activity of polyphenols and herbal extracts

Polyphenol/herbal extract	Anti-amyloidogenic activity
Investigation of the ability of EGCG to inhibit the formation of metal-free or metal-associated A β (1–40) aggregates	EGCG interacted with Cu(II)- and Zn(II)-A β monomer, dimer species. Formed more compact peptide conformations compared to EGCG-untreated A β species; ternary EGCG–metal–A β complexes were produced. This illustrates the selective modulation of the anti-amyloidogenic reactivity of EGCG towards metal-A β species (Hyung et al. 2013)
Effect of the addition of EGCG in drinking water (1.5, 3 mg/kg for 3 weeks) intake in mice	Prevented lipopolysaccharide-induced A β production by the inhibition of β -secretase activity, and improved effects on memory deficiency in liposaccharide-induced AD mice models (Lee et al. 2009)
Isothermal titration calorimetry studies on the interactions between EGCG and A β	EGCG-A β binding was enhanced by increasing temperature, salt concentration and at pH values away from the pI of A β (Wang et al. 2010)
EGCG encapsulated in nanoparticles	Improved <i>in vivo</i> efficacy, doubled bioavailability; improved chemical stability and enhanced its biological activity (Li et al. 2012; Hu et al. 2013; Smith et al. 2010)
Protonation of EGCG at low pH	Resulted in aggregation and reduced oral bioavailability of EGCG-dispersed selenium nanoparticles (Wu et al. 2013b)
Modulation of A β -induced tau hyperphosphorylation by curcumin (Cur) in human neuroblastoma SH-SY5Y cells	Cur inhibits phosphorylation of tau at Thr231 and Ser396 by modulating the phosphatase and tensin homolog (PTEN) PTEN/Akt/GSK-3 β pathway. Involves down-regulation of phosphorylation of Akt and of PTEN, a negative regulator of PIP3 induced by A β (Huang et al. 2014a)
Effects of Cur after 3-month administration to <i>APP^{Swe}/PS1^{dE9}</i> double transgenic mice, an AD model	Reduced A β (1–40) and A β (1–42) levels, and aggregation of A β -derived diffusible ligands in the mouse hippocampal CA1 area; enhanced expression of γ -secretase; increased expression of β -amyloid-degrading enzymes, including insulin-degrading enzymes and neprilysin (Esatbeyoglu et al. 2012; Wang et al. 2014)
Testing of Cur -based fluorescence imaging probes <i>in vitro</i> and <i>in vivo</i>	Near-infrared fluorescence imaging with the Cur analogue CRANAD-58 revealed interaction with A β in mouse brain; CRANAD-17 was capable of inhibiting A β 42 cross-linking induced by copper (Zhang et al. 2013c)
Targeting of endogenous neural stem cells by Cur -encapsulated nanoparticles	Cur nanoparticles: increase neuronal differentiation by activating the Wnt/ β -catenin pathway in hippocampal neural stem cells; involved in regulation of neurogenesis; rescued learning and memory impairments in an A β -amyloid induced rat model of AD (Tiwari et al. 2014)
Studies on the brain accessibility of Cur -lipid-nanoparticles	High affinity for A β in post-mortem brains samples of AD patients (Mourtas et al. 2014). Cur-loaded solid lipid nanoparticles showed 30 times higher preferential distribution into the brain (Kakkar et al. 2013)
Anti-amyloidogenic effect of an ethanol extract of Magnolia officinalis : 12.9 % magnolol, 16.5 % honokiol, 16.6 % 4-O-methylhonokiol, plus 42–45 % of other constituents	Administration of 10 mg/kg extract for 3 months inhibited amyloidogenesis, reduced A β accumulation via β -secretase 1 (BACE1) inhibition in the brain of Tg2576 mice with memory improving effects (Lee et al. 2012)
2,2',4'-trihydroxychalcone Glycyrrhiza glabra	Anti-oxidative, anti-tumor, <i>in vitro</i> inhibition of BACE1 bioactivity with IC ₅₀ 2.5 μ M, reduced A β formation in mice-AD studies (Zhu et al. 2010)

(continued)

Table 5.1 (continued)

Polyphenol/herbal extract	Anti-amyloidogenic activity
Isobavachalcone, bavachinin isolated from <i>Psoraleae Fructus</i>	Contains compounds that inhibit BACE1 (Choi et al. 2008). Isobavachalcone inhibits A β oligomerization and fibrillization, bavachinin transforms A β into non-toxic aggregates (Chen et al. 2013)
Tenuifolin, a triterpenoid saponin isolated from <i>Polygala tenuifolia</i>	2.0 μ g/mL tenuifolin significantly decreased A β -secretion from COS-7 cells without altering the ratio of A β (1–40) and A β (1–42) by BACE1 inhibition (Lv et al. 2009)
Effect of the <i>Polygonum multiflorum</i> extract component 2,3,5,4'-tetrahydroxystilbene-2-O- β -D-glucoside (TSG) on the rat A β model	Administration of TSG rescued A β (1–42) induced impairment in learning and memory, protecting synaptic structures and function; the up-regulation of Src and NR2B may be responsible for the improved learning and anti-AD properties (Zhou et al. 2012)
Effect of ethanol extract of <i>Polygonum multiflorum</i> in mouse neuroblastoma cells expressing Swedish APP (N2a-SweAPP)	Potent reduction in A β production through APP modulation, with the up-regulation of sAPP α and down-regulation of sAPP β (Liu et al. 2012)
<i>Salvia miltiorrhiza</i> lipophilic constituents: Tanshinone I (TI), Tanshinone (IIA)	Molecular dynamics simulations reveal that TI and TIIA preferentially bind to a hydrophobic β -sheet groove. TI was better than TIIA for inhibition amyloid- β aggregation; the tanshinones also affected disaggregation of amyloid fibrils, and protection of cultured cells (Wang et al. 2013)
<i>Salvia miltiorrhiza</i> water-soluble constituents: Danshensu and Salvianolic acid B	Protected PC-12 cells by blocking A β (25–35) induced Ca ²⁺ intake, lactate dehydrogenase release, cell viability decrease and apoptosis (Zhou et al. 2011)
Danshen extract (danshensu 40 mg/kg, protocatechuic aldehyde 149 mg/kg, and salvianolic acid B 50 mg/kg) was administrated intragastrically in rats	From blood and brain microdialysates collected at 15 and 30 min time intervals, danshensu and protocatechuic acid (oxidative metabolite of protocatechuic aldehyde) could be detected in the blood and brain (Zhang et al. 2011)
Examination of Salvianolic acid B (Sal B) on human islet amyloid polypeptide (hIAPP) aggregation and phototoxicity	Sal B significantly inhibited the formation of hIAPP amyloid and disaggregated hIAPP fibrils. Cytoprotective effects by Sal B on pancreatic INS-1 cells (Cheng et al. 2013a)

(outlined in Scheme 5.2). The major research challenge concerning the anti-amyloidogenic benefits of polyphenol-containing herbs and foods is to enhance their bioavailability and brain permeability (Schaffer and Halliwell 2012; Singh et al. 2008; Green et al. 2007; Lambert et al. 2006; Smith et al. 2010; van Duynhoven et al. 2011). Furthermore, the bioavailability of polyphenols from dietary input is highly variable between individuals and generally far too low to explain their bioactive antioxidant effects *in vivo* (Lotito and Frei 2006).

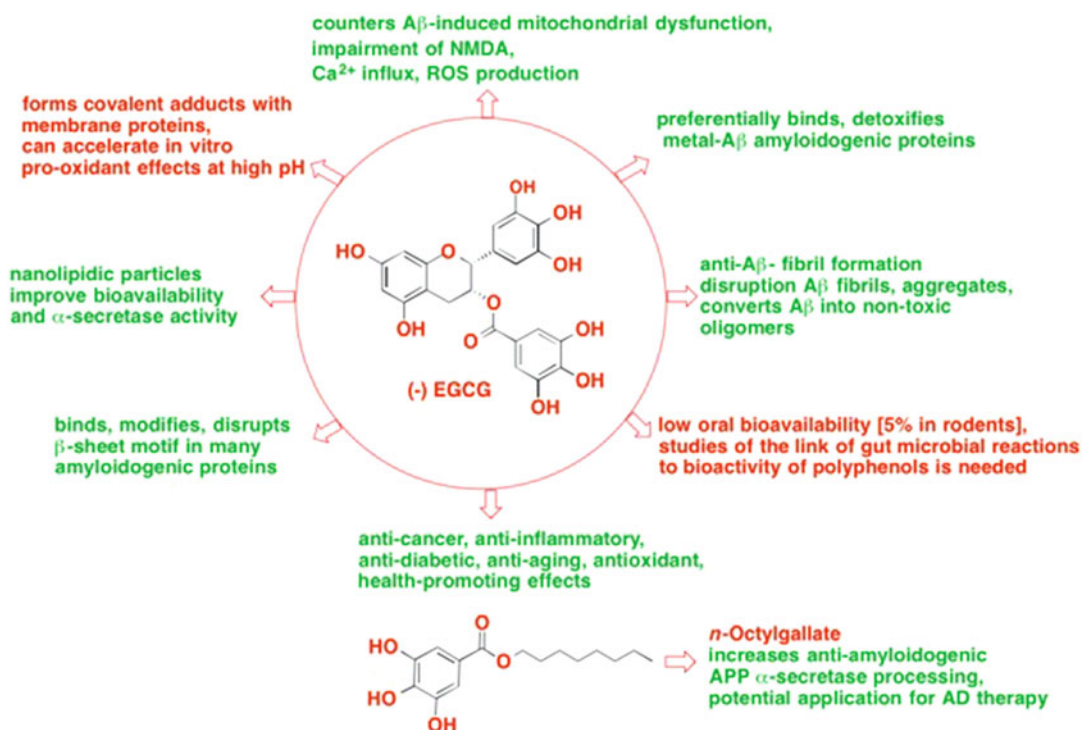
5.2.2 Curcumin

Cur is a promising neuroprotective anti-AD natural product that however has poor

brain bioavailability with incompletely defined therapeutic mechanisms. Its antioxidant, anti-inflammatory properties have been extensively documented (Esatbeyoglu et al. 2012; Wang et al. 2014). Cur-nanoparticles with improved brain permeability induced adult neurogenesis through activation of the canonical Wnt/ β -catenin pathway, and may provide opportunities for treating AD by enhancing a brain self repair mechanism (Zhang et al. 2013c).

5.2.3 *Magnolia officinalis*

The herbal constituents shown in Fig. 5.1 from *Magnolia officinalis* and other members of the *Magnoliaceae* family have diverse therapeutic applications (Lee et al. 2011b). The neolignan



Scheme 5.2 The multiple therapeutic applications of green tea constituent EGCG

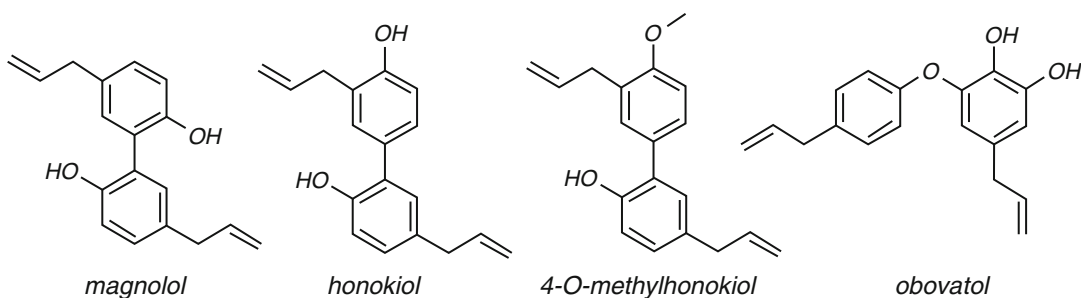
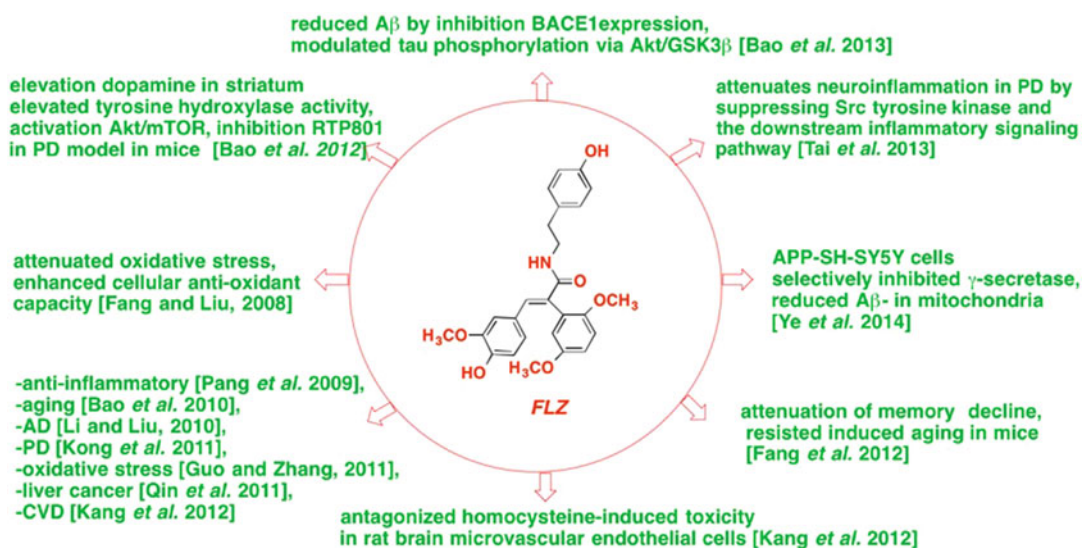


Fig. 5.1 Major bioactive constituents found in *Magnolia officinalis*

4-*O*-methylhonokiol is a potent cannabinoid receptor type-2 (CB2) ligand and has been found to attenuate memory impairment in presenilin 2 mutant mice through reduction of oxidative damage and inactivation of astrocytes and the extracellular signal-regulated kinase (ERK) pathway (Lee et al. 2011a). The various neuroprotective and anti-Alzheimer disease effects reported in rodent models (Lee et al. 2011a) may be mediated via CB2 receptors, providing evidence that the compound should be bioavailable in the brain.

5.2.4 *Annona glabrais* – Squamosamide Derivative (FLZ Compound)

Traditional Chinese medicine makes use of several constituents from the leaves and roots of *Annona glabrais*, including a natural squamosamide. Importantly, the squamosamide derivative FLZ showed enhanced antioxidant activity; in APP-SH-SY5Y expressing cells it selectively inhibited γ -secretase activity without



Scheme 5.3 Anti-amyloidogenic properties of compound FLZ, a squamoside analogue of a constituent from *Annona glabrais*

modulating the Notch pathway (Ye et al. 2014). The many positive anti-amyloidogenic studies suggest FLZ may have therapeutic potential for the treatment of AD (illustrated in Scheme 5.3) (Fang et al. 2012; Kang and Zhang 2012; Pang et al. 2009; Li and Liu 2010; Kong et al. 2011; Qin et al. 2011; Fang and Liu 2008; Bao et al. 2012, 2013; Tai et al. 2013)

5.2.5 Ginseng

The available types of ginseng, all belonging to the *Araliaceae* family, are Asian ginseng (*Panax ginseng*), American ginseng (*P. quinquefolius*) and Siberian ginseng (*Eleutherococcus senticosus*). Water extracts of the dried roots and leaves of *Panax ginseng* have been used as a stimulant/tonic, diuretic and digestive aid in traditional Chinese medicine for over 2,000 years. Ginseng phytomedicines are sold as ergogenic supplements to enhance mental and physical performance – reflective of Chinese medicine where body and mind are inseparable – to provide resistance to stress, and to prevent ‘exhaustion’ and disease. The major active principles of *P. ginseng extracts* are ginsenosides,

which are glycosylated derivatives of the triterpene dammarane such as for instance Rg₁. Rg₃ is one of the major constituents of ginseng. The ginsenosides that reduce Aβ levels in animal models and other *in vitro* studies are summarized in Table 5.2.

The diverse constituents and multiple actions of ginseng constituents in the CNS reviewed recently (Kim et al. 2013a) will not be elaborated here. The *in silico* analysis of 12 ginsenosides (see Table 5.2) revealed those with potential interactions with the BACE1 receptor active site essential for enzyme inhibition (Karpagam et al. 2013). Further studies included ADMET screening to find the drug-like ginsenosides with a specific ability to cross blood brain barrier (BBB), and to determine safety/toxicity. Also the BACE1-ginsenosides complexes were further subjected to a molecular dynamics simulation to study their stability and hydrogen bond interactions. Of the 12 ginsenosides, CK, F₁, Rh₁, and Rh₂ were predicted to pass the BBB and ADMET analysis predicted toxic effects for ginsenosides Ro and ginsenoside Rg₁, while Rf showed low oral absorption in human gastrointestinal tract. These results suggest that of the seven ginsenosides demonstrating BACE1

Table 5.2 The anti-AD bioactivities of *P. ginseng* constituents

<i>Panax ginseng</i> AD cognitive effects	Anti-A β bioactivities
	Ginsenoside Rg₃ inhibited γ -secretase activity in mouse model AD
	A β lowering by modulation/reduction of lipid kinase PI4KII α activity (Kang et al. 2013)
	Rg₃ enhanced neprilysin (NEP, rate-limiting enzyme in A β degradation) gene expression. Caused a reduction in A β (1–40) and A β (1–42). (Yang et al. 2009)
Fermented red ginseng – ginsenoside Rh ₂ neuroprotective effects. Inhibited ischemia reperfusion brain injury in rats (Bae et al. 2004)	
	<i>P. notoginseng</i> modulates protein, gene expression related to α - and β -secretases. Reductions in levels of β -secretase resulting in decline of A β generation (Huang et al. 2014b)
Fermented ginseng (FG) ameliorated memory impairment in transgenic mouse model of AD	Brain soluble A β (1–42) levels measured from the cerebral cortex of transgenic mice were significantly reduced by the FG extract treatment (Kim et al. 2013b)
	Commercially-available preparations of ginseng Rg₁ , Rg₃ , and RE , resulted in significant reductions in the amount of A β (1–42) detected in the brains of animals after single oral doses of these agents (Chen et al. 2006)
Oral administration of ginsenoside Rb₁ to mice stressed with acute immobilization; Rb ₁ modulated stress effects by attenuating the stress-induced increase in neurosteroids (Lee et al. 2006a)	
Oral administration of Rg₃ and Rb₁ to mice stressed with acute immobilization; both lowered levels of the stress-marker putrescine (Lee et al. 2006b)	
Ginsenoside Rg₁ improved learning & memory in rat model of AD (Quan et al. 2013)	Rg₁ inhibits the transcription and translation of BACE1, suppresses the activity of BACE1, and ultimately attenuates A β generation (Chen et al. 2012)
	Rg₁ promoted α -secretase cleavage of APP via estrogenic activity, indicating that it may be useful in the prevention of AD, in particular in postmenopausal females (Shi et al. 2013)
Rg₁ , applied to primary cultured cortical neurons, rescued A β -mediated mitochondrial dysfunction	May attenuate A β -induced neuronal death through the suppression of intracellular mitochondrial oxidative stress (Huang et al. 2012)
Rd attenuated β -amyloid-induced pathological tau phosphorylation	Enhanced the activity of protein phosphatase 2A (PP-2A) involved in tau dephosphorylation (Li et al. 2013a)
<i>In silico</i> approach for discovery of BACE1 inhibitors from <i>Panax ginsenosides</i> included Rb ₁ , Rd, Rf, Re, Rg ₁ , Rg ₂ , Rg ₃ , Ro, Rh ₁ , Rh ₂ , CK, and F1	Rh ₁ , Rh ₂ , CK, F1 passed the criteria of: molecular docking-evaluated interaction with BACE1 receptor proteins, complex stability, H-bond interactions, ADMET for BBB permeability, having no toxicity (Karpagam et al. 2013)
Ginsenoside Rg₅ effect on cognition and beta-amyloid deposition in STZ-induced memory impaired rats	Rg ₅ (5, 10 and 20 mg/kg) improved cognitive dysfunction in rats which was related to attenuating neuro-inflammatory responses with decreased brain levels of inflammatory cytokines TNF- α , IL-1 β ; Congo Red staining and Western blot analysis showed decreased A β deposits (Chu et al. 2014)

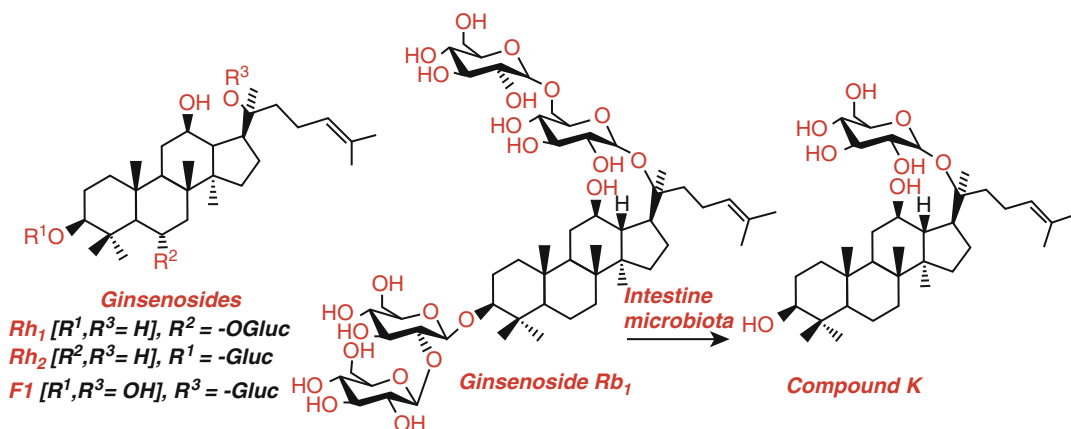


Fig. 5.2 The structures of ginsenosides *Rb*₁ and its metabolic transformation product K, and those of *Rh*₁, *Rh*₂, *F*₁

inhibition, only the four monoglycosylated ginsenosides CK, *Rh*₁, *Rh*₂, and *F*₁ pass the BBB and possess satisfactory drug-like properties. BACE1 and ginseng inhibitor complex crystal structural data to describe their binding modes would provide an accurate picture of the number and length of hydrophobic and hydrogen bond ginsenoside-enzyme interactions. These two descriptors have reliably predicted the activity of synthetic BACE1 inhibitors (Nastase and Boyd 2012). The wider implications of this research are that the brain-permeation/bioactivity of di- and multi-glycosylated ginsenosides is questionable. Intestinal microbial metabolism (Zhang et al. 2013d) similar to that of *Rb*₁ shown in Fig. 5.2 may be a pre-requisite for their neuroprotective activity.

5.2.6 Herbal Foods, Formulations and Supplements

L-3-*n*-Butylphthalide (Fig. 5.3) was first extracted from Chinese celery (*Apium graveolens* var. *secalinum*). The chemically prepared compound is used as an anti-hypertensive herbal medicine for the treatment of ischemic stroke, and has therapeutic application for the prevention of vascular dementia by up-regulation of Akt expression in the hippocampus (Huai et al. 2013; Peng et al. 2008, 2012). Potassium 2-(1-hydroxypentyl)-benzoate (dl-PPPB), a precursor

to *n*-butylphthalide, has neuroprotective effects on cerebral ischemic, vascular dementia and A β -induced animal models by inhibiting oxidative injury, neuronal apoptosis and glial activation. Further research has suggested that dl-PPPB could be an attractive multi-target neuronal protective agent for the treatment of AD (Zhao et al. 2013; Peng et al. 2014). *Z*-ligustilide found in *R. angelica sinensis* promotes the activities of superoxide dismutase and thereby reduces oxidative stress in brain tissues; protects against A β -induced neurotoxicity and is a potential therapeutic against vascular dementia (Huang et al. 2008; Kuang et al. 2006; Feng et al. 2012; Xin et al. 2013). An appreciation of the amount of *Z*-ligustilide, the bioactive component in 10 g of herb is detailed in Fig. 5.3. The pharmacokinetics and bioavailability of *Z*-ligustilide were determined by the systematic investigation in Sprague–Dawley rats. With an extraction efficiency of 62.3 %, 0.93 g *Z*-ligustilide was isolated from 100 g of *R. angelica sinensis*. Therefore, based on animal pharmacokinetic data, with the absolute bioavailability at a 50 mg/kg dose of 75.44 %, a single medicinal use of 10 g of the herb may deliver 43.7 mg of *Z*-ligustilide.

Studies on 27 herbs revealed that some lesser known herbs such as *Curcuma aromatica* and *Zingiber officinale* (ginger) extracts effectively protected cells from A β insult, followed by *Ginkgo biloba* (ginkgo), *Polygonatum*

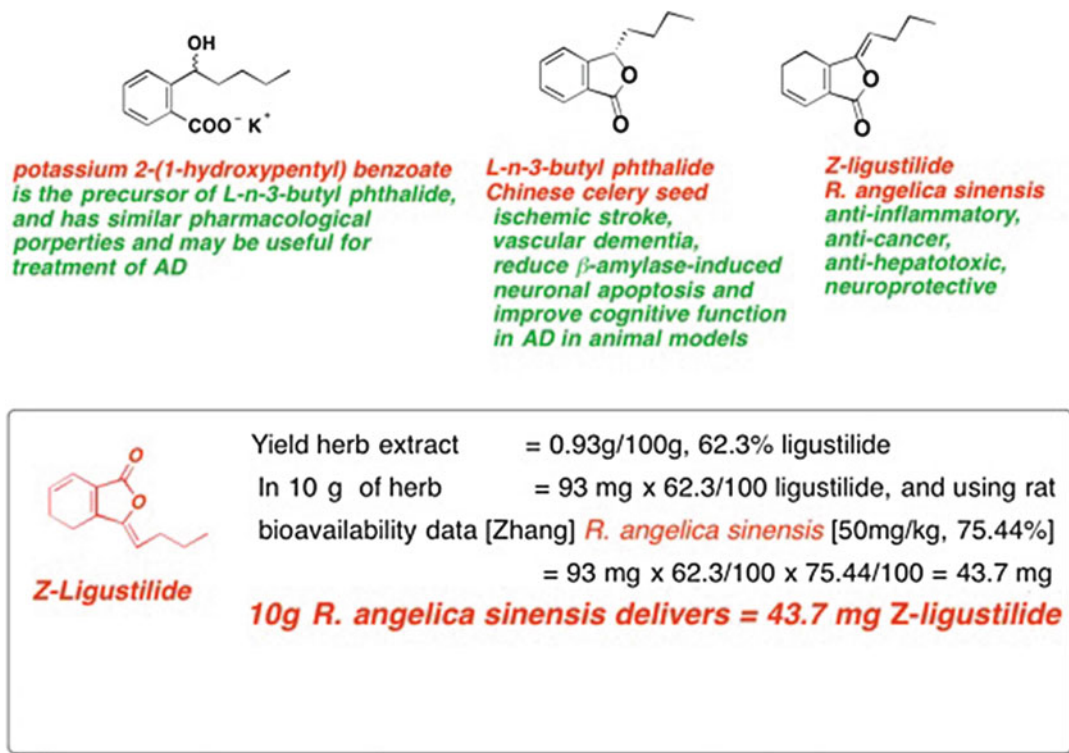


Fig. 5.3 Herbal bioactive compounds and *Z-ligustilide* bioavailability calculations (Zhang et al. 2014)

sp., *Cinnamum cassia* (Chinese cinnamon), *Rheum coreanum* (Korean rhubarb), *Gastrodia elata* (gastrodia), and *Scutellaria baicalensis* (skullcap) (Kim et al. 2007). With regards to herbs, spices and food products that disrupt, destabilize or reverse amyloid aggregation, these have been investigated for their ability: (i) to detour the generation of toxic amyloid precursors (off-pathway); (ii) to prevent the assembly of amyloid oligomers into fibrils; (iii) to inhibit fibril growth and deposition; (iv) to disassemble preformed fibrils; and (v) to promote A β clearance. The structures of the active anti-dementia constituents in Chinese herbs most widely used and investigated as potential amyloid inhibitors are presented in Fig. 5.4.

Many herbs are considered to be responsible for multiple beneficial effects such as improving vascular dementia, energy homeostasis, improving mitochondrial antioxidant capacity, and anti-inflammatory neuroprotection. The many and varied constituents in herbs can also

enhance the bioavailability and bio-effectiveness of the active constituents and thus have more therapeutic value than individual compounds. Preliminary animal model studies suggest that antioxidants in spearmint and rosemary might be useful in modulating age-associated cognitive decline. Furthermore, rosemary improves local blood circulation, relieves pain, has anticancer activity, and controls blood lipid and anti lipid peroxidation. Carnosic acid, one of the major phenolic constituents of rosemary, is a pro-electrophile specifically activated by the oxidative stress pathological state resulting in its conversion from the hydroquinone to the oxidized quinone form, before it activates the Keap1/Nrf2 pathway leading to gene induction of the antioxidant response element (ARE) and gene products that protect against oxidative stress. A survey of Chinese herbs and herbal formulas that improve cognition in dementia rated the following as the top 10 herbs for improving memory: *Poria cocos*, *Radix et*

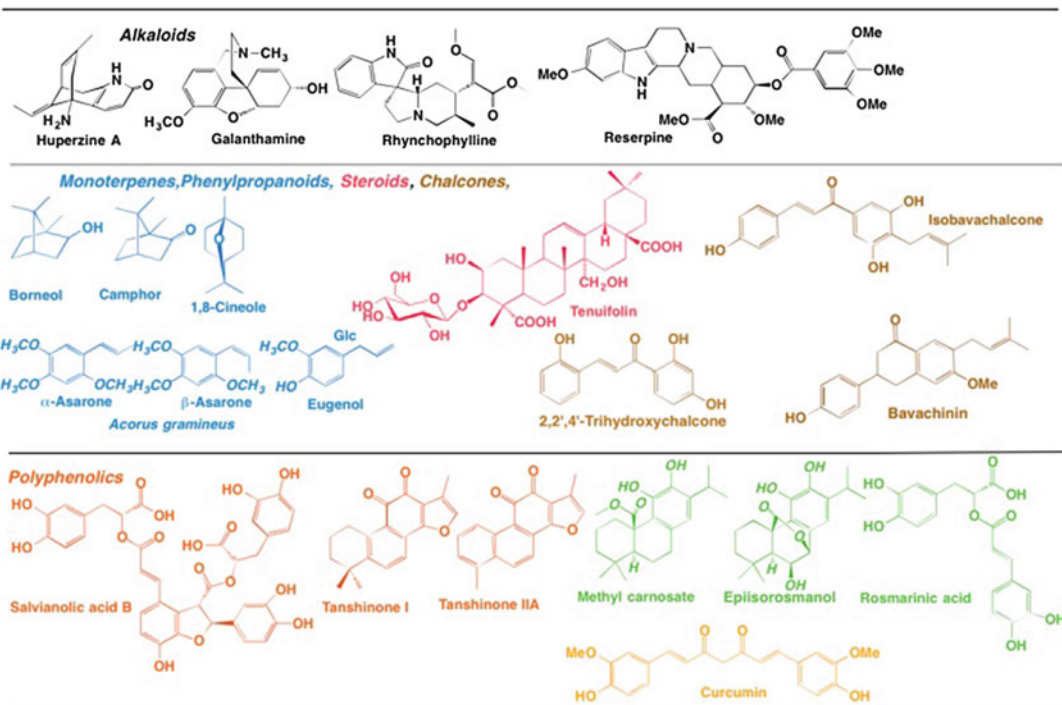


Fig. 5.4 Structures of the major chemical families of active constituents found in Chinese herbs having anti-dementia and β -amyloid anti-aggregation activities

rhizome ginseng, Radix polygalae, Radix et rhizome glycyrrhizae, Radix Angelica sinensis, Rhizoma acori tatarinowii, Semen ziziphi spinosae, Radix rehmanniae, Radix ophiopogonis and Rhizoma zingiberis (Lin et al. 2012; Shen and Chen 2013). The anti-A β bioactivity and neuroprotective mechanisms of many of these herbs are outlined in Table 5.3. In Schemes 5.4 and 5.5, the focus is on the particular herbs and spices that can effectively protect against amyloid disease. Their A β disaggregation properties and inhibition of tau protein hyperphosphorylation are highlighted (Yoshida et al. 2014; Xian et al. 2012; Fujiwara et al. 2006; Frydman-Marom et al. 2011; Kumaraswamy et al. 2013; Airoidi et al. 2013; Zeng et al. 2013).

5.2.7 Chinese Herbal Formulae for Anti-dementia Protection

Baicalin, jasminoidin, and cholic acid structures (Fig. 5.5) are the main active components of

Qingkailing (QKL, Scheme 5.6). QKL is one of the most well-known Chinese herbs and is an aqueous preparation containing extracts of 7 herbs (Cheng et al. 2012). Baicalin is a strong antioxidant; jasminoidin elicits a protective effect on neurons under a broad range of stresses and cholic acid strongly promotes the expression of growth factors in the brain. Upon further investigation of the therapeutic effects and molecular mechanisms of a combination of the three components baicalin, jasminoidin and cholic acid (CBJC) in a rat dementia model, it was found that they significantly up-regulated genes in the forebrain related to neurogenesis and antioxidant neuroprotection (Zhang et al. 2013a).

Kai-xin-san (KXS), a Chinese herbal decoction contains *Ginseng Radix rhizoma, R. Polygalae radix, R. Acori Tatarinowii,* and *Poria*. KXS has been used in China to treat stress-related psychiatric diseases with the symptoms of depression and forgetfulness. A chemically-standardized water extract of KXS applied to astrocytes significantly stimulated the

Table 5.3 The top Chinese herbs for improving memory, their major constituents, anti-dementia and neuroprotective actions

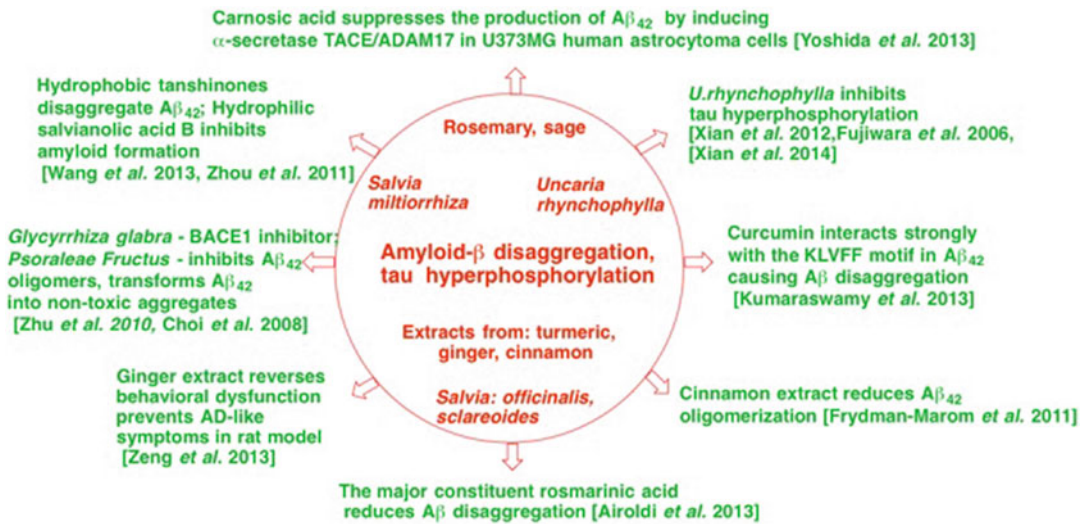
Chinese herbs and constituents	Therapeutic and anti-dementia bioactivities
<i>P. cocos</i> (a medicinal mushroom) triterpenes, pachymic acid, dehydropachymic acid.	Antioxidant; water extract enhanced hippocampal long-term potentiation, improved scopolamine-induced spatial memory impairment in rats (Cheng et al. 2013b; Hatip-Al-Khatib et al. 2004; Smriga et al. 1995)
<i>Radix ginseng</i>	Refer to Table 5.2
<i>Radix polygalae</i> (RP) oligosaccharide multi-esters, sucrose esters, triterpene onjisaponins, xanthone and xanthone C-glycosides	Sedative, antipsychotic, cognitive-improving, neuroprotective, with anti-inflammatory therapeutic effects on the central nervous system. Onjisaponin B was able to induce autophagy and accelerate both the removal of mutant huntingtin and A53T α -synuclein, associated with Huntington's and Parkinson's diseases (Ling et al. 2013; Wu et al. 2013a)
<i>Radix Glycyrrhizae</i> (RG) and the active constituent isoliquiritigenin	RG antioxidant activity related to flavonoids and total phenolics (Li et al. 2013b). Prevented A β (25–35)-induced neuronal apoptotic death by interfering with the increases of intracellular Ca ²⁺ and ROS, and RG potential therapeutic for preventing the progression of AD (Lee et al. 2012)
<i>Radix glycyrrhizae glabra</i>	Administration of 150 and 225 mg/kg improved learning and memory via antioxidant, anti-inflammatory effects in rat model studies. Glycyrrhiza (60–200 μ g/mL) contributed to the suppression of A β oligomer-induced neuronal damage, DNA fragmentation, and caspase-3 activation (Chakravarthi and Avadhani 2013; Kanno et al. 2013)
<i>R. angelica sinensis</i> (RAS); Z-Ligustilide (Lig) (Fig. 5.3) is the major constituent of the lipophilic extract of RAS	Decreased A β content and deposition in SAMP8 mice (Huang et al. 2008; Kuang et al. 2006; Hu et al. 2012b)
<i>Semen ziziphi spinosae</i> Jujuboside A (JuA) a major hypnotic-sedative	JuA has shown notable neuroprotective activities via anti-oxidative and anti-inflammatory effects in dementia animals and has potential utilization for the therapeutic treatment of AD (Liu et al. 2014)
<i>Radix Rhemanniae</i> Catalpol, iridoid glycoside	Catalpol reversed brain damage and memory deficits in mice; antioxidant, anti-inflammatory, neurogenetic, antiapoptotic, neuroprotective activities (Liang et al. 2009)
<i>Rhizoma zingiberis</i> ginger root extract (GRE)	GRE reverses behavioral dysfunction and prevents AD-like symptoms in rat model. Ginger has been shown to possess free radical scavenging, antioxidant inhibition of lipid peroxidation, dementia and multiple other therapeutic applications (Zeng et al. 2013; Haniadka et al. 2013)

expression and secretion of neurotrophic factors, including nerve growth factor (NGF), brain-derived neurotrophic factor (BDNF) and glial cell-derived neurotrophic factor (GDNF), in a dose-dependent manner: the stimulation was both in mRNA and protein expression (Zhu et al. 2013; Man et al. 2012). *Rhizoma Acori Tatarinowii* (grassleaf or sweet-flag rhizome), the rhizome of *Acorus tatarinowii* Schott, is used in TCM as an anti-convulsant; it can prevent convulsions as well as convulsion-related GABAergic neuron damage in the brain (Liao et al. 2005).

From the analysis of 1,232 traditional Chinese medicine formulae for anti-dementia (Kong et al. 2009) it was suggested that the most commonly

used herbal formulation (Fig. 5.5) was *Rhizoma Chuanxiong*, *Radix Salviae Miltiorrhizae*, *Radix Polygalae Tenuifoliae* and *Rhizoma Acori Tatarinowii*. Their major chemical constituents and anti-AD activities are summarized in Table 5.4.

Yukukansan (Yigan San) is a classical TCM formula used for dementia (Iwasaki et al. 2005b) composed of seven herbs, *Angelica acutiloba*, *Atractylodes lancea*, *Bupleurum falcatum*, *Poria cocos*, *Cnidium officinale*, *Uncaria rhynchophylla* and *Glycyrrhiza uralensis*, in a ratio of 3:4:2:4:3:3:1.5. Clinical randomized controlled trials (RCTs) revealed that Yigan San improved behavioral and psychological symptoms of dementia that include aggression,



Scheme 5.4 Anti β -amyloid effects of food spices and herbs

Scheme 5.5 Chinese herbs most commonly used against dementia and AD



agitation, screaming, wandering, hallucinations and delusions. Yigan San reduces cholinesterase inhibitor-resistant visual hallucinations in dementia patients (Iwasaki *et al.* 2005a). Yigan San improved psychiatric symptoms and sleep structure in dementia patients (Shinno *et al.* 2008). The mechanisms of action are related to regulating multiple signal pathways, such as the glutamatergic neurotransmitter system, the serotonin receptor and excitotoxicity (Ho *et al.* 2011).

A key challenge in validating and translating fundamental science of herbal medicines into better anti-dementia outcomes is to evaluate and scrutinize clinical trial outcomes using scientific research methodologies. Some animal and clinical research performed on herbs leading to improved cognitive health providing options for dementia management and prevention is presented in Table 5.5.

5.3 Summary and Future Outlook

The individual-based interventionist approach against dementia and AD for extending healthy life — better diet and regular exercise — is effective, however it needs much greater promotion, acceptance and adoption early on in life. Alkaloids, monoterpenes, diterpenes, triterpenes, flavonoids, and polyphenolic compounds represent the most prevalent classes of herbal constituents with anti-AD bioactivity. It is unclear to what extent many of these bioactive phytochemicals utilized in single or herbal formulae doses can reach the brain in sufficient concentrations, and in a biologically active form, to exert their beneficial neuroprotective effects. The majority of herbs are consumed as aqueous extracts

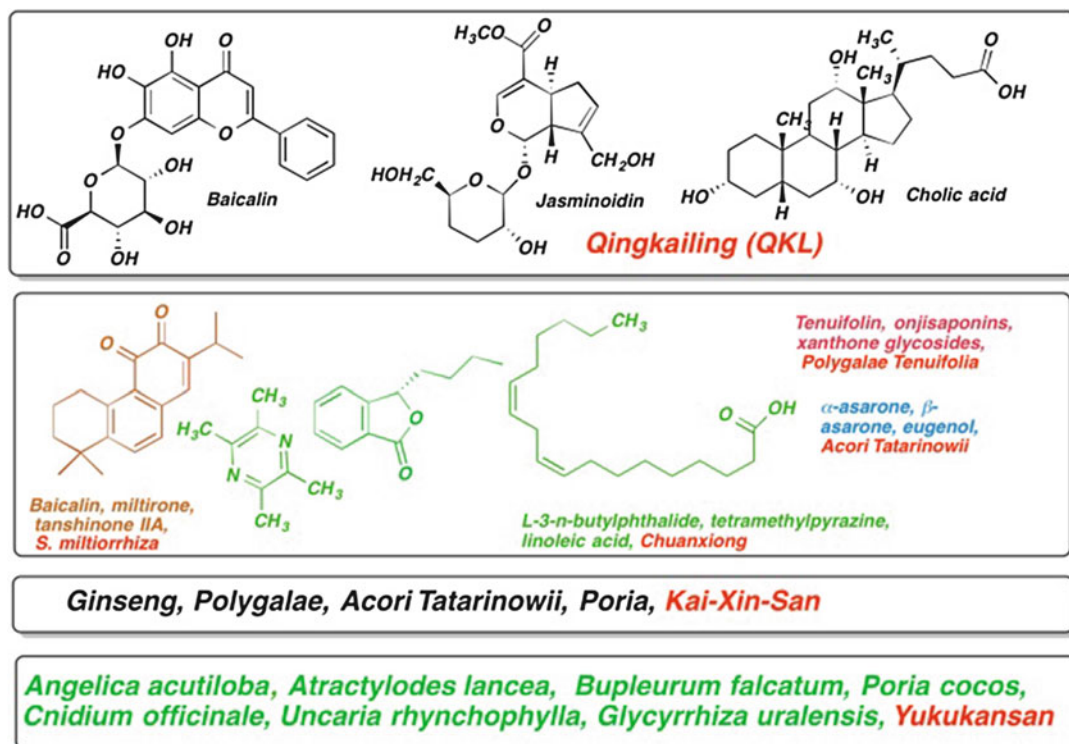
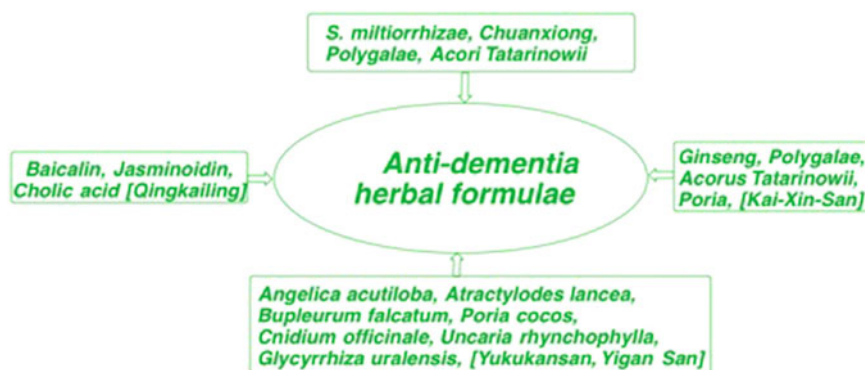


Fig. 5.5 Neuroprotective constituents of Chinese herbal formulae against dementia



Scheme 5.6 Herbal combinations and formulations used for dementia treatment

so their formulation has to provide increased bioavailability and BBB permeability (Hügel and Jackson 2014). An overview of the metabolism and strategies for enhancing polyphenol bioavailability (Lewandowska et al. 2013) include encapsulation of phospholipid-polyphenol complexes; formation of inclusion complexes with cyclodextrins or dendrimers; use of bioactive

analogues; derivatisation (e.g., amidation); use of adjuvants (e.g. piperine) as absorption enhancers; and transdermal delivery systems.

It is imperative that herbs and herbal constituents are consumed regularly and in sufficient quantities in the diet. Indeed, for *in vivo* and clinical studies, producing active compounds and extracts in large quantities is an important

Table 5.4 Neuroprotective effects of the four herb TCM formulae commonly used for dementia treatment

The constituents of a four herb anti-dementia TCM formula	Anti-AD activities
<i>Rhizoma Chuanxiong</i> Tetramethylpyrazine ligustrazine	Improved hippocampal cholinergic system function, antioxidant, enhanced learning and memory in AD mice model (Zhao et al. 2008; Shi et al. 2012)
<i>L</i> -3- <i>n</i> -butylphthalide (86,89) 9- <i>cis</i> , 12- <i>cis</i> -linoleic acid (CLA)	<i>L</i> -3- <i>n</i> -butylphthalide has been shown to reduce β -amyase-induced neuronal apoptosis, improve cognitive function, blood flow in AD animal models CLA as a μ -calpain-specific inhibitor. CLA showed neuroprotective effects against neurotoxins such as H ₂ O ₂ and A β (1–42) in SH-SY5Y cells; inhibited A β oligomerization and fibrillation. CLA decreased the levels of pro-apoptotic proteins (Lee et al. 2013)
<i>Radix Salviae Miltiorrhizae</i> Baicalin, polyphenolic acids, tanshinones	Antioxidants, anti-inflammatory, neuroprotection; inhibition of A β aggregation, oligomerization, and fibril formation (Wang et al. 2013; Zhou et al. 2011; Mei et al. 2009)
<i>Radix Polygalae Tenuifoliae</i> 3,6'-di-O-sinapoyl-sucrose (DISS) tenuifolin, onjisaponins, xanthone glycosides	DISS exerts neuroprotective effects against glutamate toxicity. Reinforces cognitive performance in aged and dysmnesia mice, elevating levels of dopamine, norepinephrine. Onjisaponins indicated cytoprotective activity in PC12 cells, exposed to serum deficiency or glutamate; improved memory in rats by enhancing cholinergic function, inhibiting A β secretion (Hu et al. 2009, 2012a; Lin et al. 2012)
<i>Rhizoma Acori Tatarinowii</i> Eugenol, α -asarone, β -asarone	Eugenol derived from <i>Rhizoma Acori Tatarinowii</i> increased BDNF mRNA expression level in hippocampus of mice. Modified Wen-Dan-Tang decoction containing <i>Acori Tatarinowii</i> attenuated the neurotoxicity of A β (25–35) and rescued neurons via suppressing apoptotic process (Liu et al. 2009)

Table 5.5 Clinical trials with herbs to counteract dementia and AD

Natural product	Animal studies; bioactivity mechanisms	Clinical trials
EGCG		300 mg/day of EGCG for 12 weeks had no adverse effect on liver function; did not enhance energy-restricted diet-induced adiposity reductions; did not improve weight-loss-induced changes in cardio-metabolic risk factors in obese Caucasian women (Mielgo-Ayuso et al. 2014)
Huperzine A (alkaloid shown in Fig. 5.3)	APPswe/PS1dE9 transgenic mice reduced A β fibrils, oligomers; inhibition of BACE1, regulating APP metabolism (Smriga et al. 1995). EGCG addition to huperzine A, significantly enhanced and prolonged the AChEI effects of huperzine A (Wang et al. 2012; Xiao et al. 2008)	Commonly used in China. USA clinical data (Ha et al. 2011) suggests 0.4 mg doses are required. Further non-Chinese clinical trials are necessary before the implementation of huperzine A for dementia and AD treatment (Yue et al. 2012) Systematic review and meta-analysis of 20 RCTs of Huperzine A for AD. Huperzine A appears to have beneficial effects on improvement of cognitive function, daily living activity, and global clinical assessment in participants with AD. The quality of some of the trials was an issue (Yang et al. 2013)

(continued)

Table 5.5 (continued)

Natural product	Animal studies; bioactivity mechanisms	Clinical trials
Curcumin	Curcumin <i>in vitro</i> inhibits: A β aggregation, A β -induced inflammation; the activity of β -secretase; AChE. In <i>in vivo</i> studies: oral curcumin inhibition of A β deposition, oligomerization, tau phosphorylation in AD animal models. Improvement in behavioral impairment in animal models (Hamaguchi et al. 2010)	Safe to use at dosage of 8 g/day for 3 months RCT study on 34 AD patients found no cognitive improvement, increase in anti-oxidant activity and vitamin E levels (Baum et al. 2008) Two CTs performed in China and USA have reported no significant differences in changes in cognitive function between placebo and curcumin groups (Gupta et al. 2013)
Korean red ginseng (KRG)		Used for adjuvant treatment for cognitive impairment in AD patients. High-dose KRG (9 g/day, $n = 15$) patients showed significant improvement on the AD Assessment and Clinical Dementia Rating Scale after 12 weeks of KRG therapy (Heo et al. 2008)
Rosemary (<i>Rosmarinus officinalis</i> L.; carnosic and rosmarinic acids)		Cognition improving effects of dried rosemary leaf powder on 28 adults (mean age 75 years). Only the lowest dose (750 mg) of rosemary had a statistically significant beneficial effect compared with placebo. Requires further work on effects of low doses over the longer term (Pengelly et al. 2012)

challenge for the utilization of natural products as therapeutic agents. Generally speaking, herbal products offer a wide range of brain-targets, nutritional benefits, safe dosage, long-term applications and efficacious treatment of AD pathology. The focus on engagement of sustainable optimal biochemical performance through diet and factors influencing it, including lifestyle choices, are key to a better mental health.

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Mario Caruana and Neville Vassallo

Abstract

Parkinson's disease (PD) is a common motor neurodegenerative disorder with multifactorial etiology that is an increasing burden on our aging society. PD is characterized by nigrostriatal degeneration which might involve oxidative stress, α -synuclein (α S) aggregation, dysregulation of redox metal homeostasis and neurotoxicity. Although the exact cause remains unknown, both genetic and environmental factors have been implicated. Among the various environmental factors tea consumption has attracted increasing interest, as besides being one of the most consumed beverages in the world, tea contains specific polyphenols which can play an important role in delaying the onset or halting the progression of PD. Green and black teas are rich sources of polyphenols, the most abundant being epigallocatechin-3-gallate (EGCG) and theaflavins. There is now consistent mechanistic data on the neuroprotective and neuroregenerative effects of tea polyphenols, indicating that they do not just possess anti-oxidant or anti-chelating properties but may directly interfere with aggregation of the α S protein and modulate intracellular signalling pathways, both *in vitro* and in animal models. EGCG in green tea has been by far the most studied compound and therefore future investigations should address more the effects of other polyphenols, especially theaflavins in black tea. Nevertheless, despite significant data on their potential neuroprotective effects, clinical studies are still very limited and to date only EGCG has

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reached phase II trials. This review collates the current knowledge of tea polyphenols and puts into perspective their potential to be considered as nutraceuticals that target various pathologies in PD.

Keywords

Parkinson's disease • Alpha-synuclein • Tea • Epigallocatechin-gallate • Theaflavins • Neuroprotection

6.1 Introduction

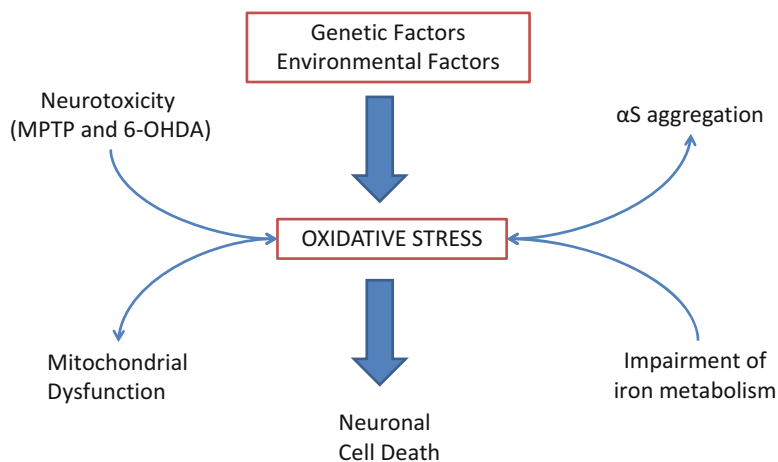
Parkinson's disease (PD) is the most common movement disorder and after Alzheimer's Disease (AD), the second most frequent progressive neurodegenerative disease (Toda 2007). The prevalence of PD worldwide ranges from 0.5 to 4 % among people aged 65 years or older. This figure is expected to rise significantly with the accelerated aging of human society (de Lau and Breteler 2006). In fact, it was predicted that by 2030, the number of PD sufferers will reach 9.3 million (Dorsey et al. 2007). PD is a debilitating disorder with varying patterns of degeneration in the dopaminergic and nondopaminergic neuronal systems (Braak et al. 2004). The major symptoms of PD include muscular rigidity, uncontrollable resting tremor, bradykinesia or akinesia, and impaired postural reflexes (Jankovic 2008). PD is distinguished from other forms of parkinsonism by the presence of Lewy bodies (LBs) and Lewy neurites (LNs), which are juxtannuclear and neuritic ubiquitinated protein aggregates composed predominantly of the presynaptic protein α -synuclein (α S) (Shults 2006). The etiology of PD in most patients remains unknown. It is assumed that both genetic and environmental factors with complex interactions are responsible for the development and progression of the disease (Fig. 6.1; Logroscino 2005).

The pathogenesis and relative selectivity of death of dopaminergic neurons in the substantia nigra (SN) *pars compacta* remains to be clarified (Kazantsev and Kolchinsky 2008). Various pathogenic mechanisms have been proposed through which dopamine-releasing neurons may be damaged in PD. These include deficiency in mitochondrial respiratory chain function (Fukae

et al. 2007), apoptosis (Hartmann and Hirsch 2001), transition metal accumulation (Barnham and Bush 2008), oxidative stress (Friedman and Galazka-Friedman 2001), deficiency in the xenobiotic mechanism (Ramsden et al. 2001), inflammation (McGeer et al. 2001), and abnormal protein handling, aggregation and misfolding (Skovronsky et al. 2006). The most favoured ones are oxidative stress due to an increased production of reactive oxygen species (ROS), and cell toxic effects of α S protein aggregation and deposition, both finally leading to neuronal cell death by apoptosis (Soto 2003). Regardless of the cause of neuronal death, the plasticity of the *pars compacta* is very robust; symptoms do not appear until 50–80 % of SN dopaminergic neurons have died. Therefore, it is not surprising that diagnosis in the early course of disease is more than rare (Jankovic 2008).

The results of studies on twins suggested that genetic factors are important in early-onset PD cases while environmental factors play a predominant etiologic role in late-onset PD patients, thus implying the importance of non-genetic factors (Tanner et al. 1999; Wirdefeldt et al. 2004). Environmental factors such as coffee drinking and smoking have been demonstrated to lower the risk of PD (Hernan et al. 2002; Costa et al. 2010; Wirdefeldt et al. 2011). The effects of tea consumption on PD risk are currently the subject of considerable scientific debate as tea components, such as polyphenols, caffeine, and theanine, have been demonstrated to be neuroprotective in PD (Tan et al. 2008; Quintana et al. 2009). The benefits of tea drinking are of relevance to PD as tea is one of the main contributors of dietary polyphenols in Western countries due to its regular consumption (Erdman et al. 2007).

Fig. 6.1 Factors that contribute to oxidative stress and ultimately neuronal cell death in PD (MPTP, N-methyl-4-phenyl-1,2,3,6-tetrahydropyridine, 6-OHDA, 6-hydroxydopamine, α S alpha-synuclein)



Thus, any evidence of the neuroprotective effects of polyphenols on PD could have a significant impact on public health. The purpose of this chapter is to provide a concise review of the most recent scientific evidence from epidemiological, experimental and clinical studies on the crucial role green and black tea polyphenols may have in the prevention and treatment of PD.

6.2 Polyphenolic Components of Tea

Tea has been consumed as a beverage for well over 2,000 years, and its worldwide consumption is perhaps second only to water. The term 'tea' refers to the dried leaves of the plant *Camellia sinensis*, an evergreen shrub of the *Theaceae* family. The three principal varieties of tea are generally categorized by the process used in their manufacture: (i) fermented (oxidized) black tea (78 %, mainly consumed in Western Europe, the United States of America, Australia, and some Asian countries); (ii) unfermented (non-oxidized) green tea (20 %, mainly consumed in China, Japan, and India); and (iii) semi-fermented (semi-oxidized) oolong tea (2 %, consumed in south-eastern China and Taiwan) (Fig. 6.2; Graham 1992; Balentine et al. 1997). Another commonly used tea is the so-called 'herbal tea'. Herbal tea is made from any of a number of a variety of plants and herbs, and therefore, cannot technically be considered a true type of tea.

While tea consists of over 2,000 different chemical substances such as methylxanthine, caffeine, lipids, amino acids, mineral substances and volatile compounds, polyphenols are the most abundant (Wheeler and Wheeler 2004). Polyphenols are a diverse class of plant secondary metabolites and more than 8,000 polyphenolic compounds have currently been identified (Porat et al. 2006; Stevenson and Hurst 2007). Polyphenols are classified into different groups depending on the number of phenol rings and the chemical groups attached to the rings. They are characterised by a polyphenol structure, which generally consists of two aromatic rings (2-phenyl-1,4-benzopyrone) each containing at least one hydroxyl group, which are connected via a three-carbon bridge and become part of a six-member heterocyclic ring (Fig. 6.3; Beecher 2003; Porat et al. 2004; D'Archivio et al. 2007). Polyphenols can be divided into two main groups, the flavonoids and the non-flavonoids, with the flavonoids making up the largest and most important single group of polyphenols present in tea (Vassallo 2008). Black and green teas both contain similar amount of flavonoids, however they differ in their chemical structure. Green teas contain more of the simple flavonoids called flavanols (known also as catechins or flavan-3-ols). The principal four flavanols found in green tea are epicatechin (EC), epicatechin 3-gallate (ECG), epigallocatechin (EGC) and epigallocatechin-3-gallate (EGCG), where the latter is the most abundant (Rietveld

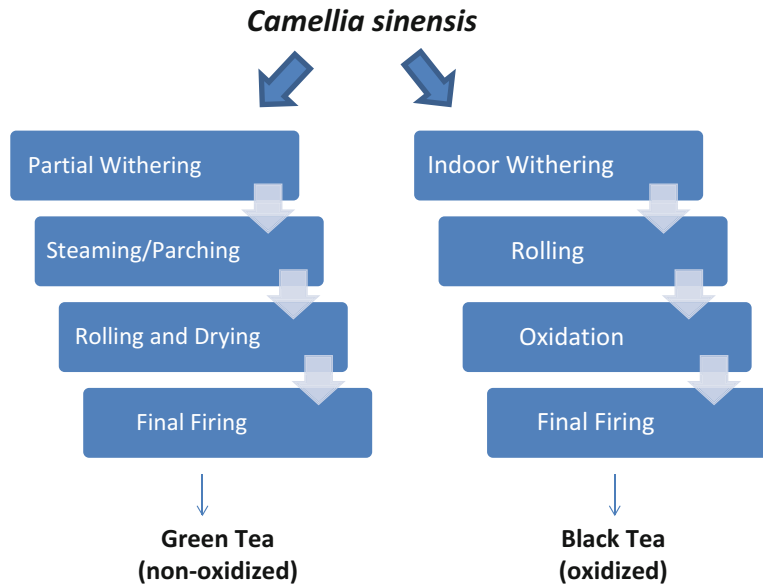


Fig. 6.2 Green and black tea processing. Tea is produced when freshly picked leaves are steamed, rolled and dried. Tea leaves contain polyphenol oxidase enzymes in separate layers of the leaf. When tea leaves are rolled or broken during industry manufacture, polyphenols known as flavanols (catechins) come in contact with polyphenol oxidase, resulting in their oxidation and the formation of flavanol dimers and polymers known as theaflavins and thearubigins. Tea leaves destined to become black tea

are rolled and allowed to ferment (oxidize), resulting in relatively high concentrations of theaflavins and thearubigins and relatively low concentrations of flavanols. Green tea is withered and then steamed to inactivate polyphenol oxidase. Consequently, green tea contains relatively high concentrations of flavanols and low concentrations of theaflavins and thearubigins (Graham 1992; Balentine et al. 1997)

and Wiseman 2003). The oxidation that the leaves undergo to make black tea converts these simple flavonoids to the more complex varieties called theaflavins and thearubigins (Khokhar and Magnusdottir 2002). The chemical composition of tea (Table 6.1) varies with the variety of plant and age of the leaf, the conditions under which it is grown, climate, season, and local agricultural practices (Aherne and O'Brien 2002).

Research interest in the benefits of tea drinking stems primarily from the presence of polyphenols which are believed to be the major component that provide health benefits (McKay and Blumberg 2002; Erdman et al. 2007). One cup of tea (2 g of tea leaves infused in hot water for 1–3 min) will provide 0.15–0.2 g of flavonoids. As little as 2–3 cups/day of tea will therefore supply a significant contribution to the total flavonoid intake in most individuals, which is estimated to average 1 g per day (Frei and Higdon 2003). In fact, it was estimated that black

tea contributes 60–84 % of dietary flavonoids in Western populations (Hertog et al. 1993; Chun et al. 2007) and it has been reported that flavonoid intakes in tea consumers are twenty times greater than in non-tea consumers (Song and Chun 2008). This intake is higher than all other known dietary anti-oxidants, estimated to be around ten times higher than the daily intake of Vitamin C, and 100 times higher than that of Vitamin E and carotenoids (Scalbert and Williamson 2000). As the total polyphenol content of green and black teas is similar, it can be assumed that the impact on plasma levels post-consumption remains fairly the same (Rietveld and Wiseman 2003).

6.3 Epidemiological Studies on Tea Consumption and PD

In the 1980s, PD prevalence was found to be low in Asian countries when compared to Europe and North America, which had significantly higher

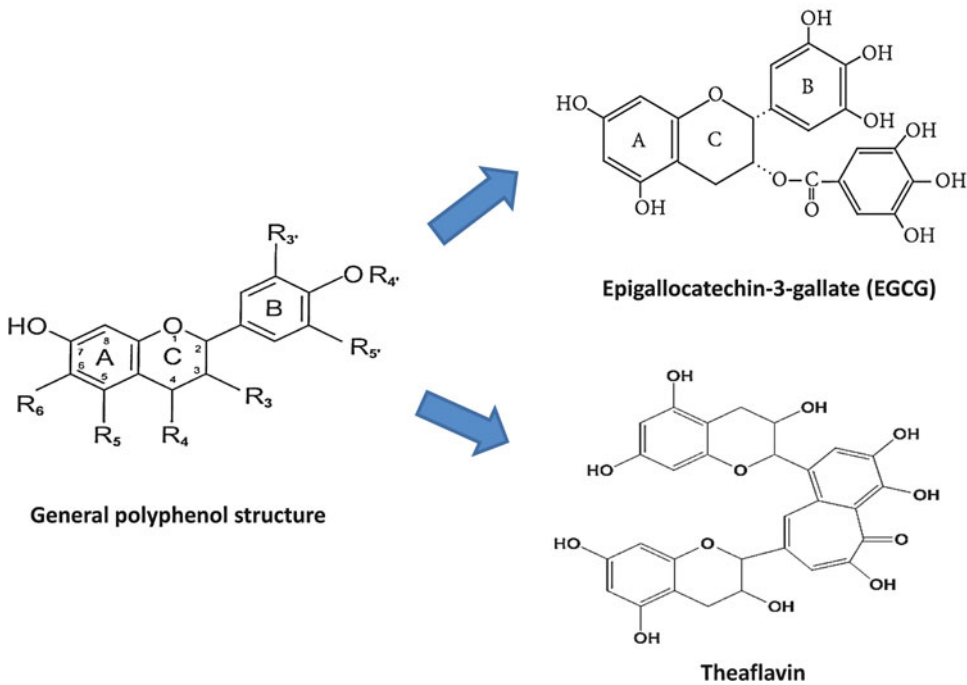


Fig. 6.3 General structure and numbering pattern for polyphenols. This figure shows the general structure and numbering pattern for common polyphenols. Every flavonoid subclass has its own unique linkages, unsaturation positions and functional groups. For most food flavonoids, R_{4'} = H, R₅ = OH and R₆ = H. Individual flavonoids within each subclass are characterised by unique functional groups at R₃, R_{3'}, and R_{5'}. Chemical

structures of epigallocatechin-3-gallate (EGCG) and theaflavin in green and black tea are shown. EGCG contains three heterocyclic rings (A, B, C) and the free radical scavenging property of EGCG is attributed to the presence of a trihydroxyl group on the B-ring and the gallate moiety at the 3' position in the C-ring. Theaflavin is the polymeric form of EGCG

rates (Li et al. 1985; Zhang and Román 1993). Apart from genetic factors, dietary habits like green tea consumption, which is more consumed by the Chinese population when compared to Caucasian, could explain this attribute (Pan et al. 2003; Gao et al. 2012). Due to this possible link, in recent years, there were more studies devoted to exploring the effects of tea consumption on PD risk. Three case–control studies (in the US, Hong Kong and Singapore) and a cohort study of male health professionals in the US have all reported an inverse association between tea drinking and PD risk (Chan et al. 1998; Checkoway et al. 2002; Tan et al. 2003). One study found such an effect for men but not for women (Ascherio et al. 2001). On the other hand, a hospital based case-control study in France reported tea consumption to be a paradoxically risk factor for PD (Preux et al. 2000).

The authors attributing the protective effect of tea suggested caffeine as the main contributor. Similarly, a biologic effect of caffeine was suggested for a positive association of tea drinking and PD in a prospective study of over 29,000 Finnish adults for 13 years (Hu et al. 2007). In another prospective study of 63,000 Chinese adults, black tea showed an inverse association with PD risk, although this time the link was not confounded by total caffeine intake or tobacco smoking (Tan et al. 2008). Surprisingly in this study, green tea consumption after adjustment for cigarette smoking and total caffeine consumption was unrelated to PD risk. The authors speculated that the protective effect of black tea may be mediated via an estrogen-related pathway. This was based on what they had reported earlier that among the women in their study cohort, levels of circulating estrogens were highest in

Table 6.1 Average values for the different constituents present in green and black tea

	Black tea	Green tea
Catechins	3–10	30–42
Theaflavins	3–6	Negligible
Thearubigins	12–18	Negligible
Flavonols	6–8	5–10
Theogallin	Negligible	2–3
Phenolic acids (caffeic acid)	10–12	1–2
Theanine	Negligible	4–6
Other amino acids	13–15	4–6
Methylxanthines	8–11	7–9
Carbohydrates	15	10–15
Protein	1	Negligible
Minerals	10	6–8
Volatiles	<0.1	0.02

The values will differ dependent on the variety of leaf, growing environment, manufacturing, particle size of ground tea leaves and infusion preparation (Graham 1992; Harbowy and Ballentine 1997; Wang and Hellwell 2001; Astill et al. 2001)

Values reflect % weight of extract solids

regular black tea drinkers, intermediate in non-tea drinkers, and lowest in regular green tea drinkers; these differences were dose-dependent and significant (Wu et al. 2005).

It was suggested that the average 1.2 l of green tea consumed daily by many people in Asia offers sufficient anti-oxidants of the polyphenolic EGCG, and in turn reduces or cures diseases with an inflammatory component, together with improving neurologic and psychological health (Sumpio et al. 2006). In terms of a dose-response relationship, only few studies have stratified their results according to the number of cups of tea consumed daily. One study showed a dose-dependent protective effect of PD in tea consumers with an odds ratio (OR) of 0.48 for daily consumption of a cup of tea versus OR of 0.27 for daily consumption of two or more cups of tea (Fall et al. 1999), whilst another study showed a similar effect in coffee and tea consumers (Tan et al. 2003). In the latter study it was concluded that one unit of coffee or tea (three cups per day for 10 years) would lead to a 22 and

28 % risk reduction of PD, respectively. On the contrary, another study could not demonstrate such dose-dependent protective effect in their hospital based case-control study (Paganini-Hill 2001). More evidence was provided by an Israeli study of 278 PD patients whose motor symptoms appeared to be delayed by 7.7 years ($p < 0.01$) when they drank more than three cups of tea per day (Kandinov et al. 2009). Two recent systematic reviews showed that tea drinking can lower the risk of PD, but no apparent dose-response relationship was found as would be expected (Quintana et al. 2009; Li et al. 2012). The latter may arise from the fact that black tea and green tea differ markedly in the nature of their polyphenols and only few studies reported stratified results according to the types of tea. It is also important to note that the contents of the bioactive compounds in tea may fluctuate because of differences in producing areas, materials, and manufacturing (Crozier et al. 2009). While the addition of milk to tea does not seem to interfere with flavonoid absorption or activity (Hollman et al. 2001), it is not obvious if other factors do – such as the frequency and timing of tea intake in relation to meals, the addition of sucrose or lemon, and variations in gut microflora. Therefore, although a positive association has turned up repeatedly in epidemiological studies between tea and PD, a clear biologic basis for this phenomenon has yet to be identified.

When reviewing the literature, the strongest and most consistent environmental associations were those between cigarette smoking, coffee/tea drinking, and a reduced risk of PD as noted in several US and European populations (Tanner et al. 2002; Hernan et al. 2002; Ritz et al. 2007; Hancock et al. 2007; Saaksjarvi et al. 2008). However, the strength of the evidence for the described inverse associations seems to be weaker for tea than for smoking or coffee drinking. The precise reasons for this are not known, although tea has not been investigated in relation to PD risk as extensively or explicitly as coffee has – perhaps because consumption of coffee is far more prevalent in North America and Europe, where most research on PD has been undertaken.

Moreover, the selection of patients and the type of control groups, for example the inclusion of patients with preclinical stage of PD, may result in conflicting results (Schrag et al. 2002). It is unclear from epidemiological studies whether the active ingredient/s mediating this neuroprotective effect in tea is actually caffeine or the polyphenols in tea. In most cases, this work can only be conducted in experimental designs not least because practical and ethical constraints limit such research in humans.

6.4 Neuroprotective Actions of Tea Polyphenols in PD

6.4.1 Tea Polyphenols and *In Vitro* Studies

Numerous *in vitro* studies have clearly demonstrated that specific tea polyphenols might contribute to prevent PD pathology and act towards neuroprotective capacities (Levites et al. 2002a; Bastianetto 2002; Bastianetto and Quirion 2004). Cell culture studies have demonstrated that flavanols reduced damage produced by hydrogen peroxide (H₂O₂), 4-hydroxynonenal, rotenone, and 6-hydroxydopamine (6-OHDA) in primary rat mesencephalic cultures, as shown by increases in cellular viability and [³H] dopamine uptake (Mercer et al. 2005; Vauzour et al. 2008). Other *in vitro* studies demonstrated that EGCG is able to rescue and reduce viability of neuroblastoma SH-SY5Y cells when administered up to 3 days after long-term serum starvation, a model of apoptotic damage (Reznichenko et al. 2005). As reviewed elsewhere, polyphenolic compounds provide neuroprotective effects through a variety of biological actions such as anti-oxidant, anti-chelating, anti-aggregating, anti-inflammatory, anti-carcinogenic, anti-viral, anti-microbial and anti-clotting activities (Scalbert et al. 2005; Ramassamy 2006; Rahman et al. 2007; Moon and Shibamoto 2009; Obrenovich et al. 2010; Albani et al. 2010; Choi et al. 2012). With regards to specific tea polyphenols, the most important plausible mechanisms cited that may be exhibiting neuroprotective effects in PD are:

(i) anti-oxidant and anti-chelating activities; (ii) inhibition of α S aggregation; and (iii) modulation of cell signalling pathways (Pan et al. 2003; Higdon and Frei 2003; Amit et al. 2008), which will be reviewed in the next sections.

6.4.1.1 Anti-oxidant and Iron-Chelating Activity

Substantial evidence of the potent anti-oxidant effects of the main tea polyphenols (flavanols and theaflavins) comes from *in vitro* studies, where they were shown to: (i) directly scavenge reactive oxygen (ROS) and nitrogen oxygen (NOS) species; (ii) inhibit 'pro-oxidant' enzymes, such as nitric oxide synthase, xanthine oxidase, cyclooxygenases and lipoxygenases; (iii) inhibit redox-sensitive transcription factors such as nuclear factor- κ B and activator protein-1; (iv) induce phase II and anti-oxidant enzymes such as glutathione S-transferases and superoxide dismutases; and (v) bind and chelate excess of divalent metals such as iron (Fe²⁺) and copper (Haenen et al. 1997; Nakagawa and Yokozawa 2002; Higdon and Frei 2003; Stevenson and Hurst 2007; Aron and Kennedy 2008; Mandel et al. 2008; Perron and Brumaghim 2009; López-Lázaro 2009). The oxygen radical absorbance capacity (ORAC) assay has demonstrated that both green and black tea have much higher capacity against free radicals than vegetables, for instance garlic and spinach (Cao et al. 1996).

The capacity of polyphenols to act as anti-oxidants is dependent upon their molecular structure, the position of hydroxyl groups, and other substitutions in their chemical structure (Tsao 2010). Although the oxidation process modifies the type of flavonoids present, the total level and their overall anti-oxidant activity, is similar in both teas (Leung et al. 2001; Luczaj and Skrzydlewska 2005). EGCG has an important anti-oxidant and iron-chelating function and this could be attributed to the 3',4'-dihydroxyl group in the B-ring, as well as the gallate group which may neutralise Fe²⁺ to form redox-inactive iron, thereby protecting cells against oxidative damage (Fig. 6.3; Kumamoto et al. 2001). In addition, the feature of tea polyphenols as potent chelators

of transitional metals, such as iron and copper, is owed to the OH at position 3' of the C-ring, the OH at positions 3' and 4' of the B-ring, or the three OH groups present in the gallol moiety of some polyphenols, such as EGCG and ECG (Nanjo et al. 1996). In a recent study examining the differential potency of a series of polyphenols to prevent DNA damage caused by Fe^{2+} and H_2O_2 , it was found that among the 12 phenolic compounds tested, EGCG was the most potent, inhibiting over 90 % of the iron-mediated DNA break (Perron et al. 2008). By correlating the pK_a and IC_{50} values of phenolic compounds for inhibition of $\text{Fe}^{2+}/\text{H}_2\text{O}_2$ -induced neurotoxicity, it was suggested that the binding of the polyphenols to iron was essential for their anti-oxidant activity (Perron et al. 2010). In one experiment performed on rat brain tissue, it was shown that lipid peroxidation was enhanced by iron ascorbate but inhibited in brain mitochondria by both black and green tea extracts (Jeong et al. 2004).

The anti-oxidant and metal-complexing properties of tea polyphenols may be of significance in the treatment of PD, since oxidative stress and accumulation of Fe^{2+} at brain areas associated with neurodegeneration have been clearly demonstrated (Zecca et al. 2004). The anti-oxidant activity could protect the dopaminergic system against free radicals, anion superoxide, lipid free radicals and hydroxyl radicals, together with neurotoxic apoptosis induced by hydroxydopamine in the cell (Weinreb et al. 2004). Also, at the central nervous system (CNS) level it may inhibit the peroxidation and lipid accumulation of Fe^{2+} compounds, and this could be the main mechanism for neuroprotection (Soto-Otero et al. 2000; Pan et al. 2004; Levites et al. 2002b). Nevertheless, it is currently more accepted that neuroprotective effects of tea polyphenols are only partly attributed to the free radical scavenging or metal chelating properties and that other properties such as targeting of specific signalling pathways and interaction with specific proteins, including αS , contribute to neuroprotection (Kaur et al. 2003; Masuda et al. 2006; Ramassamy 2006; Vafeiadou et al. 2007; Weinreb et al. 2010).

6.4.1.2 Inhibitory Activity on α -Synuclein Aggregation

Ample evidence suggests that disturbance of neuronal membranes by the soluble oligomers of the protein αS is a likely first step in the pathophysiological cascades of PD, where partial aggregated and oligomerized intracellular αS was shown to be cytotoxic and synaptotoxic (Periquet et al. 2007; Selkoe 2008). A considerable amount of scientific data shows that a possible neuroprotective characteristic of polyphenolic compounds is exerted through anti-aggregating properties (Caruana and Vassallo 2011). In relation to PD, such properties were initially tested *in vitro* on the inhibition of the assembly of αS into filaments/fibrils (Conway et al. 2001). For example, tea polyphenols such as EGCG and black tea extract, inhibited wild-type (WT) αS filament assembly and were also found to disaggregate preformed fibrils (Zhu et al. 2004; Porat et al. 2006; Masuda et al. 2006; Ono and Yamada 2006; Meng et al. 2010; Grelle et al. 2011). Recently, EGCG efficiently inhibited fibril formation of αS (Ehrnhoefer et al. 2008; Bae et al. 2010) and also transformed large αS fibrils into smaller non-toxic, amorphous protein aggregates (Hudson et al. 2009; Bieschke et al. 2010). Biophysically, EGCG was postulated to directly bind to unfolded polypeptide chains via hydrogen bonds and hydrophobic peptide backbone interactions, and inhibit beta sheet formation which is the early event in the amyloid formation cascade (Wang et al. 2010). Interestingly, this effect was evident only with flavanols carrying a gallate moiety with a high affinity for metals, such as ECG and EGCG. In this regard, another study showed that only gallate forms of flavanols were able to protect hippocampal cells against amyloid-induced toxicity (Bastianetto et al. 2006). Thus, it is possible that the anti-fibrillogenesis action of polyphenols would also result from an iron-complexing radical scavenging-mediated action.

Since it is hypothesised that small αS oligomers, rather than fibrils, may be the primary toxic species, it was also shown that polyphenolic compounds inhibit and destabilise early stage aggregates (Caruana et al. 2011). Interestingly *in vitro* studies have shown that

the anti-oxidant activities of such polyphenols are not likely to be directly involved in the inhibition progress (Zhu et al. 2004; Johnston and Brotchie 2004; Caruana et al. 2011). Specific structural features, rather than broad biochemical characteristics, determine the anti-aggregation effects of polyphenols. In fact, flavonoids with three vicinal hydroxyl groups exhibited enhanced inhibitory effects on α S aggregation (Meng et al. 2009; Berhanu and Masunov 2010; Caruana et al. 2011). Essentially, a polyphenolic inhibitor, with its polyaromatic nature is able to use aromatic recognition elements to bind the monomer/oligomer, whilst utilizing the vicinyl hydroxyl groups to electrostatically block the progress of the self-assembly process (Gazit 2002; Porat et al. 2006).

The lipophilicity of biologically active compounds is usually one of their most important pharmacological features, and interactions with membranes play an essential role in their biological activity (Hendrich 2006). It is well established that small α S oligomers can interact with and perturb membranes, thereby leading to cell death (Lashuel et al. 2002; Quist et al. 2005; Winner et al. 2011). Similarly, polyphenolic compounds, including tea polyphenols, interact with and alter lipid membranes (Blazovics et al. 2000; Oku et al. 2003; Chen et al. 2011; Duchnowicz et al. 2012; Sharma et al. 2012). Indeed, it was revealed that EGCG inhibits amyloid formation less efficiently at phospholipid interfaces than in bulk solution (Engel et al. 2012). Hence, it is relevant to know how polyphenolic compounds directly effect lipid membranes and how efficiently they can inhibit α S aggregation specifically at the phospholipid membrane interface. Furthermore, it has been established that tea flavanols can adsorb to membranes through associations with the polar headgroups of phospholipids and could protect the integrity of lipid bilayer from disrupting agents (Verstraeten et al. 2003; Sirk et al. 2008, 2011). Such polyphenol-lipid interactions may provide a level of protection for the bilayer from the α S oligomers (or/and monomer aggregation at the membrane surface), contributing to preserve the structure and function of biological membranes. Not many studies have to date exam-

ined the interaction of polyphenols with neuronal membranes and their protective effect on α S-induced membrane dysfunction. For example, black tea extract (80 % theaflavin) was found to strongly protect against membrane perturbation induced by aggregated WT and mutant α S (Caruana et al. 2012). Moreover, black tea extract inhibited permeation of mitochondrial membranes by α S oligomers (Camilleri et al. 2013). The importance of mitochondria in the pathogenesis of PD and the cytotoxicity of oligomeric α S is extensively reviewed elsewhere, and tea polyphenols have been capable of protecting such insults (Büeler 2009; Camilleri and Vassallo 2014; Caruana and Vassallo 2014). Therefore, it is important that the intracellular effects of polyphenols at the membrane level are known, enhancing our understanding of the pharmacological and therapeutic activities of such bioactive compounds.

6.4.1.3 Modulation of Intracellular Signaling Pathways

While there has been a historical spotlight on the anti-oxidant properties of tea polyphenols, there is a general consensus that such flavonoids and their corresponding *in vivo* metabolites may also exert modulatory actions intracellularly through direct action on various signalling pathways in a concentration-dependent manner (Mandel et al. 2004; Williams et al. 2004; Ramassamy 2006; Campos-Esparza and Torres-Ramos 2010). Various inhibitory or stimulatory actions of tea polyphenols on these pathways have been studied, including phosphoinositide 3-kinase (PI3K), protein kinase B (Akt/PKB), protein kinase C (PKC), and mitogen-activated protein kinase (MAPK) (Levites et al. 2002b; Vauzour et al. 2007). The effects of tea polyphenols on signaling pathways in relation to PD will be reviewed briefly.

Most studies related to modulation of cell signalling pathways have been carried out on flavanols found in green tea, but not theaflavins. Table 6.2 summarizes cell signalling pathways targeted by different tea polyphenols. Protein kinase C (PKC) is the target of many flavonoids for providing survival signalling. For example, EGCG at a dose of 2 mg/kg body weight markedly in-

Table 6.2 Cell signalling pathways targeted by different tea polyphenols in neuronal cell lines

Tea polyphenol	Activation (+) Deactivation (-)	Signalling pathways	Neuroprotection	References
EGCG; ECG	-	MAPK	Prevent apoptosis, oxidative stress and endothelial barrier dysfunction, improve mitochondrial functions	Huang et al. (2007), Hwang and Yen (2009), Yang et al. (2010)
EGCG; ECG	-	JNK	Prevent oxidative stress and apoptosis	Choi et al. (2005), Huang et al. (2007)
EGCG; ECG; EC	+	ERK1/2	Prevent oxidative stress and apoptosis	Schroeter et al. (2007)
EGCG, EC	+	PKB	Anti-oxidant defence, prevent oxidative stress-induced apoptosis, stimulate eNOS activity, regulate mitochondrial function	Schroeter et al. (2007), Vauzour et al. (2007), Na et al. (2008), Hwang and Yen (2009), Yang et al. (2010)
EGCG	+	PKC	Neuroprotective	Levites et al. (2002b)
EGCG	+	PI3K	Prevents oxidative stress-induced apoptosis, protects oxidative damage, stimulate eNOS activity	Levites et al. (2002b), Yang et al. (2010), Xi et al. (2012)

The activation of ERK, Akt/PKB, PI3K, and PKC is important to improve cell survival, and the down-regulation of p38 and JNK in preventing apoptosis. The activation of signalling pathways is shown as (+) while down-regulation of signalling pathways by tea polyphenols is shown as (-)

MAPK p38 mitogen-activated protein kinase, *PKB* protein kinase B, *ERK* extracellular signal-regulated protein kinase, *JNK* c-Jun N-terminal kinase, *PI3K* phosphatidylinositol-3 kinase, *PKC* protein kinase C, *eNOS* endothelial nitric oxide synthase, *EC* (-)epicatechin, *ECG* (-) epicatechingallate, *EGCG* (-)epigallocatechingallate

creased PKC in the membrane and cytosolic fractions of mice hippocampus. EGCG restored the reduced PKC and extracellular signal-regulated kinase (ERK1/2) activities caused by 6-OHDA toxicity, and protected against neuronal apoptosis (Levites et al. 2002a). EGCG is also involved in rapid PKC-mediated degradation of the Bcl-2-associated death promoter (Bad) by the ubiquitin proteasome system, thus neutralizing their pro-apoptotic function (Calixto et al. 2004).

Tea polyphenols have been shown to interact with ERK, c-Jun N-terminal kinase (JNK) and p38 pathways of the mitogen-activated protein kinases (MAPKs). EC was shown to modulate protein kinase signalling pathways, depending on the concentration of the compound administered (Schroeter et al. 2007). EC stimulated ERK1/2 and phosphoinositide-3-kinase (PI3K)-dependent cAMP response element-binding protein (CREB) phosphorylation at lower concentrations of 100–300 nM but this effect was no longer apparent

at the higher concentration of 30 μ M. These dose-dependent effects may be important to explain the anti- versus pro-oxidant actions of the tea polyphenols. EC also stimulated ERK and Akt phosphorylation. A 15-min exposure of EC increased the mRNA levels of the glutamate receptor subunit (GluR2) by 60 %, and resulting in increased GluR2 protein. This suggests that EC has the potential to increase CREB-regulated gene expression and increase GluR2 levels and thus modulate neurotransmission, plasticity, and synaptogenesis (Schroeter et al. 2007). EGCG inhibited H₂O₂-induced phosphorylation of JNK and p38 MAPK pathway after a 60-min exposure. EGCG also inhibited H₂O₂-induced caspase-3 activation at concentrations between 1 and 50 μ M (Choi et al. 2005). Thus, MAPK-related signalling may regulate expression of apoptotic genes, preventing apoptosis, and promoting cell survival. Another observation demonstrates that EGCG at concentrations between 5 and 25 μ M

inhibits angiotensin II-induced endothelial stress fibre formation and increased permeability via inactivation of p38/heat shock protein 27 (HSP27) pathway and suggests that EGCG may protect against endothelial barrier dysfunction and injury (Yang et al. 2010). EGCG treatment also increased the nuclear accumulation, anti-oxidant response element (ARE) binding, and transcriptional activity of nuclear factor erythroid 2-related factor 2 (Nrf2). Furthermore, EGCG activated Akt and ERK1/2. These findings suggest that Nrf2 mediates EGCG-induced expression of some representative anti-oxidant enzymes, possibly via Akt/PKB and ERK1/2 signalling, which may provide the cells with acquired anti-oxidant defence capacity to survive the oxidative stress (Na et al. 2008). In addition to MAPKs pathway, flavonoids and their metabolites have been also shown to modulate cell survival signalling due to their interaction with the PI3K/Akt pathway (Kyoung et al. 2010). The PI3K/Akt pathway is one of the strongest intracellular pro-survival signalling systems. EGCG activated Akt and ERK1/2 signalling cascade in MCF10A cells (Na et al. 2008). This effect was mediated partially via the activation of the downstream pAkt and pBad pathways.

EGCG and other flavanols found in green tea are by far the most intensely investigated, with no studies being reported on other black tea constituents. Thus, besides its free radical scavenging, iron chelating, and anti-aggregating properties, EGCG can exert its action on different sites of the apoptotic pathways, including altering the expression of anti- and pro-apoptotic genes. These studies further implicate that green tea extract may also exert protection through controlling calcium homeostasis, activation of MAPK, PKC, anti-oxidant enzymes and survival genes, thus potentially preventing progression of PD.

6.4.2 Tea Polyphenols and *In Vivo* Models of PD

A number of studies have reported on the protective effects of tea polyphenols against brain damage in various animal models of PD

(Dajas et al. 2003; Mandel and Youdim 2004; Mandel et al. 2008). Studies have used either a single compound such as EGCG, or a complex mixture of extracts from tea (Mercer et al. 2005; Masuda et al. 2006; Weinreb et al. 2008; Chen et al. 2008). Green or black tea polyphenol extracts, as well as individual EGCG, attenuated striatal dopamine depletion and SN dopaminergic neurons loss when given chronically to mice, rats or monkeys treated with the parkinsonism-inducing neurotoxins, 1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine (MPTP) or 6-OHDA (Levites et al. 2001; Chaturvedi et al. 2006; Chen et al. 2014). One study concluded that EGCG does not protect against 6-OHDA-induced loss of nigrostriatal neurons in rats (Leaver et al. 2009). More recently, it was demonstrated that stand-alone polyphenols including EC, EGC, and EGCG protect, rescue and most importantly restore the impaired movement activity (climbing capability) induced by paraquat in *Drosophila* models of PD (Jimenez-Del-Rio et al. 2010). Significantly, these findings receive further support from a recent *in vivo* preclinical neurorescue/neurorestorative drug cocktail study, demonstrating that synergistically EGCG and rasagiline (whilst individually having no profound protective effect) almost completely restored nigrostriatal dopaminergic neuron degeneration caused by MPTP (Reznichenko et al. 2010). Therefore, in a combination therapy regime, EGCG may have the potential to complement the pharmacological activities of current drugs in PD (Chen et al. 2008). Table 6.3 summarizes the most relevant studies concerning the neuroprotective and neurorestorative activities of tea polyphenols in animal models concerning PD.

Understanding the *in vivo* effects of tea consumption is far from complete. Evidence that tea polyphenols are acting directly or indirectly as anti-oxidants *in vivo* exists, but is far more limited when compared to *in vitro* studies. Administration of green tea extract and, in one case, black tea extract, attenuated decreases in superoxide dismutase (SOD) activity caused by infection, ethanol or the carcinogen, 3-methylcolanthrene (Frei and Higdon 2003; Higdon and Frei 2003).

Table 6.3 Neuroprotective and neurorestorative activities of tea polyphenols in animal models of PD

PD model	Tea polyphenol and oral dose	Neuroprotection	References
6-OHDA rat	(i) Black tea extract (1.5 % <i>ad libitum</i>)	Improvement of spontaneous locomotion, striatal dopamine and anti-oxidant enzymes, prevention and rescue of SN dopaminergic neurons	Chaturvedi et al. (2006)
	(ii) Green tea (150 mg/kg/day)	Protected dopaminergic neurons and preserved the free radical scavenging capability of both the midbrain and the striatum.	Guo et al. (2007)
MPTP mice	(i) EGCG (2 and 10 mg/kg/day), or Green tea extract (0.5 and 1 mg/kg/day)	Prevented dopamine neuron loss and depletion in striatal dopamine and hydroxylase protein levels	Levites et al. (2001)
	(ii) EGCG (25 mg/kg/day), and green tea (5 ± 0.7 ml)	Prevented the loss of tyrosine hydroxylase (TH)-positive cells in the SN and of TH activity in the striatum, preserved striatal levels of dopamine and its metabolites	Choi et al. (2002)
	(iii) EGCG (5 mg/kg/day)	Rescue of striatal dopamine depletion and SN dopaminergic neurons loss-induced by MPTP	Reznichenko et al. (2010)
	(iv) Theaflavins (10 mg/kg/day)	Reduces oxidative stress, improves motor behaviour and expression of dopamine transporter and vesicular monoamine transporter 2 in striatum and SN	Anandhan et al. (2012a, b)
MPTP monkey	Green tea extract (40 mg/kg/day)	Alleviates motor impairments and dopaminergic neuronal injury in the SN, inhibition of MPTP-induced accumulation of neurotoxic α S oligomers in the striatum and other brain regions	Chen et al. (2014)

6-OHDA 6-hydroxydopamine, MPTP N-methyl-4-phenyl-1,2,3,6-tetrahydropyridine, SN substantia nigra

Another study showed that green tea consumption prevented decline in glutathione peroxidase, indicating a protection in age-related oxidative damage in the brain (Kishido et al. 2007). While green and black tea administration improved the resistance of lipoproteins to *ex vivo* oxidation in several animal models, the improvement was generally much less than that conferred by supplementation with other anti-oxidants. Such an observation raises the question of whether tea polyphenols are present in sufficient quantities *in vivo* to work through an anti-oxidant mechanism (Lotito and Frei 2006; Stevenson and Hurst 2007; Spencer et al. 2009). Some studies have shown that blood concentrations of polyphenols are not high enough to add significantly to the body's total anti-oxidant capacity (D'Archivio et al. 2007; Ghosh and Scheepens 2009). In fact, it was estimated that, after ingestion, up to 95 %

of polyphenols undergo structural modification, that in turn may change the 'biological activities' of polyphenols as observed in the *in vitro* studies (Lotito and Frei 2006; Stevenson and Hurst 2007). In turn, there is evidence that polyphenols may be working through other mechanisms, for instance, by protecting endogenous anti-oxidant enzymes such as ascorbic acid in the human body against oxidation consequently improving the overall anti-oxidant level *in vivo* (Aron and Kennedy 2008). This indirect contribution to anti-oxidant effects requires polyphenols at concentrations much lower than would be essential for chemical anti-oxidant protection *in vitro* (Spencer et al. 2009). In other words, polyphenols may act beyond their anti-oxidant activity when not present at suitable concentrations to exert anti-oxidative effects (Saura-Calixto et al. 2007).

It is well known that tea polyphenols differ in their bioavailability and bioactivity. The rather poor bioavailability of EGCG needs to be considered when results obtained *in vitro* are extrapolated to situations *in vivo*. Most of the ingested EGCG is actually not absorbed in the blood, since absorption takes place in the small intestine and substantial quantities pass from the small to the large intestine where it undergoes further degradation by the action of local microbiota (Auger et al. 2008; Stalmach et al. 2009; Roowi et al. 2010). Bioavailability studies for EGCG indicate that peak plasma concentrations are reached after 1–2 h in healthy subjects with one oral dose (800 mg) in the morning after an overnight fasting period; these levels diminish gradually to undetectable levels within 24 h (Chow et al. 2005). The elimination half-life of EGCG is around 3.4 ± 0.3 h (Lee et al. 2002). It was argued that since green and black tea display similar anti-oxidant potential *in vivo*, despite containing different classes of polyphenols, it can be assumed that at least some of the thearubigins and theaflavins are absorbed (Leung et al. 2001; Rietveld and Wiseman 2003). However, the bioavailability of individual thearubigins and theaflavins has thus far not been directly evaluated in human studies. Nevertheless, although the bioavailability of tea flavonoids is low, repeated consumption of tea drinks resulted in a significant accumulation of flavanols in most body organs with relatively high peak plasma levels (Henning et al. 2008). The lack of a precise analytical method to estimate the presence of the more bioavailable flavanols in green tea compared to theaflavins and thearubigins in black tea *in vivo* may lead to an underestimation of the bioactivity of black tea when compared to green tea (Kumar and Pandey 2013). There has been extensive debate about whether the addition of milk to tea affects the bioavailability of flavonoids. Studies have clearly shown that plasma levels of polyphenols such as flavanols increased significantly after tea consumption and were unaffected by the addition of milk even when considering *in vivo* anti-oxidant potential (Kyle et al. 2007). Assuming that small micromolar quantities of tea polyphenols can exhibit bioactivity *in vivo*, cur-

rent data suggests that long-term consumptions of tea can result in the absorption and retention of sufficient amounts of polyphenols to exert the required effects in plasma and tissues.

Despite the increasing amount of evidence favouring the bioavailability of polyphenols in the systemic circulation, less information is available regarding their ability to cross the blood-brain barrier (BBB) and reach the CNS (Williamson and Manach 2005; Crozier et al. 2009). Flavanols were shown to cross a cellular model of the BBB in a time-dependent and stereo-selective manner (Faria et al. 2011). Multiple animal models have demonstrated that EGCG and EC cross the BBB, reaching a concentration of 0.5 nmol/g in rat brain in the case of EGCG consumption (500 mg/kg) and to co-localise within the brain tissues independently of their route of administration (Nakagawa and Miyazawa 1997; Abd El Mohsen et al. 2002; Adachi et al. 2006). These findings suggest that tea polyphenols are potential biologically active nutrients for direct neuroprotective and neuromodulatory actions. Although the uptake and distribution of dietary polyphenols within the brain is somewhat documented, more uncertainty revolves around the dosage, absorption, metabolism, tissue distribution, and intracellular accumulation and excretion of such compounds. Thus future work is required to investigate this further (Schaffer and Halliwell 2012; Vauzour 2012).

6.5 Clinical Studies with Tea Polyphenols in PD

Despite numerous efforts in the search for disease-modifying therapies in PD, currently the only approved treatment, apart from rasagiline, are agents that target symptoms without modifying the actual pathophysiology of the disease (Olanow et al. 2009). A double-blind, randomized, placebo-control delayed clinical study to evaluate the safety, tolerability, and efficacy of green tea polyphenols in slowing disease progression in patients with early PD, was conducted by the Chinese Parkinson Study Group (CPSG); 410 untreated people with early

PD were enrolled at 32 Chinese Parkinson Study Group sites. Participants were randomized to 0.4, 0.8, or 1.2 g of green tea polyphenols daily or placebo in the first phase of the study, and at 6 months the placebo group switched to 1.2 g of green tea polyphenols daily for 6 more months. Although insomnia was slightly increased, it was found that green tea polyphenols were well tolerated and provided a mild symptomatic relief in early untreated PD (Chan et al. 2009). Data from Chow and colleagues also confirmed that a daily dose of 800 mg caffeine-free EGCG for 4 weeks is safe and well tolerated in healthy human subjects (Chow et al. 2003).

Nonetheless, to date clinical trials so far have failed to identify compounds such as tea polyphenols with compelling proof for disease-modifying properties. One important reason is the lack of a reliable biomarker that can be used to track disease progression (Gerlach et al. 2012). An urgent need for suitable biomarkers in PD together with well-designed controlled studies to assess a risk reduction of PD with tea polyphenols should be prioritized so as to support the evidence derived from *in vitro* and *in vivo* studies.

6.6 Conclusion

Tea is one of the most frequently consumed beverages in the world and its medicinal effects have a long, rich history. In this review we have shown that in the last decade there has been an extensive interest in tea polyphenols as a potential therapeutic agent in PD (Mandel and Youdim 2004; Spencer 2008). Indeed, there is convincing evidence to suggest that the consumption of green and black tea exerts a beneficial effect in reducing the risk of PD, due to its polyphenolic content which exhibits numerous biochemical activities. From the epidemiological data reviewed, it was determined that the dose for the daily intake of tea should be around 2–3 cups/day, in order to induce neuroprotection. The observed beneficial effect, mostly from case–control studies, of tea drinking should now be investigated further in large prospective cohort studies.

There seems to be a consensus that the efficacy of green tea is likely to be mediated by the effects of EGCG, whilst the main bioactive constituents of black tea are the theaflavins. It is clear that EGCG has been the polyphenolic compound of choice most extensively investigated both *in vivo* and *in vitro*, and therefore future studies should now address the effects of other important tea polyphenols, especially theaflavins. While many of the mechanisms underpinning their beneficial effects have been highlighted, it has become clear that apart from the classical anti-oxidant properties, tea polyphenols may in part incur neuroprotection in PD through specific cell signalling pathways and prevention of α S aggregation.

Finally, the extent of their contribution *in vivo*, and at physiological relevant concentrations remains to be ascertained. Conflicting epidemiological inferences and discrepancies between *in vitro* and *in vivo* studies may be due to erratic bioavailability of tea polyphenols. Although more urgent work needs to be done to prove whether tea polyphenols can be translated in PD patients and to clarify their absorption, metabolism, and potential toxicity in humans, their multiple biological activities, and especially in combination with other compounds that possess neuroprotective moieties, may offer a superior therapeutic effect in delaying the initiation and progression of PD.

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The Effect of (–)-Epigallo-catechin-(3)-gallate on Amyloidogenic Proteins Suggests a Common Mechanism

7

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Abstract

Studies on the interaction of the green tea polyphenol (–)-Epigallocatechin-3-gallate (EGCG) with fourteen disease-related amyloid polypeptides and prions Huntingtin, Amyloid-beta, alpha-Synuclein, islet amyloid polypeptide (IAPP), Sup35, NM25 and NM4, tau, MSP2, semen-derived enhancer of virus infection (SEVI), immunoglobulin light chains, beta-microglobulin, prion protein (PrP) and Insulin, have yielded a variety of experimental observations. Here, we analyze whether these observations could be explained by a common mechanism and give a broad overview of the published experimental data on the actions of EGCG. Firstly, we look at the influence of EGCG on aggregate toxicity, morphology, seeding competence, stability and conformational changes. Secondly, we screened publications elucidating the biochemical mechanism of EGCG intervention, notably the effect of EGCG on aggregation kinetics, oligomeric aggregation intermediates, and its binding mode to polypeptides. We hypothesize that the experimental results may be reconciled in a common mechanism, in which EGCG binds to cross-beta sheet aggregation intermediates. The relative position of these species in the energy profile of the amyloid cascade would determine the net effect of EGCG on aggregation and disaggregation of amyloid fibrils.

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Keywords

Epigallocatechin-3-gallate (EGCG) • Amyloid polypeptides • Aggregation • Fibrils

7.1 Introduction

The polyphenol (–)-Epigallocatechin-3-gallate (EGCG) is produced by the tea plant (*Camellia sinensis*) and accounts for about 10 % of the dry weight of its leaves (Graham 1992). The flavonoid EGCG belongs to the subclass of flavan-3-ols which have an unsaturated C-ring with the B-ring attached to their C2-atom (Fig. 7.1a). In EGCG, the B-ring is attached in epi conformation and the 3-ol of the C-ring is substituted by an 3-O-gallate function (Beecher 2003). EGCG has been found to inhibit formation of amyloid fibrils and to bind to existing amyloid fibrils and remodel them into non-amyloid aggregates (Fig. 7.1b) (Ehrnhoefer et al. 2006, 2008; Bieschke et al. 2010). Its anti-aggregation mechanism has since been studied for 14 amyloidogenic peptides and proteins. At first glance, the data seem to suggest that EGCG has quite different effects on the different polypeptides.

Our review aims to compare these effects on a phenomenological and mechanistic level. To do so, we will concentrate on disease-related polypeptides and prions. In the first part, we will give a broad overview of the published experimental data on the action of EGCG, while in the second part we will discuss how the experimental results might be reconciled in a general mechanism that leads to the inhibition of amyloidogenic aggregation and the disaggregation of amyloid fibrils.

7.2 Phenomenological Overview of EGCG

About 10 years ago, it was first reported that green tea extract and EGCG reduced the cytotoxicity of the Amyloid-beta peptide (A β) involved in Alzheimer's Disease (AD) (Levites

et al. 2003) and also reduced the amount and size of amyloid deposits in APP transgenic mice (Rezai-Zadeh et al. 2005). Likewise, it diminished the formation of toxic aggregates in yeast and *Drosophila* models of Huntingtin protein (Htt) aggregation (Ehrnhoefer et al. 2006). These early results prompted a large number of studies on the effect of EGCG on various amyloidogenic proteins. Tables 7.1, 7.2, and 7.3 present an overview of the effects of EGCG extracted from data that were published prior to August 2014. Empty fields mark characteristics for which no data are available for the respective polypeptide. Some characteristics have been addressed in an extensive number of studies; in these cases, the complete reference list is included in the overview table while selected studies are referenced in the text.

7.2.1 Amyloid Toxicity

Amyloid fibrils may have functional roles (Chapman et al. 2002; Berson et al. 2003; Fowler et al. 2006). Several functional prions that possess amyloid structural elements have been identified in yeast and other fungi (Shorter and Lindquist 2005; Coustou et al. 1997). However, in many polypeptides amyloid formation is toxic to cells. Indeed, deposition of misfolded polypeptides in amyloid or amyloid-like aggregates is characteristic of some of the most prominent neurodegenerative diseases: the Amyloid-beta peptides (A β) derived from the Amyloid Precursor Protein (APP) and the tau protein in Alzheimer's disease (AD) (Golde et al. 1993; Kosik et al. 1986), Huntingtin protein (Htt) in Huntington's disease (HD) (Trottier et al. 1995; Scherzinger et al. 1997), alpha-Synuclein (α -Syn) in Parkinson's disease (PD) (Takeda et al. 1998), and the prion protein PrP in transmissible spongiform encephalopathies (TSE) (Prusiner 1998). Amyloid deposits also form in systemic diseases: globular

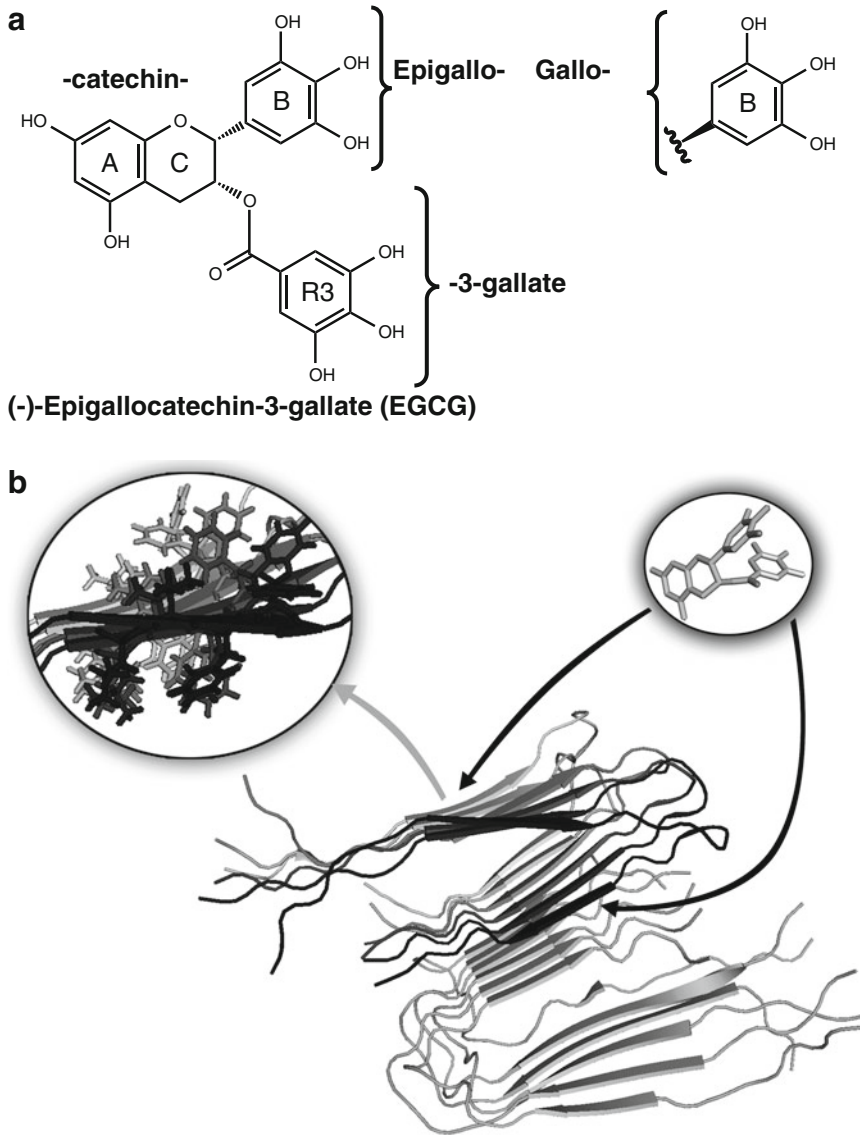


Fig. 7.1 (–)-Epigallocatechin-3-gallate (EGCG) and its interaction with A β amyloid fibrils **(a)** EGCG is a flavonoid of the subclass flavan-3-ol having an unsaturated C-ring with the B-ring attached to its C2-atom. In EGCG, the B-ring is attached in epi conformation and the 3-ol of the C-ring is substituted by a 3-O-gallate function. The phenyl rings are di- or trisubstituted by phenol groups, thus making EGCG a polyphenol (Beecher 2003). **(b)** In A β fibrils, the monomers form a hairpin structure, with the N-terminal domain facing the fibril surface, while the C-terminal domain is directed towards the fibril core. Two

or three monomers build the fibril base. The monomers within the fibrils are stacked on top of each other with adjacent monomers building intramolecular cross- β -sheets along the fibril axis (Lu et al. 2013; Petkova et al. 2006; Luhrs et al. 2005; Colletier et al. 2011). EGCG binding is indicated by black arrows. EGCG likely binds along the LVFF-motif (enhanced with side chain structure) (Grelle et al. 2011; Lopez del Amo et al. 2012; Wang et al. 2012b). This figure was made using Pymol and structure coordinates were taken from PDB entry 2LMO (Petkova et al. 2006)

Table 7.1 Effect of EGCG on disease-related polypeptides and prions (part 1)

Influence of EGCG	Htt	A β	α -Syn	IAPP
Aggregate morphology				
Prevents fibrils/aggregate formation		5	12, 23	25
Redirects into small amorphous aggregates		Y		
Leads to monomer depletion				
Disaggregates fibrils	1	12, 21	23	25, 26, 28
Remodels into large amorphous aggregates		12, 21	12	25, 26, 28
Kinetics				
Slows ThT kinetics		5, 14, 16, 17	5, 12	25, 27, 28
Accelerates ThT kinetics		N (5)	N (5, 12)	N (25,27, 28)
Reduces or inhibits seeding-competence		5, 7, 18, 21	5, 12	25, 26
Inhibits prion propagation				
Structural				
Redirects into non-SDS-stable species	1	N (12)		
Redirects into semi-SDS-stable species		5, 16	5, 12	
Redirects into SDS-stable species		12, 14, 21	12, 14	
Reduces proteinase K-resistance				
Redirect into non-A11-specific oligomers		17	5	
Uncovers antibody binding epitope				
Reduces ThT/CR fluorescence amplitude		5, 12, 14, 16, 17, 21	5, 12	25, 27, 28
EGCG structure and binding				
Non-epi-gallo-moiety reduces efficiency	1	8, 12		
Gallate-moiety necessary for effectivity	1	8, 12		
Binds to oligomers		5	5	28
Direct remodeling		21, 12	12	
Interacts with specific residues				
Hydrophobic interactions		9, 18, 19		
H-Bonding				
π - π stacking		18		N (26)
Schiff-base formation		N (9, 19, 21)		
Disulfide bridge				
Destruction of salt-bridges		N(9, 18, 19, 21)		N (26)
Toxicity				
Reduces membrane permeabilization				
Inhibits cytotoxicity (cell models)	1	16, 20	20, 24	25
Inhibits deposits in vivo (animal models)		10 papers ^a	5, 12, 24	25
		3, 4, 6, 7, 10, 11, 15		

(continued)

Table 7.1 (continued)

Influence of EGCG	Htt	A β	α -Syn	IAPP
Reduces cognitive impairment in mice		4, 6		
Inhibits deposits in vivo (humans)				
Phase 2 or 3 clinical trial	Y	Y	Y	

N() indicates studies with negative results, blank spaces indicate that no experimental results are available

A β Amyloid-beta, Htt Huntingtin, IAPP Islet Amyloid Polypeptide, α -Syn alpha-Synuclein; 1 (Ehrnhoefer et al. 2006), 2 (Levites et al. 2003), 3 (Rezai-Zadeh et al. 2005), 4 (Rezai-Zadeh et al. 2008), 5 (Ehrnhoefer et al. 2008), 6 (Lee et al. 2009b), 7 (Lee et al. 2009a), 8 (Lin et al. 2009), 9 (Wang et al. 2010), 10 (Giunta et al. 2010), 11 (Abbas and Wink 2010), 12 (Bieschke et al. 2010), 13 (He et al. 2011), 14 (Grelle et al. 2011), 15 (Dragicevic et al. 2011), 16 (Gauci et al. 2011), 17 (Sinha et al. 2012), 18 (Lopez del Amo et al. 2012), 19 (Wang et al. 2012b), 20 (Camilleri et al. 2013), 21 (Palhano et al. 2013), 22 (Lee et al. 2013), 23 (Caruana et al. 2011), 24 (Lorenzen et al. 2014), 25 (Meng et al. 2010), 26 (Cao and Raleigh 2012), 27 (Suzuki et al. 2012), 28 (Young et al. 2014)

^a2, 5, 8, 12, 13, 14, 15, 17, 18, 21

monoclonal immunoglobulin light chains (LC) and Transthyretin (TTR) deposits form in the heart and in other tissues in systemic Light Chain Amyloidosis (AL) (Solomon et al. 1982) and in Transthyretin-Amyloidosis, respectively (Saraiva et al. 1984). In Diabetes Mellitus type II (DM II) the Islet Amyloid Polypeptide (IAPP, amylin) forms amyloid deposits in the insulin producing islet- β -cells (Johnson et al. 1989). *In vitro*, it was observed that also insulin is capable of forming amyloid fibrils under physiological conditions, albeit at high concentrations of 345 μ M (Wang et al. 2012a). Diabetes mellitus has large economic implications; about 347 million people worldwide suffer from DM type I and type II (Danaei et al. 2011), which dwarfs even prevalence of AD (estimated 44 million cases in 2013) (Prince et al. 2014).

Patients undergoing long-term hemodialysis are at risk of developing dialysis-related amyloidosis (DRA), where β_2 -Microglobulin (β_2 m) is deposited, a light chain of the major histocompatibility complex I (MHC-I) (Grey et al. 1973) that is prone to form amyloid deposits (Gejyo et al. 1985, Linke 1985).

Interestingly, two other devastating diseases are also indirectly connected to amyloid formation. An amyloid forming-protein derived from human semen was found to enhance the infectious potential of HIV and was therefore named semen-derived enhancer of virus infection (SEVI). The parasite plasmodium falciparum

involved in malaria pathogenesis produces a highly abundant surface protein, the merozoite surface protein 2 (MSP2) which is considered as a candidate for a malaria vaccine (Genton et al. 2002). It was shown that MSP2 has also an amyloidogenic nature (Yang et al. 2007). While this is by no means a complete list of amyloid diseases, experimental data on EGCG have been collected for all of these polypeptides, illustrating the broad interest in EGCG that led to a large body of experimental data that are available for mechanistic analysis.

7.2.2 Reduction in Cytotoxicity of Amyloidogenic Proteins

EGCG reduced the toxicity of amyloidogenic polypeptides related to a number of neurodegenerative diseases and systemic diseases that were listed above. EGCG treatment reduced cytotoxicity of A β (Levites et al. 2003; Ehrnhoefer et al. 2008; Bieschke et al. 2010) [see also Table 7.1], Htt (Ehrnhoefer et al. 2006), α -Syn (Ehrnhoefer et al. 2008; Bieschke et al. 2010; Lorenzen et al. 2014), IAPP (Meng et al. 2010) and tau (Wobst et al. 2015) in cellular models. Experiments probing the effect of EGCG on TTR toxicity in cell culture were inconclusive (Miyata et al. 2010).

Amyloid toxicity is believed to be linked to the presence of oligomeric aggregation intermediates (Walsh et al. 2002a; Haass and Selkoe 2007).

Table 7.2 Effect of EGCG on disease-related and polypeptides and prions (part 2)

Influence of EGCG	Sup35 NM25	Sup35 NM4	tau	MSP2	SEVI
Aggregate morphology					
Prevents fibrils/aggregate formation	30	N (30)	31		34, 36
Redirects into small amorphous aggregates			31		
Leads to monomer depletion					
Disaggregates fibrils	30	N (30)	31	33	34
Remodels into large amorphous aggregates	30	N (30)		33	34
Kinetics					
Slows ThT kinetics			31	32, 33	34
Accelerates ThT kinetics			N (31)	N (32, 33)	N (34)
Reduces or inhibits seeding-competence	30	N (30)			
Inhibits prion propagation	30	N (30)			
Structural					
Redirects into non-SDS-stable species					
Redirects into semi-SDS-stable species					
Redirects into SDS-stable species				32, 33	
Reduces proteinase K-resistance					
Redirect into non-A11-specific oligomers			31		
Uncovers antibody binding epitope			31		
Reduces ThT/CR fluorescence amplitude	30	30	31	32, 33	34
EGCG structure and binding					
Non-epi-gallo-moiety reduces efficiency					34, 36
Gallate-moiety necessary for effectivity	30	30			34, 36
Binds to oligomers			31	32	34
Direct remodeling					
Interacts with specific residues				32	34
Hydrophobic interactions					
H-bonding					
π - π stacking					
Schiff-base formation				N (32)	-34
Disulfide bridge				N (32)	
Destruction of salt-bridges					
Toxicity					
Reduces membrane permeabilization					
Inhibits cytotoxicity (cell models)			31		
Inhibits deposits in vivo (animal models)					
Reduces cognitive impairment in mice					
Inhibits deposits in vivo (humans)					
Phase 2 or 3 clinical trial					

N() indicates studies with negative results; blank spaces indicate that no experimental results are available
Sup35 NM25 prion strain (head region 21–38, center region 39–90, tail region 91–106), *Sup35 NM4* prion strain (head region 21–38, center region 39–78, tail region 79–96), *tau* HisK18 Δ K280 fragment of tau protein, *MSP2* Plasmodium falciparum merozoite surface protein 2, *SEVI* semen-derived enhancer of virus infection, 30 (Roberts et al. 2009), 31 (Wobst et al. 2015), 32 (Chandrashekar et al. 2010), 33 (Chandrashekar et al. 2011), 34 (Hauber et al. 2009), 36 (Popovych et al. 2012)

Table 7.3 Effect of EGCG on disease-related polypeptides and prions (part 3)

Influence of EGCG	LC	TTR	β 2-m	PrP	Ins
Aggregate morphology					
Prevents fibrils/aggregate formation		43	N (49)		51
Redirects into small amorphous aggregates		44			51
Leads to monomer depletion		43			N (51)
Disaggregates fibrils					
Remodels into large amorphous aggregates		44			
Kinetics					
Slows ThT kinetics	N (41)		49		
Accelerates ThT kinetics	41		N (49)		
Reduces or inhibits seeding-competence				50	
Inhibits prion propagation					
Structural					
Redirects into non-SDS-stable species	N (41)				
Redirects into semi-SDS-stable species	41	43			
Redirects into SDS-stable species	N (41)				
Reduces proteinase K-resistance				50	
Redirect into non-A11-specific oligomers					
Uncovers antibody binding epitope					
Reduces ThT/CR fluorescence amplitude	41	43	49		
EGCG structure and binding					
Non-epi-gallo-moiety reduces efficiency					
Gallate-moiety necessary for effectivity				50	
Binds to oligomers		43			
Direct remodeling					
Interacts with specific residues					
Hydrophobic interactions					
H-bonding					
π - π stacking					
Schiff-base formation					
Disulfide bridge					
Destruction of salt-bridges					
Toxicity					
Reduces membrane permeabilization					
Inhibits cytotoxicity (cell models)					
Inhibits deposits in vivo (animal models)		45, 47			
Reduces cognitive impairment in mice					
Inhibits deposits in vivo (humans)	37, 38	46			
Phase 2 or 3 clinical trial	Y	Y			

N() indicates studies with negative results; blank spaces indicate that no experimental results are available

LC immunoglobulin light chain protein, TTR transthyretin (studies with variants: wt/V30M/E54K/Y38F/L55P), β 2-m β 2-Microglobulin, PrP prion protein, Ins Insulin 37 (Mereles et al. 2008), 38 (Mereles et al. 2010), 41 (unpublished data), 43 (Miyata et al. 2010), 44 (Ferreira et al. 2011), 45 (Ferreira et al. 2012b), 46 (Kristen et al. 2012), 47 (Ferreira et al. 2012a), 49 (Woods et al. 2011), 50 (Rambold et al. 2008), 51 (Wang et al. 2012a)

However, the mechanism of toxicity has not yet been completely clarified. For A β and α -Syn, it was found that aggregates can permeabilize vesicular and mitochondrial membranes (Rodrigues et al. 2000; Volles et al. 2001). EGCG was shown to inhibit permeabilization of model membranes and mitochondrial membranes (Gauci et al. 2011; Caruana et al. 2011; Camilleri et al. 2013). In an *ex vivo* study with human semen samples from 47 individuals, the majority of samples contained SEVI species and EGCG was able to efficiently inhibit the SEVI-mediated HIV activity (Hauber et al. 2009).

7.2.3 Reduction of Amyloid Deposits and Toxicity in Higher Organisms

EGCG was shown to reduce amyloid deposition in animal models of protein misfolding disorders. Treatment with green tea extract rich in EGCG lowered the load of amyloid aggregates in *Caenorhabditis elegans* (Abbas and Wink 2010) and in AD mouse models (Rezai-Zadeh et al. 2005; Giunta et al. 2010; Dragicevic et al. 2011) [see also Table 7.1], where it also reduced cognitive impairment (Rezai-Zadeh et al. 2008; Lee et al. 2009b). Depletion of amyloid deposits was observed in mouse models of Transthyretin Amyloidosis (Ferreira et al. 2012a, b), while depletion of amyloidogenic deposits in the human heart was observed in a phase II clinical trial after EGCG treatment (Kristen et al. 2012). Treatment with EGCG of a case of AL-Amyloidosis after several unsuccessful cycles of chemotherapy resulted in reduction of amyloid deposits in the patients' heart (Mereles et al. 2008). The same group performed a retrospective study on the influence of green tea and EGCG consumption on amyloid deposition in AL patients (Mereles et al. 2010). Phase II clinical studies by two European AL-Amyloidosis Treatment Centers are currently in progress (Schönland 2013; Merlini 2013). Phase II clinical studies have also been performed on AD, PD and HD (Chan 2007; Friedemann and Dörr 2009; Priller 2011).

7.2.4 Pharmacological Aspects of Aggregation Inhibition

Therapeutic use of EGCG is complicated by several pharmacokinetic drawbacks, most notably the widely variable bioavailability after oral consumption (Hunstein 2007). Nevertheless, it may be a treatment option worth evaluating, especially in rare diseases that are otherwise unprofitable for drug development. Exchanging the B-ring epigallo-moiety (Beecher 2003) by a gallo-moiety (Fig. 7.1) reduced the efficiency of the catechin on Htt aggregation (Ehrnhoefer et al. 2006), on A β aggregation (Bieschke et al. 2010) and A β cytotoxicity (Lin et al. 2009), and on SEVI-mediated infectivity (Hauber et al. 2009; Popovych et al. 2012). Depleting the C3-atom gallate function from the C-Ring showed comparable effects in the mentioned studies. In contrast to EGCG, Epigallocatechin (EGC) was neither able to disaggregate fibrils of the Sup35 NM25 strains prionogenic N-terminal domain (NM25) nor to inhibit its prion propagation (Roberts et al. 2009). EGCG treatment led to lysosomal degradation of PrP^C and subsequent accumulation of PrP^{Sc} in transiently transfected cell lines. While EGCG efficiently interfered with the PrP^{Sc} accumulation, four times the amount of EGC was necessary to achieve the same effect (Rambold et al. 2008).

7.3 Changes in Aggregate Morphology of Amyloid Precursors and Amyloidogenic Species

7.3.1 Folding State of Amyloid Precursors and Amyloidogenic Species

Protein misfolding can result in the formation of rope-like or straight amyloid fibrils that have highly stable cross- β sheet structures, which are aligned along the fibril axis (Chiti and Dobson 2006). The amyloidogenic species could either be a fragment of a precursor polypeptide like A β ,

Htt-exon1 and Insulin (Permutt et al. 1981), or entire proteins like α -Syn (Ulmer et al. 2005), LC protein (Redegeld and Nijkamp 2003) and TTR (Wojtczak et al. 1992). The native structure of amyloid precursors range from highly unstructured proteins like Htt (Zhang et al. 2013) and α -Syn (Dedmon et al. 2005) to highly structured β -sheet rich proteins like β 2m (Bjorkman et al. 1987), TTR (Wojtczak et al. 1992) and LC (Edmundson et al. 1993), or α -helix rich proteins like the PrP (90–232) (Riek et al. 1996).

Irrespective of the starting point, the formation of the cross- β -sheet structure requires at least a partial unfolding or structural rearrangement of the amyloid precursor. The presence of EGCG during aggregation reduced the β -sheet content of all the protein aggregates for which secondary structure was measured, namely α -Syn (Ehrnhoefer et al. 2008; Bieschke et al. 2010), MSP2 (Chandrashekar et al. 2010, 2011), PrP (Rambold et al. 2008) and Insulin (Wang et al. 2012a).

7.3.2 Redirection of Aggregation into Non-amyloidogenic Species

The question of whether EGCG prevents the formation of amyloidogenic fibrils and aggregates was addressed in most studies with amyloidogenic polypeptides for which EGCG data was available. Its effect on *de novo* amyloid formation depends on the amyloidogenic protein (Fig. 7.2a). In most proteins, EGCG prevented amyloid fibril formation, and redirected the assembly process to generate amorphous aggregate species. This was observed for Htt (Ehrnhoefer et al. 2006), A β (Ehrnhoefer et al. 2008; Lopez del Amo et al. 2012), α -Syn (Bieschke et al. 2010; Suzuki et al. 2012), the HIV-mediating SEVI (Hauber et al. 2009; Popovych et al. 2012), IAPP (Meng et al. 2010; Suzuki et al. 2012), and Insulin (Wang et al. 2012a). EGCG was on the other hand unable to prevent fibril formation by β 2m (Woods et al. 2011), the prion protein (Rambold et al. 2008), or heparin-induced fibril formation of tau (Wobst et al. 2015). However, EGCG completely

prevented the formation of β -sheet rich aggregates of an aggregation-prone mutant tau (His-K18 Δ K280) in the absence of heparin (Wobst et al. 2015). The crucial rate limiting step in TTR-amyloid formation involves monomerization of the TTR tetramer (Hammarstrom et al. 2003). Here too, EGCG redirected the aggregation of mutant TTR-L55P and TTR-Y78F from large amorphous aggregates into small amorphous aggregates (Ferreira et al. 2011). In another study, it prevented fibril formation of wt-TTR and TTR-V30M by preventing the monomerization of TTR (Miyata et al. 2010).

Remarkably the fibril formation of Sup35 NM25 was also redirected into amorphous aggregates, while fibril formation was not impaired by EGCG in the NM4 prion N-terminal domain (Roberts et al. 2009).

7.3.3 Remodeling of Pre-formed Amyloid Fibrils into Non-amyloidogenic Species

As discussed above, it is established that EGCG generally redirects the aggregation of an amyloid precursor. One could therefore ask whether it can also disrupt pre-formed amyloid fibrils, and whether such a disruption would result in similar supramolecular assemblies to those observed in aggregation in the presence of EGCG. Disaggregation of amyloid fibrils into comparable amorphous aggregate species was in fact reported in presence of EGCG for A β (Bieschke et al. 2010; Palhano et al. 2013), α -Syn (Bieschke et al. 2010; Caruana et al. 2011), tau His-K18 Δ K280 (Wobst et al. 2015), IAPP (Meng et al. 2010; Cao and Raleigh 2012; Young et al. 2014), SEVI (Hauber et al. 2009) and the variants TTR-L55P and TTR-Y38F (Ferreira et al. 2011). Even fibrils of Sup35 were either disrupted and remodeled into large amorphous aggregates by EGCG (NM25), or not altered (NM4) corresponding to the results of *de novo* aggregation in presence of EGCG (Roberts et al. 2009). Additionally, EGCG was able to remodel preformed fibrils of MSP2 into large amorphous aggregates (Chandrashekar et al. 2011).

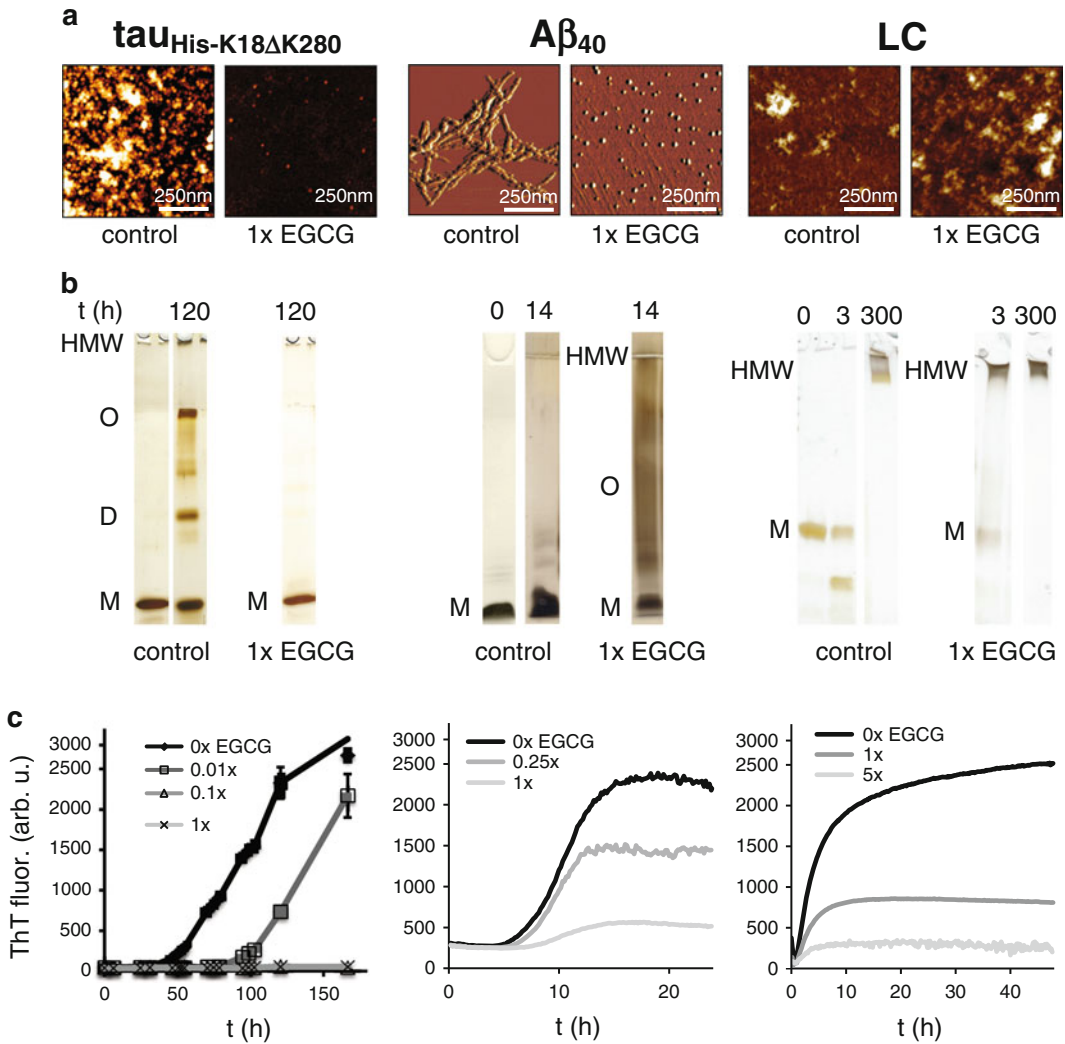


Fig. 7.2 Influence of EGCG on aggregation kinetics, aggregate morphology and SDS-stability of tau, A β and LC. **(a)** Aggregation kinetics in absence (*black*) and presence (*gray*) of EGCG, where 1 \times mean equimolar concentrations of the amyloid precursor and EGCG. **(b)** Aggregate morphology after aggregation in absence and presence of EGCG. **(c)** SDS-stability (tau, A β) and semi-SDS-stability

(LC, samples were neither reduced nor boiled) of aggregates formed in absence and presence of EGCG. With tau, EGCG prevents formation of SDS-stable aggregates while with A β it leads to SDS-stability indicating structural changes induced by EGCG. In LC, the formation of semi-SDS-stable aggregates was accelerated

7.3.4 Seeding Competence of EGCG-Induced Aggregates

A hallmark of amyloidogenic aggregates is their ability to incorporate monomeric polypeptides into amyloid structures. This process is referred to as ‘seeding-competence’, which is also a key property in prion propagation (Jarrett and Lansbury 1993). A simple test for seeding competence

is to compare aggregation kinetics in the presence of a suspected amyloid precursor to those of unseeded aggregation (Ehrnhoefer et al. 2008). In mechanisms where nucleation is the rate-limiting step, the addition of seeding- or propagation-competent species will shorten or abrogate the lag phase.

Aggregates that resulted either from redirected monomer aggregation or from remodeling of

mature fibrils by EGCG were either seeding-incompetent or showed reduced seeding-competence of A β (Ehrnhoefer et al. 2008; Bieschke et al. 2010; Palhano et al. 2013), α -Syn (Ehrnhoefer et al. 2008; Bieschke et al. 2010) and IAPP (Meng et al. 2010; Cao and Raleigh 2012). The prion protein (Rambold et al. 2008) as well as the yeast prion NM25 were also transformed into propagation-incompetent species, while the propagation-competence of NM4 was not impaired (Roberts et al. 2009).

7.3.5 Altered Aggregate Stability and Conformation

Resistance against denaturation by boiling in the presence of Sodium Dodecyl Sulfate (SDS-stability) is a rapid and useful indicator of aggregate stability (Wanker et al. 1999). Upon fibril formation, many amyloidogenic polypeptides become resistant to SDS denaturation and to proteolytic digestion by proteinase-K (PK-resistance). Changes in both properties would indicate structural changes in the aggregation process, although the structural basis of these effects is poorly understood. The SDS-stability can be measured either by a filter retardation assay (FRA) (Wanker et al. 1999) or SDS-PAGE (Ehrnhoefer et al. 2008, Bieschke et al. 2010). The latter is performed in the presence of SDS but without heating or at a temperature gradient, and can resolve differences in stability more sensitively. It is worth noting that apparent molecular weight in (semi-) denaturing gels does not necessarily reflect the size of the aggregates in solution.

The response of different amyloidogenic proteins to EGCG with respect to SDS and PK-stability varies considerably (Fig. 7.2b). EGCG prevented the formation of SDS-stable aggregates of Huntingtin (Ehrnhoefer et al. 2006) and an aggregation-prone tau mutant His-K18 Δ K280, (Wobst et al. 2015). In contrast, EGCG induced highly SDS-stable aggregates of A β and α -Syn (Ehrnhoefer et al. 2008; Bieschke et al. 2010; Grelle et al. 2011; Gauci et al. 2011; Palhano et al. 2013). Similarly, we observed that EGCG accelerates the formation of SDS-stable species

of immunoglobulin light chains (LC, Fig. 7.2b). SDS-stable aggregates of A β and α -Syn are structurally different from SDS-resistant fibrils these polypeptides form in the absence of EGCG. In LC aggregation, EGCG accelerates the formation of semi-SDS-stable aggregates. MSP2 was also reported to produce SDS-stable oligomer species in the presence of EGCG (Chandrashekar et al. 2010, 2011). Rambold et al. (2008) found that EGCG reduced the PK-resistance of infectious PrP aggregates.

7.3.6 Conformational Change Can Be Probed by Antibody Binding

The anti-oligomer antibody A11 recognizes a common intermediate oligomeric structure in A β 40 and A β 42 (Kayed et al. 2003), Sup35 (Shorter and Lindquist 2004), β 2m (Ribeiro et al. 2012), α -Syn (Ehrnhoefer et al. 2008) and tau (Flach et al. 2012; Wobst et al. 2015). EGCG prevented the binding of A11 to A β (Sinha et al. 2012), α -Syn (Ehrnhoefer et al. 2008) and tau (Wobst et al. 2015) indicating that either structural changes in the protein aggregates or EGCG binding obscured the antibody binding sites.

7.4 Towards a Holistic Molecular Mechanism for EGCG-Amyloid Interaction

7.4.1 Altered Protein Aggregation Kinetics by EGCG

The benzothiazole dye Thioflavin T (ThT) binds amyloid, amyloid fibrils and cross- β -sheet aggregation intermediates with a characteristic red-shift of its emission spectrum, which does not occur in presence of unstructured monomers or oligomers (LeVine 1997). Therefore, it is frequently used to observe the aggregation kinetics leading to formation of amyloid fibrils by numerous amyloidogenic polypeptides (LeVine 1999).

EGCG reduced the ThT amplitude in most amyloidogenic polypeptides in a dose-dependent

manner (see Table 7.1). This could be either due to a change in aggregate morphology, to partial sequestration of the amyloidogenic polypeptide or to an inhibition of ThT binding by competitive EGCG binding. After aggregation under influence of EGCG, the ThT fluorescence of Sup35 prion strains and TTR was reduced (Roberts et al. 2009; Miyata et al. 2010). By measuring kinetics of ThT fluorescence, it was shown that EGCG also slowed down the aggregation kinetics of A β (Ehrnhoefer et al. 2008; Grelle et al. 2011; Gauci et al. 2011; Sinha et al. 2012), α -Syn (Ehrnhoefer et al. 2008; Bieschke et al. 2010), IAPP (Meng et al. 2010; Suzuki et al. 2012; Young et al. 2014), tau (Wobst et al. 2015), MSP2 (Chandrashekar et al. 2010, 2011), SEVI (Hauber et al. 2009) and β 2m (Woods et al. 2011). These findings suggest that inhibition of aggregation kinetics might be due to a general mechanism of action of EGCG. EGCG however did not affect the kinetics of fibril formation of the tau protein when this was induced by heparin (Wobst et al. 2015). This is in contrast to the efficiency of EGCG in inhibiting the formation of ThT-positive tau oligomers in the absence of heparin, even at high substoichiometric concentrations (Fig. 7.2c) (Wobst et al. 2015).

When examining the effect of EGCG on different monoclonal immunoglobulin light chains isolated from urine of patients suffering either from AL Amyloidosis (clinical amyloid deposits) or from Multiple Myeloma (no clinical amyloid deposits), the formation of ThT-positive species was observed for all light chains (Andrich and Bieschke 2014). Typical aggregation kinetics showed a rapid phase of less than two hours followed by a slow phase of several days. In the presence of EGCG, only the ThT signal during the slow aggregation phase was reduced (Fig. 7.2c). The fact that light chains from patients without clinical amyloid deposits showed also the formation of ThT-positive species suggests that not every ThT-positive species of LC necessarily forms amyloid fibrils in patients.

We will discuss below how these different effects of EGCG on aggregation kinetics – inhibition, neutral, or acceleration – might be reconciled within a single model.

7.4.2 EGCG Interacts with Oligomeric Amyloid Precursors

EGCG binds to oligomeric species of A β , α -Syn and tau (Ehrnhoefer et al. 2008; Wobst et al. 2015). EGCG binding to oligomers was also reported for IAPP (Young et al. 2014), MSP2 (Chandrashekar et al. 2010) and SEVI (Hauber et al. 2009).

A more difficult question to answer is, whether EGCG is able to bind to monomeric amyloid precursors. Nitro blue tetrazolium chloride (NBT) is a dye that can be used to stain EGCG (Paz et al. 1991). In aggregation assays, EGCG-specific staining by NBT was found for A β and α -Syn, and tau species that ran as monomers on SDS-PAGE (Ehrnhoefer et al. 2008). However, peptide species running as monomers on an SDS gel are not necessarily monomeric in solution. SDS-labile oligomers could bind EGCG and still disassemble during SDS-PAGE. These species would be distinct from monomers within the aggregation cascade. EGCG binding to native proteins, such as albumin has been observed at higher stoichiometric ratios (Ehrnhoefer et al. 2008; Bae et al. 2009; Nozaki et al. 2009).

EGCG binding to the aggregation-prone tau mutant His-K18 Δ K280 resulted in monomers with a slightly lower electrophoretic mobility than the monomers prior aggregation, however the sub-stoichiometric effect of EGCG makes it likely that it preferentially binds to oligomeric aggregation intermediates rather than monomeric proteins (Wobst et al. 2015). Similarly, EGCG binding to the monomeric A β peptide was observed at a broad range of stoichiometries (Wang et al. 2010, 2012b), suggesting that multiple binding modes of EGCG exist to the A β peptide, possibly with a high affinity binding to oligomeric aggregation intermediates and lower affinity binding to the monomeric peptide. We postulate that the former high affinity interaction, rather than the latter low affinity interaction, is responsible for the inhibitory effect of EGCG on amyloid formation (Fig. 7.3a).

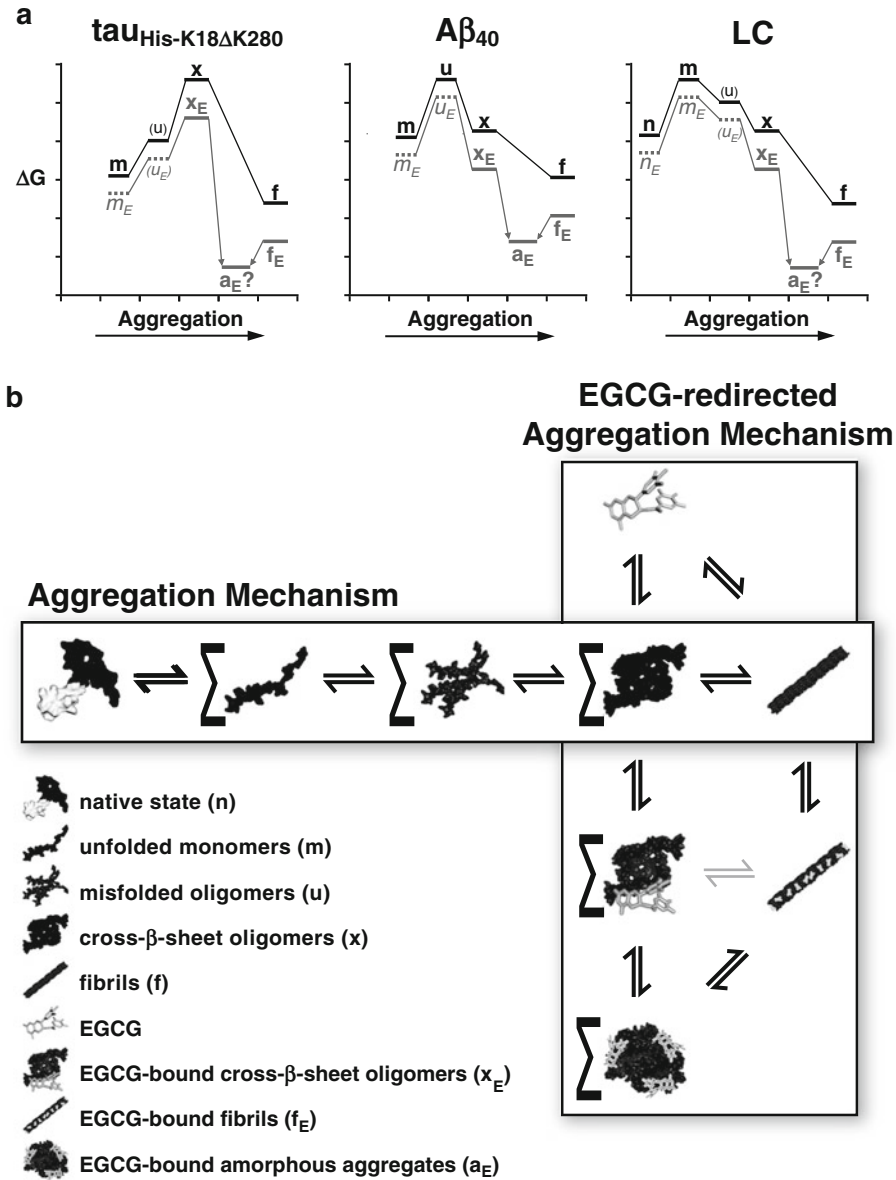


Fig. 7.3 Redirection of amyloid formation by EGCG. (a) Simplified two-dimensional energy diagrams for the aggregation in absence (black) and presence (gray) of EGCG, with n indicating the native state, m partially unfolded monomers, u misfolded oligomers, x cross- β -sheet oligomers, f fibrils and the subscript E indicating an EGCG-bound state. Dashed energy levels imply that EGCG binds only with low affinity, for bracketed intermediates it is unclear whether they exist in a non-transient manner, question marks refer to species for which the

stability in respect to the fibrillar state is not clear. (b) The formation of amyloid can be summarized in a common mechanism. Different intermediate subspecies are denoted by a summation sign. We propose that EGCG binds preferably to the cross- β -sheet oligomers and the cross- β -sheet motif on the fibril surface in the aggregation cascade. Hence, EGCG disturbs the highly ordered fibril structure and induces the formation of EGCG-bound amorphous aggregates

7.4.3 EGCG Remodels A β Fibrils Without Releasing Monomers or Oligomers

In the case of the A β peptides, it is well established that oligomeric intermediate species are cytotoxic (Walsh et al. 2002b; Haass and Selkoe 2007). We found that EGCG remodels fibrillar A β into spherical aggregates that are indistinguishable from those formed by monomeric A β in the presence of EGCG (Bieschke et al. 2010). Correspondingly, the remodeled aggregates exhibit reduced cytotoxicity. This would suggest that no toxic oligomeric species are produced during the remodeling of A β by EGCG. This hypothesis was tested by generating A β fibrils, each labeled by incorporating A β monomers that were linked either to green or red fluorophores. Mixtures of fibrils with both labels were incubated in presence of EGCG and the remodeling of fibrils was observed using fluorescence microscopy. If the remodeling process had involved the random dissociation of fibrils it would have resulted in oligomers in which red- and green-labeled monomers were statistically distributed. Instead, we observed that oligomers were clustered into either predominantly green or red, indicating that they had resulted from the remodeling of individual fibrils. Therefore, we concluded that EGCG directly remodels A β fibrils rather than disaggregating them into monomeric peptides (Bieschke et al. 2010). These findings are supported by Palhano et al. (2013), who were unable to detect by mass spectrometry A β 40 monomers during remodeling.

7.4.4 Elucidating the Binding Mode of EGCG to Amyloid Fibrils and Precursors

7.4.4.1 Investigations of the Binding Mode of EGCG

A number of possible non-covalent and covalent binding modes for EGCG to misfolded polypeptides have been postulated: non-site specific interaction with exposed hydrophobic surfaces through the hydrophobic effect (Ehrnhoefer

et al. 2008), site-specific hydrogen bonding (H-bonding) (Maiti et al. 2006), aromatic π - π -stacking (Scheraga et al. 1962), site-specific covalent binding, such as formation of disulfide-bridges via cysteines (Lambert et al. 2008) and formation of Schiff bases via primary amines (Ishii et al. 2011). EGCG binding to amyloidogenic polypeptides has been analyzed using various biophysical and biochemical methods. For instance, isothermal titration calorimetry (ITC) was used to measure overall binding energies (Wang et al. 2010, 2012b). This approach however depends on highly homogenous preparations of interaction partners, which would be challenging when measuring interaction of EGCG with oligomeric amyloid precursors. For that reason, studies have so far been limited to monomeric proteins and peptides (Wang et al. 2010; Li et al. 2013). Peptide fragments have been engineered to address the question of site-specific binding (Wang et al. 2012b).

This approach must however, be interpreted with caution, since the results are only applicable to the native full-length polypeptide if the structure of the fragments resembles the structure of the parent polypeptide or if EGCG binds independently of the secondary structure formation. While monomeric A β , as many other amyloidogenic polypeptides, is unstructured as a monomer, this is not necessarily true for oligomeric aggregation intermediates.

Single amino acids have been substituted or chemically modified in A β and IAPP to analyze site-specific interaction of EGCG with the respective amino acids. Specifically, lysines were removed to probe for Schiff base formation (Wang et al. 2010, Wang et al. 2012b, Palhano et al. 2013), phenylalanines and tyrosines for π - π -stacking (Cao and Raleigh 2012, Lopez del Amo et al. 2012) and cysteines for SH-linkage (Cao and Raleigh 2012). Of course, this approach also relies on the assumption that the substitution does not alter the structure of the polypeptide within the amyloid oligomers and fibrils.

Solution-state and, more so, solid-state NMR spectroscopy provide technically challenging, but non-disruptive methods for probing the structure of EGCG-bound polypeptides (Lopez del Amo

et al. 2012). Here, the solubility and size of the amyloid polypeptide as well as sample homogeneity and spectral resolution are limiting factors. Since all these methods have their constraints, it is necessary to combine the findings of different methods to draw a reliable picture.

7.4.4.2 EGCG is Binding to Cross- β -Sheet Motif in A β End-Stage Oligomers and Fibrils

Solid-state NMR (Lu et al. 2013; Petkova et al. 2006), X-ray crystallography (Colletier et al. 2011) as well as hydrogen-deuterium (H/D) exchange and mutational analysis (Luhrs et al. 2005) have elucidated A β fibrillar structure. Within the core of a fibril strand, the A β monomers form a hairpin structure (Fig. 7.1b). These hairpins are stacked on top of each other, in a way that peptide groups of adjacent monomers form intramolecular cross- β -sheet H-bonds while their amino acid side chains alternate towards the fibril surface and towards the fibril center. The side chains between the arms of the hairpins interdigitate to further stabilize the fibril structure (Colletier et al. 2011).

The outwardly directed N-terminal β -sheet region of A β contains an LVFF-motif with Leu17 and Phe19 orientated towards the inward C-terminal β -sheet region, while Val18 and Phe20 point out towards the fibril surface. Lysine16 could in principle provide a Schiff-base anchor directly upstream to the LVFF-motif. The N-terminal amino acids 1–14 of A β are believed to be mostly unstructured and not part of the cross- β -sheet structure (Luhrs et al. 2005, Petkova et al. 2006). The N-terminal tail of A β contains a high degree of aromatic, acidic and basic amino acids, which could interact with EGCG through π - π -stacking or H-bonding. In contrast, hydrophobic side chains are concentrated on the C-terminus of the peptide that lies on the second arm of the hairpin (Petkova et al. 2006).

The structure of EGCG-induced A β oligomers was examined using solid-state NMR spectroscopy, which also yielded insights towards the binding model of EGCG (Lopez del Amo et al. 2012). As expected, no clear structure

could be observed for the flexible N-terminal domain. In contrast, the LVFF motif was well-structured in the presence of EGCG. However, its structure was distinct from that in the absence of EGCG and the β -sheet structure of the KLVFFA region was lost in the presence of EGCG, while the β -sheet on the C-terminus of A β remained intact. Additionally the rotation of the phenyl rings (Phe19/20) as well as the rotation of adjacent His13/14 was hindered upon EGCG binding, strongly suggesting an interaction via π - π -stacking between the outwardly directed rings and the aromatic rings of EGCG. A cross-correlation for Phe19/Leu34 was found in presence and absence of EGCG, suggesting that the hydrophobic zipper stabilizing the hairpin structure may be partly intact in presence of EGCG.

Data stemming from ITC experiments conducted by Wang et al. (2010, 2012b) suggest that EGCG is forming H-bonds with the N-terminal domain of A β whilst binding to the LVFF motif via hydrophobic interactions. The authors proposed that the mode of EGCG binding gradually shifts from H-bonding to hydrophobic interaction at higher binding stoichiometries, which remains to be confirmed using complementary methods. The mean binding energies found in these ITC studies are lower than would be expected for the formation of covalent bonds, such as Schiff bases.

EGCG binding to A β 40 via Schiff base formation was specifically investigated by acetylation of the two lysine residues Lys16 and Lys28 (Palhano et al. 2013). Lysine acetylation did not alter the effect of EGCG on ThT fluorescence, seeding competence or toxicity. Acetylation of A β results in blocking of Lys16 and Lys28 and prevents the salt-bridge formation of Aps23/Lys28 that is present in A β fibrils (Luhrs et al. 2005). Lys16 and Lys28 were mutated into arginine to prevent Schiff base formation, but and at the same time exclude an effect of the salt-bridge on EGCG activity. In the same study, Superoxide Dismutase (SOD1) was used to prevent EGCG oxidation. Neither modification prevented the remodeling of A β fibrils, yet both N-acetylation and Lys-to-Arg mutations prevented the formation of SDS-resistant aggregates. Moreover, treatment with SOD1 resulted in a delayed remodeling of pre-

formed A β fibrils into SDS-stable aggregates (Palhano et al. 2013). It should be noted, however, that mutant SOD1 itself is capable of forming amyloid-like fibrils associated with amyotrophic lateral sclerosis (ALS) (Falconi et al. 2013), so that SOD1 might compete with A β for EGCG binding. Taken together, these data suggest that covalent cross-linking by EGCG does occur, but is not required for remodeling of A β by EGCG.

7.4.4.3 Observations Towards the EGCG Mechanism in IAPP, α -Syn, MSP2 and SEVI

Fewer data are available for EGCG binding to other polypeptides. In IAPP, mutational analysis revealed that site-specific interactions (π - π -stacking, Schiff base formation, disulfide bridge formation) are not necessary for EGCG binding, which implies a non-covalent, non-site-specific binding mode via hydrophobic interactions (Cao and Raleigh 2012). Similarly, when the α -synuclein protein was incubated with EGCG about 30 % of all amino acids lost their NMR-resonances, suggesting an interaction that was not site-specific but likely driven by hydrophobicity (Ehrnhoefer et al. 2008). In an NMR-study with MSP2 only weak interaction of EGCG, mostly with the N-terminal and C-terminal region of the protein could be observed (Chandrashekar et al. 2010). A pull-down experiment with EGCG-sepharose also found only weak interaction with MSP2; however, interaction might have been sterically hindered, since EGCG was coupled directly to the resin (Chandrashekar et al. 2010). In contrast, Popovych et al. (2012) found that EGCG interacted with specific residues (amino acids 246–286) of the SEVI peptide, and that prevention of Schiff base formation in SEVI prevented the inhibition of fibrillation.

In conclusion, EGCG seems to interact with polypeptides both through hydrophobic modalities and by site-specific side-chain interactions, most likely through Schiff base formation. Whether covalent interaction or hydrophobic effect drives its anti-amyloid activity seems to depend on the specific polypeptide.

7.5 Conclusion – A Mechanism for Aggregation Redirection by EGCG

A large and growing body of scientific literature demonstrates that EGCG is a potent inhibitor of amyloid formation and amyloid toxicity. Early studies on Htt, α -synuclein and A β have identified a mechanism of EGCG that is distinct from other aggregation inhibitors, namely that EGCG redirects amyloidogenic polypeptides into highly stable off-pathway aggregates. Further studies have since characterized a variety of effects of EGCG on different polypeptides. This raises the question whether EGCG acts on amyloidogenic polypeptides by a plethora of mechanisms, or whether these effects may be combined into a unified picture.

Formation of amyloid fibrils proceeds by a common mechanism starting with an amyloidogenic monomer, that can be present under physiological conditions (α -Syn, PrP, Sup35), be released from a precursor as is the case for A β and TTR. Amyloidogenicity may result from elevated protein concentrations, as for immunoglobulin light chains and β -microglobulin. Aggregation proceeds through a partially or fully unfolded monomeric state (*m*) to partially unstructured oligomer species (*o*) and then to cross- β -sheet oligomer species (*x*), which in the end progress into amyloid fibrils (*f*) (Fig. 7.3b). However, the relative stabilities of these aggregation intermediates result in aggregation kinetics with different rate-limiting steps for each polypeptide.

The variety in the details of the aggregation mechanisms may explain the apparent variety of mechanisms for the action of EGCG. The experimental data suggest that EGCG is indeed interacting with a common binding motif, but that the importance of the EGCG-binding species in the aggregation mechanism may be different depending on the polypeptide.

Data from several polypeptides indicate that EGCG binds to a cross- β -sheet structure, possibly at the same site and in competition with amyloidophilic dyes like ThT. Due to the sterical constraints in the cross- β -sheet structure, the hy-

dophobic side chains of the incorporated amino acids provide a distinct binding surface (Biancalana and Koide 2010). The fact that EGCG is able to remodel preformed fibrils of several amyloidogenic polypeptides implies its ability to bind the common cross- β -sheet structure. In our unified model, binding of EGCG to the cross- β -sheet oligomer (x) results in the species (x_E), which may then be remodeled into an amorphous aggregate (a_E) in case that this species is more stable than x_E (Fig. 7.3b).

Notably, the impact of the binding of EGCG to the amyloid cross- β -sheet motif depends on the energetic aggregation landscape of the specific amyloidogenic polypeptide (Fig. 7.3a). The shape of this landscape depends in turn on the relative stabilities of aggregation competent monomers, aggregation nuclei, oligomeric intermediates and fibrils.

Three modes of aggregation have been observed in studies involving EGCG: (1) the formation of the aggregation-competent species is rate-limiting (e.g. TTR, LC), (2) primary nucleation is rate limiting (e.g. tau), or (3) fibril propagation / secondary nucleation is rate limiting (e.g. A β , IAPP). The experimental data for these three mechanisms suggest different energetic states of the EGCG-induced aggregates (x_E) and different positions with respect to the rate-limiting step of aggregation (Fig. 7.3a).

If the formation of a cross- β -sheet nucleus is rate-limiting, then in binding to this rate-limiting species, the effect of EGCG on aggregation kinetics can be profound. The aggregation rate of tau, for instance, depends on the formation of an aggregation nucleus and it is likely that EGCG binds with strong affinity to the cross- β -sheet oligomers x . If EGCG binding remodels the nucleus into a non-amyloid aggregate, EGCG will inhibit aggregation at very low stoichiometric ratios, as is observed in the case of heparin-independent tau aggregation. In Fig. 7.2b, we see that EGCG prevents the large-scale formation of SDS-stable dimers (D), oligomers (O) and high-molecular-weight species (HMW) of tau, likely by sequestering a rate-limiting but non SDS-stable aggregation nucleus, leaving most of the protein in its monomeric state. We could not

observe fibril formation under these conditions and therefore were unable to observe if fibril remodeling occurs. Therefore, we cannot be sure whether EGCG redirects tau nuclei into aggregates that have a lower energy than the fibrils.

In contrast, if the cross- β -sheet oligomer is formed after the rate-limiting step, the effect of EGCG on aggregation kinetics is much less potent. Whether EGCG has any effect on aggregation would depend on the relative stabilities of the EGCG-induced aggregate and the amyloid fibril. If the fibril is more stable, then EGCG can still bind to the cross- β -sheet motif, reducing ThT binding, but it will have no effect on aggregation kinetics; such is the case for heparin-induced formation of tau fibrils (Wobst et al. 2015). The same may be true if formation of an aggregation competent species, for example a monomeric amyloid precursor, is rate-limiting. In this case, the rate-limiting species has no cross- β -structure and so EGCG binding does not affect aggregation kinetics. This is likely the case for TTR as well as for LC proteins.

In TTR, the rate-limiting step is the monomerization of the TTR tetramer (Hammarstrom et al. 2003). Under the experimental conditions of LC aggregation that we tested (neutral pH, mildly reducing conditions) the rate-limiting step in aggregation is the monomerization of LC dimers via reduction of disulfide bridges. Without a reduction of the disulfide bridges of light chain dimers, we did not observe aggregation over the course of one month, whereas the reduced protein formed ThT positive aggregates within hours. EGCG can then bind to this aggregate species and accelerate the formation of SDS-stable HMW-species (Fig. 7.2b).

In these cases, even if EGCG did bind to the amyloidogenic precursor, EGCG binding at substoichiometric concentrations cannot deplete its supply and is therefore not able to significantly slow down aggregation kinetics. Instead, the polyphenol seems to accelerate early aggregate formation but inhibit the slower aggregation into fibrillar species, which would be consistent with a nucleation-dependent mechanism for the second step of aggregation, although exact mechanistic details are yet to be explored.

Regarding the A β peptide, aggregation into non- β -sheet oligomers can occur rapidly, but nucleus formation is not rate limiting under most experimental conditions. Instead, secondary nucleation processes initiated by fibrillar species dominate the aggregation kinetics (Knowles et al. 2009; Cohen et al. 2013). Under these conditions, EGCG will inhibit aggregation kinetics if it can remodel fibrils into stable aggregates that are not seeding-competent, thereby removing the secondary nuclei. We found that fibril formation of A β is quantitatively redirected into SDS-stable amorphous aggregates, which are the most stable species on the aggregation pathway (Ehrnhoefer et al. 2008). Correspondingly, A β aggregation in presence of EGCG leads to formation of non-fibrillar aggregates (a_E , Fig. 7.2a). If these are more stable than the fibril, EGCG binding will also result in remodeling of EGCG bound fibrils (f_E) into amorphous aggregates (a_E). In the case of A β , EGCG bound to cross- β -sheet species partially disrupts the regular fibril structure and thus promotes the formation of aggregates with only partial β -sheet structure (Bieschke et al. 2010; Lopez del Amo et al. 2012). A similar mechanism may be applicable to IAPP, leading to the remodeling of fibrils into more stable EGCG-induced aggregates.

Our model would predict that the effect of EGCG on aggregation kinetics is lost if the fibrils are more stable than the EGCG-induced aggregates. This was indeed observed for heparin-induced tau fibril formation (Wobst et al. 2015) and may be the case for other amyloidogenic polypeptides for which EGCG would have no effect on aggregation kinetics, such as the NM4 strain of Sup35.

To conclude, the green tea polyphenol EGCG is an intriguing molecule that alters the amyloid aggregation process in novel ways. It has prompted a surprisingly large number of studies that scrutinize its activity on disease-related amyloidogenic proteins and peptides. Its pleiotropic effects on the aggregation cascade of amyloidogenic polypeptides illustrates how subtle differences in the aggregation mechanism may yield very different outcomes of drug intervention. Thus, it remains to be explored

whether the mechanistic insight provided by EGCG can be effectively translated into new therapeutic strategies.

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