# Amyloid, APP, and Electrical Activity of the Brain

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Dimitri Hefter<sup>1,2</sup>, Susann Ludewig<sup>3,4</sup>, Andreas Draguhn<sup>1,\*</sup>, and Martin Korte<sup>3,4,\*</sup>

#### Abstract

The Amyloid Precursor Protein (APP) is infamous for its proposed pivotal role in the pathogenesis of Alzheimer's disease (AD). Much research on APP focusses on potential contributions to neurodegeneration, mostly based on mouse models with altered expression or mutated forms of APP. However, cumulative evidence from recent years indicates the indispensability of APP and its metabolites for normal brain physiology. APP contributes to the regulation of synaptic transmission, plasticity, and calcium homeostasis. It plays an important role during development and it exerts neuroprotective effects. Of particular importance is the soluble secreted fragment APPs $\alpha$  which mediates many of its physiological actions, often counteracting the effects of the small APP-derived peptide A $\beta$ . Understanding the contribution of APP for normal functions of the nervous system is of high importance, both from a basic science perspective and also as a basis for generating new pathophysiological concepts and therapeutic approaches in AD. In this article, we review the physiological functions of APP and its metabolites, focusing on synaptic transmission, plasticity, calcium signaling, and neuronal network activity.

#### Keywords

APP, amyloid, Alzheimer's disease, synaptic transmission, plasticity, oscillations

### Introduction

The Amyloid Precursor Protein (APP) is a ubiquitously expressed transmembrane protein with a long extracellular and a short intracellular domain. The predominant isoform in the central nervous system (CNS) consists of 695 amino acids (aa; Gralle and Ferreira 2007; Müller and Zheng 2012; Müller and others 2017). APP forms a protein family together with the homologous APP-like proteins 1 and 2 (APLP1 and APLP2, respectively). APP is highly conserved throughout the animal kingdom, beginning in nematodes, emphasizing its indispensable role for neuronal function. Indeed, triple knock-out (KO) mice lacking all three proteins of the APP/APLP family, as well as APP/APLP2 double knock-outs (DKO) are not viable. They show severe cortical malformations (Herms and others 2004), demonstrating a crucial role of APP/APLP during development. In fact, APP has been shown to promote synapse formation, dendritic sprouting, and neuronal migration (Müller and others 2017). Single APP-KO mice are viable due to compensation by the homologues, but show deficits in long-term potentiation (LTP) at old age (Dawson and others 1999; Ring and others 2007), learning and memory formation as well as higher susceptibility to seizures and hypoxia-ischemia (Hefter and others 2016; Koike and others 2012; Steinbach and others 1998). These findings point to several physiological functions of APP which are addressed in more detail below.

APP is cleaved by the  $\alpha$ - or  $\beta$ -secretase (also known as beta-site amyloid precursor protein cleaving enzyme [BACE]) and consecutively by the  $\gamma$ -secretase (Haass and others 2012; Fig. 1A-C). Recently, a third, novel,  $\eta$ -secretase pathway with yet unclear function has been

<sup>1</sup>Institute of Physiology and Pathophysiology, Heidelberg University, Heidelberg, Germany

 <sup>2</sup>RG Animal Models in Psychiatry, Department of Psychiatry and Psychotherapy, Central Institute of Mental Health, Medical Faculty Mannheim, University of Heidelberg, Heidelberg, Germany
<sup>3</sup>Helmholtz Centre for Infection Research, Neuroinflammation and Neurodegeneration Group, Braunschweig, Germany
<sup>4</sup>Cellular Neurobiology, Zoological Institute, Technical University Braunschweig, Braunschweig, Germany

\*Equal contribution.

#### **Corresponding Authors:**

Andreas Draguhn, Institute of Physiology and Pathophysiology, Heidelberg University, Im Neuenheimer Feld 326, Heidelberg 69120, Germany.

Email: andreas.draguhn@physiologie.uni-heidelberg.de

Martin Korte, Neuroinflammation and Neurodegeneration Group, Helmholtz Centre for Infection Research, Inhoffenstraße 7, Braunschweig, Niedersachsen 38124, Germany. Email: m.korte@tu-braunschweig.de



Figure 1. Proteolytic processing of APP by the secretases. (A) Schematic structure of the membrane-bound, full-length Amyloid Precursor Protein (APP). On the left side the shorter C-terminal intracellular domain is depicted in green, on the right side the longer N-terminal extracellular domain is in blue. The amyloid sequence is shown in red. Arrowheads point to the secretase cleavage sites. The length of the fragments is not proportional to the length of the respective amino acid sequence. (B) Cleavage by  $\alpha$ -secretase and consecutively by  $\gamma$ -secretase. CTF $\alpha$  = C-terminal fragment alpha; APPs $\alpha$  = extracellularly secreted soluble APP alpha fragment; AICD = APP intracellular domain. (C) Cleavage by  $\beta$ -secretase and consecutively by  $\gamma$ -secretase. CTF $\beta$  = C-terminal fragment beta;  $APPs\beta = extracellularly secreted$ soluble APP beta fragment;  $A\beta$  = amyloid beta. (D) Cleavage by  $\eta$ -secretase and consecutively by  $\alpha$ ,  $\beta$ , and  $\gamma$ -secretases.  $\text{CTF}\eta=\text{C-terminal}$  fragment eta;  $\text{APPs}\eta=\text{extracellularly}$ secreted soluble APP eta fragment; A $\eta$ - $\alpha$ , A $\eta$ - $\beta$  = amyloid eta-alpha and eta-beta, respectively.

discovered (Willem and others 2015; Fig. 1D). Cleavage by the  $\alpha$ -secretase initiates the so-called non-amyloidogenic pathway, which results in the APP-intracellular domain (AICD) and the soluble extracellularly secreted APPs $\alpha$ fragment. APPs $\alpha$  was shown to mediate most of the known neuroprotective and neurotrophic effects of APP (Hefter and others 2016; Mockett and others 2017). In contrast,  $\beta$ secretase cleavage is the starting point for the amyloidogenic pathway. Besides, the intracellular fragment AICD it produces a secreted APPsß fragment and, importantly, amyloid  $\beta$ , a small peptide varying from 38 to 43 aa in length. AB exists in different monomeric or multimeric soluble forms and can aggregate to fibrils and plaques. Such aggregates are most easily formed by AB42, which is less common than the AB40 isoform and is more prone to precipitate. The extracellular accumulation of amyloid plaques, along with the intracellular deposition of tau fibrils, is the histopathological hallmark of Alzheimer's disease (AD) and has been replicated in various mouse models of the disease (Sasaguri and others 2017). Amyloid deposition may be one of the initial steps of the pathophysiological cascade of AD as it usually precedes tau pathology as well as pathophysiological alterations and clinical symptoms. Thus, it is widely being considered crucial for AD pathogenesis, and enormous efforts have been put into development of strategies targeting amyloid (Selkoe and others 2016). However, the correlation between amyloid burden and clinical symptoms is quite weak. In line with this fact, clinical trials based on anti-amyloid treatment strategies for AD received heavy setbacks in recent years (van Dyck 2018). This therapeutic failure makes it even more important to understand the normal functions of APP family proteins and their metabolites in animal models and in humans.

APP metabolites affect synaptic transmission and network function in health and disease, partially counteracting each other. Of special interest are neuroprotective and neurotrophic functions of APP—or its fragment APPs $\alpha$  which may be essential for normal neuronal function and for resilience toward neurodegenerative diseases. In the following sections we give an overview of physiological functions of APP at the synaptic and network levels.

# **APP and Synaptic Transmission**

APP and its metabolites affect both excitatory and inhibitory synaptic transmission in health and disease. At the postsynaptic site APP interacts with glutamatergic (Hoe and others 2009) and cholinergic (Richter and others 2018) receptors. At the presynaptic site it regulates the function of GABA<sub>B</sub> receptors (GABA<sub>B</sub>-R; Dinamarca and others 2019; Fig. 2B), synaptic vesicle release (Klevanski and others 2015), and the abundance of presynaptic proteins (see section "Short-Term Plasticity" for more details; Fig. 2A). AB oligomers disrupt synaptic activity and shift the excitation/inhibition ratio toward excitation, altering paired-pulse plasticity and abolishing LTP (Puzzo and others 2008). Application of synthetic AB on cultured hippocampal neurons leads to downregulation of dynamin, a protein that is essential for synaptic vesicle endocytosis and interrupts synaptic vesicle recycling (Kelly and others 2005; Kelly and Ferreira 2007). Various direct and indirect interactions with receptors



**Figure 2.** Synaptic actions of APP and its metabolites. (A) Putative and conformed actions of APP and AICD on presynaptic vesicle release. (B) Putative actions of APP dimers at the synapse. APP form dimers with a second APP molecule in the trans-(1) or cis-formation (2). Cis-dimeric APP may act as a receptor for APPs $\alpha$  and A $\beta$ . (3) APP colocalizes in transport vesicles for GABA<sub>B</sub>-R and is part of the GABA<sub>B</sub>-R complex at the presynaptic site where it decreases the releasable vesicle pool and increases paired pulse facilitation. (C) Interactions between APP and cholinergic receptors. APP, APPs $\alpha$ , AICD, and picomolar concentrations of A $\beta$ . Present evidence indicates a positive modulation of cholinergic transmission via interactions with  $\alpha$ 7-nAchR, whereas nanomolar and higher concentrations of A $\beta$  block these receptors. M1-AchR activation promotes cleavage of APP by  $\alpha$ -secretase and APPs $\alpha$  secretion; nAchR promotes amyloidogenic processing. Ach = acetylcholine; M1-AchR = M1-type muscarinic acetylcholine receptors; nAchR = nicotinic acetylcholine receptors; Gq = G-Protein q. (D) Interactions between APP and NMDAR. Left: synaptic NMDAR signaling, predominantly GluN2A. Right: extrasynaptic NMDAR signaling, predominantly GluN2B. A $\beta$  inhibits GluN2A-NMDAR-mediated survival pathways. A $\beta$  inhibits GluN2A-NMDAR signaling, Full-length APP positively modulates membrane trafficking and AICD increases expression of GluN2B-NMDAR. GluN2B-NMDAR signaling reduces abundance of APP at the membrane and promotes A $\beta$  secretion. Glu = glutamate.

have been proposed to mediate the toxic activity of  $A\beta$ (Strooper and Karran 2016). APP and APLP molecules can form cis- and trans-dimers. Cis-homodimerized APP has been proposed to function as a cell-surface G-protein coupled receptor with A $\beta$  and APPs $\alpha$  as putative ligands (Ludewig and Korte 2017). Thus, a recent study showed that synaptic effects of amyloid require the presence of full-length APP (Wang and others 2017). In their transforms, transcellular dimers of APP family proteins mediate cell-to-cell adhesion and support synaptic connectivity (Schilling and others 2017). A $\eta$ - $\alpha$ , one of the products of the recently discovered  $\eta$ -secretase pathway, was shown to inhibit neuronal activity and LTP in hippocampal slices (Willem and others 2015). While APP and its fragments affect synaptic activity in multiple ways, APP cleavage and amyloid levels are themselves dynamically regulated by neuronal activity (Bero and others 2011; Cirrito and others 2005). Intriguingly, even the relative abundance of amyloid isoforms is regulated by patterns of neuronal firing. Thus, firing of single spikes was shown to favor the formation of the supposedly more toxic AB42 peptide, while burst firing facilitated the formation of  $A\beta 40$ . These effects are mediated by an activity-dependent conformational change of presenilin1, the catalytic partner of  $\gamma$ -secretase (Dolev and others 2013). In the following section we will focus on the interplay of APP and three major neurotransmitter systems which are known to be altered in AD.

# Glutamatergic Transmission and NMDA Receptor Signaling

Ionotropic glutamate-receptors comprise α-amino-3hydroxy-5-methyl-4-isoxazole-propionic acid receptors (AMPAR) and N-methyl-D-aspartate receptors (NMDAR). The characteristic voltage-dependent gating of NMDAR plays a pivotal role in synaptic plasticity, learning, and memory as well as in pathophysiological cascades in AD and hypoxia-ischemia (Paoletti and others 2013). The NMDAR family comprises a heterogeneous group of tetramers consisting of a combination of the obligatory GluN1 subunit and the facultative GluN2-A, B, C, or D or GluN3 subunits. During development, the expression pattern in the forebrain shifts from the predominant GluN2B to GluN2A subunits (Monyer and others 1994). In the adult brain, GluN2B is commonly associated with excitotoxicity and neuronal degeneration while GluN2A is linked to plasticity and pro-survival pathways (Bading 2017; Hardingham and Bading 2010). Chronic overactivation of NMDAR has been hypothesized as a pathological mechanism in AD, in line with the use of the noncompetitive NMDAR blocker memantine as one of the few approved drugs for treatment of AD (Liu and others 2019). APP as well as APLP1 and APLP2 were found to affect NMDAR trafficking and to enhance its surface expression (Cousins and others 2009; Cousins and others 2015; Fig. 2D). The effects on expression are mutual: APP increases membrane levels and function of GluN2B-containing NMDAR while NMDAR activation decreases surface expression of APP and promotes amyloidogenesis (Hoe and others 2009). Furthermore, APP was found to regulate the homeostasis of D-serine, a potent endogenous co-agonist of NMDAR. APP-deficient transgenic mice show D-serine-dependent impairments in structural plasticity of dendritic spines (Zou and others 2016).

Several studies using intrathecal injection of exogenous APPs $\alpha$  in mice suggest that APPs $\alpha$ , but not APPs $\beta$ , potently regulates NMDAR function. Indeed, suppression of NMDA currents in hippocampal neurons by APPs $\alpha$ occurs already at picomolar concentrations (Furukawa and Mattson 1998). Another study did not find effects of APPs $\alpha$  on baseline NMDAR currents, but showed its requirement for the induction of LTP (Taylor and others 2008). The effects on NMDAR function appear to be agedependent. In healthy aged rats APPs $\alpha$  levels were found to be reduced alongside with NMDAR function and LTP. In these animals, exogenous application of APPs $\alpha$  did potentiate LTP while it had no such effect in younger animals (Moreno and others 2015).

The intracellular fragment AICD was shown to promote expression of GluN2B-containing NMDAR and to facilitate GluN2B-mediated synaptic transmission. While APPknockdown attenuates GluN2B currents, increased AICD production enhances NMDAR function and disrupts LTP (Pousinha and others 2017). On the other hand, Aβ oligomers inhibit NMDAR-activity-dependent trophic cascades leading to synapse loss in the hippocampus (Shankar and others 2007). Vice versa, several studies suggest modulation of APP processing by NMDAR signaling. Activation of synaptic NMDAR stimulates  $\alpha$ -secretase processing and inhibits AB production (Hoey and others 2009) while activation of extrasynaptic NMDAR promotes amyloidogenesis (Rush and Buisson 2014). In another study, pharmacological block of NMDAR prevented toxic effects of A $\beta$  on dendritic spines (Wei and others 2009). Taken together, an increasing body of evidence indicates that activation of synaptic GluN2A-NMDAR promotes plasticity and survival via facilitation of APPsa production. In contrast, extrasynaptic GluN2B-NMDAR promote cell death by  $\beta$ -secretase activation (Rush and Buisson 2014).

AMPAR are the principal mediators of excitatory glutamatergic transmission in the CNS. Their dysregulation by A $\beta$  is one of the hallmarks of synaptic failure in AD (Paula-Lima and others 2013). Reduced expression and function of AMPAR were found in neurons that overexpress wildtype APP or the APP Swedish double mutation as well as in neurons following exogenous application of (Almeida and others 2005; Chang and others 2006; Hsieh and others 2006). One of the major underlying mechanisms is disrupted AMPAR trafficking mediated by NMDA and metabotropic glutamate receptor signaling (Guntupalli and others 2016). A study on mice overexpressing A $\beta$  showed that its detrimental effect on synapses requires the presence of the AMPAR subunit GluA3 (Reinders and others 2016). On the other hand, AMPAR activation was shown to increase the clearance of A $\beta$  in a dose-dependent manner, thus decreasing its extracellular levels (Hettinger and others 2018). Simultaneously, it promotes non-amyloidogenic cleavage of APP (Hoey and others 2013). These findings demonstrate a complex, reciprocal relationship between APP metabolites and glutamate receptors that depends on receptor composition, location and developmental stage.

#### GABAergic Transmission

GABA receptors are divided into three main types: the ionotropic GABA<sub>A</sub> and GABA<sub>C</sub> (GABA<sub>A</sub>- $\rho$ ) and the metabotropic GABA<sub>B</sub>-R, all of which are composed from a variety of distinct subunits (Chebib and Johnston 1999). Disrupted GABAergic synaptic transmission, interneuron dysfunction and consecutive aberrant network activity are implicated in the pathophysiology of AD (Verret and others 2012). APP was shown to modulate paired-pulse inhibition and tetanic potentiation in striatal and hippocampal GABAergic interneurons via inhibition of L-type calcium channels (LTCC; Yang and others 2009). Interestingly, APP is highly expressed in GABAergic interneurons, regulating both phasic and tonic inhibitory function. Its selective deletion in GABAergic, but not glutamatergic, neurons disrupts adult hippocampal neurogenesis (Wang and others 2014). APP and APLP2 were found to be part of the GABA<sub>B</sub>-R complex (Schwenk and others 2016), suggesting a role in presynaptic release dynamics or plasticity (Fig. 2B). Indeed, evidence from two very recent studies supports this hypothesis: Binding of APP to GABA<sub>B</sub>-R was shown to promote its axonal trafficking, thus facilitating cell-surface expression of these receptors and presynaptic inhibition. At the same time, GABA<sub>B</sub>bound APP was less likely to be cleaved by the amyloidogenic  $\beta$ -secretase (Dinamarca and others 2019). These findings link dysfunctional GABA<sub>B</sub>-R trafficking and deficits in presynaptic inhibition with increased amyloid cleavage in AD. Furthermore, all secreted APP fragments  $(\alpha, \beta, and \eta)$  were shown act as a GABA<sub>B</sub>-R ligands. APPsa binding to the sushi domain of GABA<sub>B</sub>-R inhibited presynaptic vesicle release, thus suppressing synaptic transmission and enhancing short-term facilitation in the hippocampus (Rice and others 2019). Intriguingly, APPsβ binding exerted similar effects. Thus, this study is one of very few to find a physiological role of APPsB. However, it must be taken into consideration that APP proteins bind to GABA<sub>B</sub>-R with relatively low affinity (dissociation

constants  $K_D$  of ~200 and ~400 nM, respectively), and 1  $\mu$ M concentrations have been used in this study, while the physiologically active concentrations of these proteins are in the picomolar to low nanomolar range (Hick and others 2015; Puzzo and others 2008). Therefore, APP's effects under physiological circumstances rather arise from other interactions than with GABA<sub>B</sub>-R, while functional implications of these findings require further investigation. Nevertheless, these findings link APP and GABAergic signaling both in health and disease and present an additional regulatory mechanism of presynaptic excitability by APP.

## Cholinergic Transmission

Dysfunction of the cholinergic system is strongly implied in the pathogenesis of AD. Several clinically approved antidementives inhibit the acetylcholine-cleaving enzyme acetylcholinesterase, aiming to facilitate cholinergic transmission (Knight and others 2018). Studies in APP-deficient mice show that APP is important for peripheral cholinergic transmission at the neuromuscular junction (Caldwell and others 2013; Weyer and others 2011) as well in the autonomous nervous system (Cai and others 2016; Fig. 2C). Transgenic mouse models of AD confirm that APP overexpression and aberrant amyloid deposition are associated with degeneration of cholinergic neurons beginning with damage of their axons in the cortex (Foidl and others 2016). In support of these findings, inhibition of  $\beta$ -secretase rescued cholinergic dysfunction in a mouse model of AD (Ohno and others 2004). Of special importance for cholinergic function is the intracellular AICD peptide. Mice with knock-in of a specific, dysfunctional motif within AICD show impaired muscular and cognitive performance (Matrone and others 2012). APPs $\alpha$  has been suggested to function as an endogenous positive allosteric modulator of cholinergic signaling by binding to  $\alpha$ 7-type nicotinic acetylcholine receptors ( $\alpha$ 7nAChRs; Richter and others 2018). The aa sequence required for associating APPsa with a7-nAChRs is also present in A $\beta$ . Therefore, it is not surprising that A $\beta$  also potently binds to these receptors, both on presynaptic and postsynaptic sites. The resulting modulation of cholinergic signaling depends on its concentration and the conformation of nAchR (Lasala and others 2019). While physiological concentrations of A $\beta$  in the picomolar to low nanomolar range were shown to activate nAchR and increase synaptic vesicle recycling, higher concentrations inhibit cholinergic currents (Lazarevic and others 2017). This may be of great importance for AD, where levels of AB are increased. Similar to GABAergic and glutamatergic transmission, several studies have shown that cholinergic signaling modulates APP processing regulating interstitial amyloid levels. Thus, activation of muscarinic (M1) receptors was shown to inhibit BACE and facilitate non-amyloidogenic cleavage

and APPs $\alpha$  secretion (Jiang and others 2014) while activation of nicotinic receptors was found to increase interstitial A $\beta$  levels (Wei and others 2009).

## **Calcium Signaling**

Neuronal activity is accompanied by fluctuations of intracellular calcium concentration. Calcium modulates a plethora of enzymatic cascades, gene transcription, as well as pro-survival and apoptosis pathways; thus, calcium signaling is a major activity-dependent factor affecting the intermediate- and long-term cellular fate (Bading 2013). Furthermore, neuronal excitability and thus network function are heavily influenced by calcium concentrations. Dysregulation of the intracellular calcium homeostasis is considered a common pathomechanism in various neurological conditions, prominently AD and stroke, where cell death and degeneration are involved (Alzheimer's Association Calcium Hypothesis Workgroup 2017; Green and LaFerla 2008). Cytosolic calcium can derive from extracellular sources such as NMDAR or voltage-gated calcium channels (VGCC) as well as from intracellular stores such as mitochondria and the endoplasmic reticulum (ER; Clapham 2007; Lerdkrai and others 2018). APP and its metabolites regulate calcium homeostasis at multiple levels both in trophic, plasticityrelated pathways (via APPs $\alpha$ ) and in a destabilizing way in AD (via A $\beta$ ; Fig. 3). The interactions between APP and NMDAR have been highlighted above; in the following section we focus on VGCC and internal calcium stores.

## Voltage-Gated Calcium Channels

VGCC are a family of voltage-gated ion channels with selective permeability for Ca<sup>2+</sup>. Excessive calcium influx through VGCC is suggested to contribute to calcium overload in degenerative and neurovascular diseases (Cataldi 2013), while VGCC antagonists, on the other hand, show protective effects against cognitive decline (Lovell and others 2015). In several studies amyloid oligomers were shown to activate one subtype of VGCC, Cav1.2 subunit-expressing L-type calcium channels (LTCC). The expression of LTCC is driven by  $A\beta$  under hypoxic conditions (Webster and others 2006). Their blockage has been identified as a potential pharmacotherapeutic target for AD (Anekonda and Quinn 2011; Lovell and others 2015). APP regulates LTCC in cultured GABAergic neurons of the hippocampus and striatum, with APP deletion leading to aberrant activity of LTCC and altered short-term plasticity (STP; Yang and others 2009). As expression of APPsα on an APP-KO background was sufficient to rescue deficits linked to enhanced calcium influx through LTCC, these effects are likely mediated by the APPs $\alpha$  fragment (Hefter and others



Figure 3. APP and calcium homeostasis. Calcium can enter the cytosol from extracellular space via NMDAR and voltagegated calcium channels including L-type calcium channels (LTCC) or from intracellular stores such as mitochondria and the endoplasmic reticulum via several calcium transporters. Intracellular calcium triggers various signaling cascades, translocates to the nucleus, and affects gene transcription. APP and APPs $\alpha$  inhibit LTCC. AICD is important for the control of calcium homeostasis by regulating transport to internal stores and transcription of calcium-binding proteins. A $\beta$  oligomers are present in mitochondria where they destabilize mitochondrial membrane potential and function, disturb adenosine triphosphate (ATP) production, and increase reactive oxygen species (ROS) levels.

2016). Expression of human APP in rat primary cortical neurons lead to increased LTCC currents and inhibited calcium oscillations (Santos and others 2009). On the other hand, A $\beta$  appears to inhibit P/Q-type VGCC (Cataldi 2013). The mechanisms of interaction between amyloid and VGCC remain elusive—suggestions include channel phosphorylation by MAP-kinases, free radical formation, and effects on channel trafficking.

#### Internal Calcium Stores

It has been suggested that intracellular calcium stores play a major role as sources for dysregulation of neuronal calcium levels in aging and AD (Thibault and others 2007). Dysfunction of presynaptic calcium stores leads to neuronal hyperactivity and likely more frequent NMDAR activation in an in vivo mouse model of AD (Lerdkrai and others 2018). A $\beta$  has been suggested to disrupt mitochondrial calcium homeostasis and thus lead to perturbations of cerebral

glucose and cholesterol metabolism as well as to accumulation of reactive oxygen species (Barbero-Camps and others 2014; Chen and Yan 2006; Chen and Zhong 2013). In support of this hypothesis, A $\beta$  oligomers were found to promote cell death via endoplasmic reticulum stress, endosomal/lysosomal leakage, and mitochondrial dysfunction (Umeda and others 2011). In contrast to these toxic effects of A $\beta$ , AICD is implicated in regulation of intracellular calcium and energy homeostasis. Possible mechanisms are modulation of calcium transporters between the cytoplasm and intracellular calcium stores (Hamid and others 2007) and transcription of calcium-binding proteins (Cao and Sudhof 2001). Thus, cells lacking AICD exhibit increased calcium levels, hyperpolarized mitochondria, and reduced ATP content (Hamid and others 2007).

In a recent study full-length APP was found to play a critical role in mitochondrial calcium homeostasis in astrocytes. In vivo  $Ca^{2+}$  imaging of astrocytic microdomains revealed a critical role of the amyloid precursor protein for mitochondria (Montagna and others 2019). As astrocytes potently modulate neuronal function by regulating the local supply of ions and energy metabolites, these findings point toward a novel, APP-related mechanism modulating brain function via glial cells.

## **APP and Synaptic Plasticity**

Since many interaction partners of APP mentioned above such as NMDAR and calcium channels are major mediators of short- and long-term synaptic plasticity, it is not surprising that APP intervenes with plasticity-related pathways in various ways (Ludewig and Korte 2017). As synaptic plasticity is the physiological foundation of learning and memory, deficits of which constitute the core symptoms of AD, analysis of APP functions may reveal leverage points to improve these deficits. Therefore, in this section we will highlight the effects of APP on short- and long-term synaptic plasticity, structural plasticity, learning, and memory formation (Fig. 4).

#### Long-Term Potentiation and Depression

LTP and LTD (long-term depression) are plasticity processes over the time scale of several hours which are mediated by NMDAR and the subsequent intracellular calcium cascades. LTP deficits are a prominent feature of transgenic AD mouse models (Pozueta and others 2013). On the other hand, LTP is also abolished in APP-deficient transgenic mice (Dawson and others 1999; Seabrook and others 1999). Aged APP-KO mice (9 to 12 months) show impaired learning behavior and LTP (Ring and others 2007; Tyan and others 2012). The age-dependent alterations in synaptic plasticity and other biological functions indicate that APLP2—and, possibly, APLP1—may



Figure 4. APP and synaptic plasticity. Neuronal plasticity takes place at different time scales: from short-term plasticity (STP) in the order of milliseconds to hundreds of milliseconds to long-term plasticity in the order of hours, days, and years. These functional changes induce structural plasticity such as formation of synapses and dendritic spines. APP and picomolar concentrations of AB facilitate STP by interaction with the presynaptic vesicle release machinery, GABAergic and cholinergic receptors, and calcium channels, whereas higher, pathological concentrations of  $A\beta$  inhibit STP and long-term potentiation (LTP). APP is necessary for LTP, which is mediated by APPs  $\!\alpha$  and AICD. Physiological concentrations of A $\beta$  promote long-term depression (LTD). APP and APPs $\alpha$  exert trophic effects on spine formation and dendritic arborization promoting structural plasticity, while A $\beta$  oligomers as in AD act toxic and lead to reduced numbers of spines and dendritic branches.

perform redundant functions within the protein family. Interestingly, adult and aged APLP2-KO mice behave like wildtype mice (Midthune and others 2012; Weyer and others 2011), while aged APLP1-KOs have impaired basal synaptic transmission and reduced miniature excitatorypostsynaptic currents frequency, while LTP is normal (Schilling and others 2017; Vnencak and others 2015). It seems likely that endogenous APP is able to compensate for the genetic ablation of APLP2 with age, while vice versa APLP2 is incapable to compensate the loss of APP in aged animals. Furthermore, APLP1 is an essential player in the maintenance of synaptic transmission.

Several studies suggest that the APP intracellular domain (AICD) plays essential roles at synapses.

APP $\Delta$ CT15-DM mice lacking the last 15 aa of APP including the highly conserved YENPTY motif on an APLP2 deficient background show impaired induction and maintenance of LTP. Impaired synaptic plasticity was accompanied by severely altered hippocampusdependent behavior. Acute hippocampal slices of this mouse strain showed altered postsynaptic properties and a trend toward defective protein synthesis-dependent late-LTP, presumably caused by abolished YENPTY interactions (Klevanski and others 2014). Further known APP interaction partners comprise Dab1, Shc, Grb, and Mint/X11 proteins. Mint/X11 proteins promote clathrin-mediated endocytosis of APP and are involved in the transport of APP to the cellsurface (Aydin and others 2012; van der Kant and Goldstein 2015). Of particular importance might be the interaction with the adapter protein family Fe65. Fe65/ Fe65L1 double-deficient mice show a phenotype of cortical dysplasia similar to APP triple-KO animals (Guenette and others 2006). Co-localization of Fe65 with APP in the endoplasmic reticulum (ER) and Golgi apparatus alters the translocation of APP to the cell surface (Sabo and others 1999). As APP family members are enriched in intracellular compartments under resting conditions, this co-localization might be coupled to activity-dependent synaptic plasticity and thus be important for synaptic function (Thinakaran and Koo 2008). The complex of AICD and Fe65 is known to bind Tip60, building a transcription factor and thus regulating gene transcription after shuttling into the nucleus (Cao and Sudhof 2001). This process is essential for protein synthesis-dependent L-LTP. Recently, a study showed that Fe65 proteins bind to all APP family members and might act downstream of cell-surface APP/APLPs at central synapses (Strecker and others 2016). Moreover, a single aa difference between the intracellular domains (ICD) of APP and APLP2 enabled induction of LTD via the AICD, but not the APLP2-ICD (Trillaud-Doppia and others 2016). More recently, the same group reported a role of AICD in metaplasticity (Trillaud-Doppia and Boehm 2018). In this study, neurons expressing full-length chimeric APP (the  $A\beta$ domain was exchanged for its homologous domain from APLP2) showed facilitated LTD and completely blocked LTP induction while basic synaptic transmission and plasticity were unaltered. These effects within a very short time frame suggest that AICD might also act independent of gene transcription.

Several studies highlight the LTP-facilitating function of APPs $\alpha$  in chicks and mice (Ishida and others 1997; Meziane and others 1998; Mileusnic and others 2004). Expression of the APPs $\alpha$  fragment on an APPknockout background (Ring and others 2007) or an

APP/APLP2-knockout background (Weyer and others 2014; Zhang and others 2013) was sufficient to rescue LTP deficits. Even acute application of APPs $\alpha$  (10 nM), but not APPsB (50 nM), was sufficient to restore impairments in LTP in conditional APP/APLP2 DKO mice (Hick and others 2015). Moreover, in contrast to APPs $\alpha$ the knock-in of APPs $\beta$  in mice was unable to rescue the perinatal lethality seen for APP and APLP2 DKO mice (Li and others 2010; Li and others 2011; Weyer and others 2011), confirming functional differences between these two large extracellular domains. Several approaches have been used to reduce the APPs $\alpha$  content, for example, conditional KO of the major  $\alpha$ -secretase ADAM-10 (Prox and others 2013) or application of conventional  $\alpha$ secretase inhibitors in vitro and in vivo (Taylor and others 2008). These manipulations resulted in strongly impaired LTP. Recently, it was shown that viral-driven expression of APPsa rescues synaptic plasticity in conditional APP/ APLP2 DKO mice (Richter and others 2018; Tan and others 2018). The same approach has restored impaired LTP in the APP/PS1 tg mouse model (Fol and others 2016). In contrast, the modulatory role of APPs $\beta$  for synaptic plasticity is less clear (Hick and others 2015; Richter and others 2018; Taylor and others 2008). Of course, the effects of AB peptide accumulation and plaque formation during AD progression are relevant and intensely studied (Zott and others 2018). While this disease-related focus points to a negative modulatory function on synaptic plasticity, several studies indicate that picomolar concentrations of monomeric AB play an important physiological role. By modulating presynaptic vesicle release, picomolar amounts of A $\beta$  have been proven to be essential for processes of synaptic plasticity like LTP and STP (Abramov and others 2009; Wang and others 2012) via interaction with nicotinic acetylcholine receptors (Lawrence and others 2014). LTP is enhanced following exogenous administration of 200 pM AB42 (Puzzo and others 2008). Furthermore, A $\beta$ 42 alters the counterpart of LTP, LTD. In vivo injection of 1 to 2 pmol A $\beta$ 42 was shown to facilitate LTD as well as reversal of LTD in the CA1 region of the hippocampus (Kim and others 2001). The underlying mechanism is likely related to NMDAR and AMPAR. Synthetic AB42 peptides as well as naturally secreted AB from APP/PS1 transgenic mice promote the endocytosis of NMDAR from the postsynaptic surface, depressing the NMDAR current in wild-type cortical neurons (Snyder and others 2005). Furthermore, AB induced synaptic depression involves enhanced loss of AMPA receptors from dendritic spine surfaces via endocytosis. Interestingly, this is mediated by signaling pathways similar to that occurring during LTD, involving calcineurin and p38 MAP kinase activity (Chen and others 2013; Zhao and others 2004).

Recently an in vivo study on the  $\beta$ -secretase BACE-1 confirmed the essential role of proteolytic peptides generated within the amyloidogenic pathway. Loss of BACE-1 resulted in increased granule cell excitability and prolonged paired-pulse-inhibition as well as altered network gamma oscillations and impaired synaptic plasticity (Vnencak and others 2019).

In contrast, nanomolar concentrations of A $\beta$ 42 have deleterious, neurotoxic effects. Especially a ~7 kDa A $\beta$ extract from human AD brains seems to be particularly toxic (Brinkmalm and others 2019). Application of nanomolar amounts of synthetic A $\beta$ 42, A $\beta$ 40, or A $\beta$ 25-35 (200 nM to 1  $\mu$ M) in acute slices inhibits LTP induction in the hippocampus (Chen and others 2000; Wang and others 2017; Zhao and others 2004). One of the underlying mechanisms is inhibition of AMPAR phosphorylation (Zhao and others 2004). Moreover it has been demonstrated that the intraneuronal injection of 1 to 1000 nM A $\beta_{42}$  (Ripoli and others 2014) as well as cerebral microinjection of heterogeneous human A $\beta$  (~3 pg) in rats in vivo (Walsh and others 2002) depressed basal synaptic transmission and LTP.

Interestingly, Wang and others (2017) showed that Aβ-mediated impairments of synaptic plasticity depend on the expression of APP. The synaptotoxic effects of A $\beta$  go along with presynaptic alterations disturbing the excitation/inhibition (E/I) balance. The recently discovered peptides An- $\alpha$  and An- $\beta$ , both generated via the  $\eta$ -secretase pathway, have been shown to have opposing functions on LTP, similar to the APPsa and AB counterbalance (Willem and others 2015). While both peptides had no influence on baseline synaptic transmission, hippocampal LTP was severely impaired by An- $\alpha$  but not by A $\eta$ - $\beta$ . Willem and others (2015) used either A $\eta$ - $\alpha$ conditioned medium or 100 nM synthetic A $\eta$ - $\alpha$  and observed a reduction of LTP. Interestingly, a concentration-dependent action of murine or human recombinant APPs $\alpha$  has been shown (Hick and others 2015; Kuchibhotla and others 2008; Richter and others 2018; Tan and others 2018; Taylor and others 2008). All studies highlight a facilitating action of recombinant APPs $\alpha$  in a range of 1 to 11 nM on LTP, while for An- $\alpha$  no other concentrations have been tested so far. Moreover, the application of higher APPsa amounts had no effect or resulted even in reduced LTP when reaching 3300 nM (Taylor and others 2008). Interestingly, by analogy to APPs $\alpha$  and APPs $\beta$ , A $\eta$ - $\alpha$  and A $\eta$ - $\beta$  differ in their sequence by 16 C-terminal aa, which are only present in APPs $\alpha$  and A $\eta$ - $\alpha$ . This peptide sequence contains a predicted neuroprotective domain (Furukawa and others 1996) and was recently shown to be sufficient to facilitate LTP in a manner similar to APPsa (Richter and others 2018).

#### Short-Term Plasticity

STP takes place on a timescale of milliseconds to hundreds of milliseconds and is dependent on presynaptic factors such as depletion of the transmitter vesicle pool as well as postsynaptic factors such as feed-forward inhibition. APP was found to affect STP on GABAergic interneurons (Yang and others 2009). These authors reported increased L-type voltage gated Ca2+ channel (LTCC) activity, and Hefter and others (2016) showed that APP, possibly via the APPsα fragment, stabilizes Ca<sup>2+</sup> homeostasis by regulating the inhibition of LTCCs. Besides APP, its homologue APLP1 is also implicated in GABAergic transmission. APLP1-KO and aged APP-KO mice exhibit reduced GABAergic mediated paired-pulse depression (PPD; Seabrook and others 1999; Vnencak and others 2015) suggesting that APP/APLP1 interaction with GABA provides a negative-feedback mechanism to preserve homeostatic control of neural circuits. APLP1 localized at the presynaptic active zone, similar to APP and APLP2 (Lassek and others 2016; Rice and others 2019). It contributes to neuronal adhesion (Kaden and others 2009; Mayer and others 2016; Schilling and others 2017) and interacts with signaling molecules of the synaptic vesicle release machinery (Barthet and others 2018; Del Prete and others 2014; Lassek and others 2014; Sullivan and others 2014). Indeed, the intracellular regions of APP, APLP2, and CTF-β interact with several presynaptic proteins involved in vesicle dynamics including Rab, AP-2 subunits, synaptotagmins, clathrin, and complexin (Del Prete and others 2014; Fanutza and others 2015). Depletion of APP causes a reduction in synaptophysin, synaptotagmin-1, and SV2A protein levels pointing toward a role of APP in controlling the synaptic vesicle protein content in the presynaptic active zone. Contrary, an increase in synaptic vesicle proteins is observed, when APLP1 or APLP2 are depleted together with APP (Lassek and others 2014). These results were confirmed in a more recent study by Lassek and others (2016), using the conditional APP/APLP2 DKO mouse model. Moreover, Hick and others (2015) observed altered paired-pulse-facilitation (PPF) behavior at the Schaffer collateral-CA1 pathway for a conditional APP/ APLP2 double KO (DKO) mouse model, a deficit that was restored by the viral-mediated re-expression of the APPsa fragment (Richter and others 2018). It seems likely that APPsa interacts with a7-nAChRs at pre-synapses to increase glutamate release via a 16 aa long sequence located within the C-Terminus of APPsa. Consistently as APPs $\beta$  does not contain these 16 aa, it fails to enhance  $\alpha$ 7-nAChRs mediated currents, to restore LTP and PPF (Richter and others 2018). In addition, high amounts of APPsa were found to interact with the

 $GABA_B$ -R, enhancing short-term facilitation via inhibition of synaptic vesicle release in mouse hippocampal synapses (Rice and others 2019). A $\beta$  also regulates STP by modulation of presynaptic nAchR, presynaptic calcium channels, and vesicle proteins (please refer to sections "APP and Synaptic Transmission" and "Calcium Signaling" for more details).

# Structural Plasticity-Dendritic Spine Morphology/Arborization

Long-term plasticity over days to years is reflected in structural changes including synaptogenesis, formation of dendritic spines, and dendritic sprouting. These processes are not only important during development but also take place throughout the whole adult life and are indispensable for healthy brain function (Holtmaat and Svoboda 2009). In this respect it is noteworthy that several studies provide evidence that APP is involved in processes of neurite outgrowth and branching (Perez and others 1997; Young-Pearse and others 2008). Whereas spine density is reduced in aged APP-KO animals, it remains unaltered in APLP2-KO mice as well as in organotypic hippocampal cultures (OHCs) of APLP2-KOs in vitro (Lee and others 2010; Midthune and others 2012; Weyer and others 2011). Moreover, the quantitative analysis of adult hippocampal CA1 neurons of conditional APP/APLP2 DKO mice yielded pronounced reductions in total neurite length, dendritic branching, reduced spine density, spine head volume, and proportion of mushroom spines, which are thought to represent mature synapses (Hick and others 2015). Weyer and others (2011) therefore elucidated which secreted APP-fragment (APPs $\alpha$  or APPs $\beta$ ) contributes to this effect. They observed normal neuronal morphology in organotypic hippocampal cultures and revealed unaltered neurite length and dendritic branching in APPsa DKO mice (no full length APP or APLP2). In addition, this study reported no significant alteration in spine density in the mid-distal portion of apical dendrites, corresponding roughly to the region in which CA3 axons terminate on CA1 dendrites (Weyer and others 2011; Weyer and others 2014). Overall, these experiments suggest that lack of transmembrane APP/APLP2 isoforms in presence of intact APPs $\alpha$  and APLP1 expression does not affect dendritic structure and spine density of CA1 hippocampal neurons of mice. It can be concluded that APPs $\alpha$  is essential for spine density regulation and dynamics, as highlighted by the effects of exogenous application of the extracellular domain (Tyan and others 2012) or by the knock-in of APPs $\alpha$ (Weyer and others 2011). Weyer and others (2014) reported that CA1 neurons from APLP1-KO or APLP2-KO mice showed unaltered neuronal morphology and no changes in spine density, whereas APP-KO mice revealed a highly reduced dendritic complexity in mid-apical dendrites. Despite unaltered morphology of APLP2-KO pyramidal hippocampal neurons, APP/ APLP2-DKO showed an additional branching defect in proximal apical dendrites, indicating redundancy and a combined function of APP and APLP2 for dendritic morphology. Hippocampal principal neurons in conventional APP-KO showed a significant reduction in spine density and a deficit in the number of large mushroom spines. No further decrease in spine density, however, was observed in APP/APLP2-DKO mice. Expression of APPs $\alpha$  in these DKO mice, which lack the transmembrane APP and express only the secreted APPs $\alpha$  fragment, was sufficient to rescue the defects in spine density observed in APP-KO mice. Collectively, these studies reveal a combined role of APP and APLP2 for dendritic architecture and a unique function of secreted APPs $\alpha$  for normal synaptic density. These findings are of high relevance for AD, since it has been reported that  $A\beta$  release from axons and dendrites reduces local spine number and plasticity (Wei and others 2009).

Authors from the same group reported increased spine numbers in the somatosensory cortex of 4-month-old APP-KO mice (Bittner and others 2009; Zou and others 2016). In a subsequent study, they specified that this finding explicitly included mushroom spines only, while the fraction of thin spines was decreased (Zou and others 2016). Furthermore, they found significantly decreased turnover rates of dendritic spines in APP-KO mice. These alterations of spine morphology might contribute to impaired spine plasticity as thin spines are less stable. This is supported by the finding that 5 weeks exposure to an enriched environment has no effect on spine density in APP-KO mice, while physiologically enhancing it in controls (Montagna and others 2019; Zou and others 2016).

Taken together, APP plays a role in developmentaland plasticity-related changes in spine structure, resulting in a striking decrease in spine density in APP-KO mice.

## Learning and Memory

Core symptoms of AD include impaired working, episodic and spatial memory in parallel to hippocampal atrophy, one of the first brain regions affected by the disease (Scheltens and others 2016). These symptoms could be reliably reproduced in several mouse models of AD and show a correlation to the amyloid plaque burden (Sasaguri and others 2017). However, in some models memory deficits arise before there is a detectable amyloid pathology burden (Sasaguri and others 2017). Cognitive deficits in human patients do also lack a strict correlation with detectable amyloid pathology (Jung and others 2017). An important memory-supporting role of APP is suggested by studies in APP-knockout mice, which show spatial memory impairments in the Morris water maze as well as impairments in conditioned avoidance and novel object recognition tests (Dawson and others 1999; Ring and others 2007). Conditional DKO mice lacking both APP and APLP2 exhibit impaired hippocampus-dependent behavior and learning, including deficits in the nesting test, in the Morris water maze and the radial arm maze (Hick and others 2015). Several studies have shown a strong positive correlation between APPsa levels and cognitive performance. Intraventricular application of picomolar concentrations of APPsa enhanced memory both in mice with pharmacologically induced memory deficits as well as in healthy mice (Meziane and others 1998). In aged rats, exogenous intraventricular APPsa injection ameliorated age-dependent LTP and memory deficits, enhancing object location memory (Xiong and others 2017). Remarkably, in an AD mouse model with amyloid plaques and deficient synaptic function and memory, application of APPsa via viral gene transfer could rescue most of these deficits, decreasing plaque load, increasing dendritic spine density, rescuing LTP, and restoring performance in the Morris water maze (Fol and others 2016). In another study of the same group expression of APPs $\alpha$  on the background of conditional knockout of APP and APLP2 also rescued deficits in spine density, LTP, and PPF, as well as spatial reference memory. Notably, APPs had no significant effects in these animals (Richter and others 2018). While these results highlight the importance of APPs $\alpha$ , it is important to keep in mind that the AICD has also been shown to harbor important functions in plasticity and memory. Mice lacking the AICD and APLP2 (to prevent compensatory mechanisms) exhibit impairments in LTP, in hippocampus-dependent behavior and malformations of the neuromuscular junction (Klevanski and others 2015). A recently discovered interaction of AICD with heterotrimeric G-protein subunits occurring in lipid raft microdomains was shown to provide a positive, regulatory feedback loop enhancing non-amyloidogenic APP processing, thereby preserving spatial memory impairments in 5xFAD mice (Deyts and others 2019). While having detrimental effects on plasticity and memory in nanomolar concentrations as shown in previous sections, physiological (picomolar) concentrations of AB may be involved in memory formation. Thus, enhanced levels of hippocampal A $\beta$  production were quantified during memory induction for contextual fear learning (Puzzo and others 2011).

# **APP and Network Oscillations**

The brain exhibits rhythmic activity over a wide range of frequencies from delta oscillations below 1 Hz to sharp

wave-ripple oscillations (SWP-R) of >200 Hz (Buzsáki and Draguhn 2004). These oscillations represent distinct cognitive and behavioral states and entrain neuronal activity within and between brain areas. In the previous sections we outlined multiple functions of APP and its metabolites both at the pre- and postsynaptic sites and in the cytosol. The APP protein family can both negatively and positively modulate main transmission systems including glutamatergic, GABAergic, and cholinergic signaling and tune neuronal excitability by regulating intracellular calcium levels. As these are pivotal factors

including glutamatergic, GABAergic, and cholinergic signaling and tune neuronal excitability by regulating intracellular calcium levels. As these are pivotal factors contributing to network activity, the cellular functions of APP suggest its importance for the rhythmic activity of the brain and cognition. Indeed, AD patients exhibit prominent changes in the function of large brain networks, particularly those involved in memory formation (Sperling and others 2009; Zott and others 2018). These changes include deceleration of the alpha rhythm (8-12 Hz) toward theta frequency (4-8 Hz in humans), aberrant gamma (30–80 Hz) activity (Hamm and others 2015; Kitchigina 2018), and reduction of slow oscillations during sleep (Lucey and Holtzman 2015). Hyperactivity of local networks such as the hippocampus in parallel to dysfunctional connectivity between large brain networks are suggested to underlie cognitive dysfunction in AD (Zott and others 2018). These clinical findings of neuronal hyperactivity are well in line with results from animal studies discussed in previous sections, showing that pathological AB deposition along with reduced non-amyloidogenic signaling lead to aberrant neuronal activity. Mouse models lacking APP or expressing mutated forms of the protein exhibit alterations in their network function (Korte and others 2012), demonstrating that APPdependent changes at synapses and cells cause effects at the network level. In the following section we review the effects of APP on physiological theta/gamma and SPW-R oscillations as well as pathophysiological network states (Fig. 5).

## Theta/Gamma Oscillations

Theta-nested gamma oscillations are associated with active waking and exploratory behavior in rodents and with cognitive activity in humans (Tamura and others 2017). Compelling evidence from both in vivo and in vitro studies shows reductions in the power of gamma oscillations in mouse models of AD (Korte and others 2012; Verret and others 2012). For example, mice expressing human APP with an AD-associated mutation show impaired gamma activity in vivo. This deficit is associated with reduced expression of the voltage-dependent sodium channel subunit Nav1.1 in parvalbumin-positive interneurons and can be rescued by increasing

	Theta-Gamma THeta-Gamma Theta-Gamma 1000 1000 1000 1000 1000 1000 1000 1000 10		SPW-R N 250 200 u m bay 100 200ms		Epilepsy	
	in vivo	in vitro	in vivo	in vitro	in vivo	in vitro
АРР-КО	= power <b>↓</b> coupling	= power	?	=		
APPsα–KI	=	=	?	=	?	?
	?	=	?	=	?	?
AD models	power coupling	power	?	=		ł

**Figure 5.** APP and network activity. Left: Theta-gamma activity in the murine hippocampus in vivo. The local field potential (LFP) trace (bottom) shows ~5 Hz theta activity with superimposed gamma oscillations on each theta peak. These gamma oscillations are clearly distinguishable as peaks in the ~60 Hz range in the time-frequency plot (spectrogram, upper part). Middle column: Sharp wave-ripple activity (SWP-R) in the murine hippocampus in vivo. Sharp waves are of <100 ms duration and superimposed by ~200 Hz ripple activity. Right column: epileptiform activity recorded in an acute hippocampal mouse slice. APP-KO mice exhibit reduced gamma power in slices and reduced gamma-theta coupling in vivo. In AD models gamma power and gamma-theta coupling are reduced. Mice lacking APP or overproducing amyloid show higher susceptibility to seizures. SPW-R oscillations in vitro are, however, unaltered in these mice. APPs $\alpha$ -KI mice on an APP/APLP2-DKO background exhibit unaltered network activity. APPs $\alpha$ -DM mice expressing APPs $\alpha$  on an APP/APLP2-double knockout background also show largely unaltered network activity in vitro, but their network function has not been studied in vivo.

expression of this subunit (Verret and others 2012). Due to compensation by the homologues, gamma activity is only slightly affected in mice lacking either APP or APLP2 (Zhang and others 2013). In this ex vivo study, gamma frequency in the hippocampal CA3 region in acute slices from APLP2-KO mice was significantly reduced as compared to WT or mice expressing APPs $\alpha$  on the otherwise lethal APP/APLP2-KO background. These data suggest that APP and APLP2 are not essential for gamma oscillations, but rather fine-tune this network activity, for example, modulating its frequency.

The rescue of baseline synaptic transmission, network function, and short-term plasticity by APPs $\alpha$  once again emphasizes the pivotal role of this fragment. While the power of theta and gamma oscillations remained unaltered in APP-deficient mice in vivo, the coupling between these oscillations was strongly diminished specifically in the parietal cortex and the hippocampus, suggesting perturbed communication between large brain networks in these animals (Zhang and others 2016). The role of APLP2 in theta/gamma oscillations in vivo has not been studied yet.

## Sharp Wave-Ripple Oscillations (SPW-R)

SPW-R represent a form of network activity that is prominent in the hippocampal formation during sleep and awake immobility. During ripples (~200 Hz oscillations) distinct populations of neurons (ensembles) fire with submillisecond precision, possibly strengthening connections between co-active neurons and their targets. In line with this notion, SPW-R are crucial for memory consolidation (Buzsáki 2015). Similar to gamma, SPW-R oscillations prevail in APP-deficient mice in vitro (Hefter and others 2016; Zhang and others 2013). In the Swedish mutant mouse model of AD SPW-R activity was also unaltered in comparison to WT despite reduced synaptic excitation in the hippocampal CA1 (Hermann and others 2009). Whether APP affects SPW-R in vivo remains elusive. Further in vivo studies on the influence of APP on SPW-R as a sleep-related activity highly relevant for memory formation might yield promising results as reciprocal connections between sleep disturbances, memory deficits, and amyloid levels have been established in AD patients (Zott and others 2018).

## Neuronal Hyperactivity and Epilepsy

Epilepsy is a pathological condition of recurrent hypersynchronized network activity. AD patients are prone to develop epileptic seizures (Born 2015; Kitchigina 2018), similar to mouse models of AD (Born 2015; Verret and others 2012). On the other hand, mice deficient of APP are also more susceptible to seizures, as shown in the kainate model of epilepsy (Steinbach and others 1998). Neurons in vicinity of amyloid plaques exhibit aberrant activity (Busche and others 2008), which might lead to the development of an epileptic focus. This hyperactivity may, in turn, favor amyloid deposition and compromise the metabolic homeostasis of neurons (Bero and others 2011). In a recent study on another transgenic AD mouse model hyperactivity was found in several brain regions including the hippocampus, beginning at a relatively early stage of the disease. This was associated with the extent of the subsequent tau pathology (Liu and others 2018). Reduction of AB levels by BACE inhibition could rescue circuit function and memory by ameliorating neuronal hyperactivity in a mouse model of AD (Keskin and others 2017). However, the use of BACE inhibitors as potential therapeutic tools for AD is restricted by the importance of its cleavage products in physiological concentrations for synaptic plasticity and network function (as described in section "Long-Term Potentiation and Depression"). In line with this finding, decreasing interstitial amyloid levels is not sufficient to normalize the neuronal hyperactivity typical for AD. Thus, application of amyloid antibodies in a mouse model of AD successfully decreased AB levels, but even exacerbated neuronal dysfunction by facilitating hyperactivity (Busche and others 2015).

# An Integrative Approach to APP Functions in the CNS

In the sections above, we highlighted various functions of APP family proteins at the synaptic, cellular, network, and systemic levels. The current body of knowledge on APP derives to a large extent from transgenic mouse lines with full or partial depletion of APP/APLP or with APP overexpression and increased amyloidogenesis. While this approach has significantly advanced our understanding of particular APP functions, our knowledge about the physiological role of APP in the nervous system is still far from complete. Moreover, the relationship between altered protein levels and disturbed functions may be indirect and nonlinear. Indeed, a large body of literature suggests a U-shaped relationship between APP levels and various observables. For instance, both mice lacking as well as overexpressing APP exhibit age-dependent deficits in LTP and learning, disturbed calcium homeostasis, altered oscillatory activity, and increased vulnerability to seizures and metabolic stress (for details, see sections above). Further complicating factors are the mutual compensation of deficient functions by the homologous proteins APP, APLP1, and APLP2 (Herms and others 2004), differences between functions of full-length APP or its fragments, and the complex regulation of APP processing by secretases. The unexpected discovery of the nsecretase pathway (Willem and others 2015) after three decades of research on APP demonstrates how much is still to be explored. Importantly, APP is not confined to neurons, but is ubiquitously expressed und thus is also suggested to affect the function of astrocytes, microglia, other immune and endothelial cells (Montagna and others 2019). It is likely that APP does also influence brain function via its effects in non-neuronal cells. The diverse expression of  $\alpha$ - versus  $\beta$ -secretases may heavily influence effects of APP in different neuronal subtypes, together with the different availability of downstream interaction partners (e.g., the predominance of GluN2A vs. GluN2B NMDAR). Finally, research on cognitive performance in rodents may not be fully applicable to the situation in humans, despite the high conversation of APP and its isoforms (Sasaguri and others 2017).

In face of all these obstacles, it is not surprising that many questions remain open. Nevertheless, some functions and mechanisms have been firmly established: Expression and processing of APP is controlled by neuronal activity; APP and its metabolites potently modulate synaptic function in various neurotransmitter systems; APP regulates intracellular calcium levels via interaction with calcium channels as well as intracellular stores in an activity-dependent manner; it affects neuronal excitability, gene expression, cell proliferation, and pro-apoptotic pathways. It is known that APP family proteins or their fragments unfold their action in the nucleus (AICD), mitochondria, ER, and lysosomes (A $\beta$ ), in the cytosol (AICD and A $\beta$ ), at the cell membrane (full-length APP), or in the extracellular space (APPs $\alpha$ ). The effects of the APP metabolites APPs $\alpha$  and A $\beta$  mostly antagonize each other (inhibition vs. excitation of synaptic transmission, increasing vs. decreasing calcium levels and cellular excitability, pro-survival vs. pro-apoptotic pathways). In line with this functional



**Figure 6.** Summary of APP functions on multiple biological scales. APP expression and processing are tightly regulated by neuronal activity. Under physiological conditions (e.g., during learning) and compensable stress, upregulation of APP expression and  $\alpha$ -secretase cleavage leads to increased secretion of APPs $\alpha$  and can promote synaptic transmission and plasticity, regulate calcium signaling and homeostasis, thus influencing network activity and promoting learning and memory formation. Furthermore, AICD, which is a product of all APP processing pathways, also modulates these processes. t physiological concentrations, A $\beta$  also appears to harbor important functions in regulation of synaptic transmission and plasticity. Chronic and severe insults such as hypoxia-ischemia, on the other hand, might lead to overexpression of APP and/or shift the balance to amyloidogenic signaling. A $\beta$  at high concentrations forms oligomers and plaques that act toxic at multiple scales, inhibiting synaptic transmission and plasticity, leading to chronically increased cellular calcium levels, impairments in network function, and ultimately to degeneration and dementia.

antagonism, APPs $\alpha$  lowers A $\beta$  levels by direct inhibition of BACE (Obregon and others 2012). Therefore, regulation of APP is a key mechanism for development and function of the nervous system, and possibly a key target for therapeutic interventions.

Its multiple functions make APP a perfect candidate for acting as an acute-phase protein under conditions of cellular stress, for example, in hypoxia-ischemia (HI) or traumatic brain injury (TBI). Indeed, APP is rapidly upregulated in response to metabolic stress and following TBI (Van Den Heuvel and others 2007), and APPknockout mice experience increased mortality in response to HI and TBI (Koike and others 2012; Plummer and others 2016). These protective effects were shown to be once again carried out by APPs $\alpha$  (Hefter and others 2016; Plummer and others 2016) via inhibition of LTCC (Hefter and others 2016). However, presence of membranebound APP (but not APLP1 or APLP2) was required to mediate neuroprotective effects of APPs $\alpha$  by activation of the Akt survival pathway via its C-terminal domain (Milosch and others 2014).

Several lines of evidence indicate a relationship between AD and HI conditions: stroke, heart failure, and metabolic diseases comprise risk factors for AD. Vice versa, AD patients are at a higher risk to suffer a stroke (Tolppanen and others 2013). TBI is also a well-known risk factor for AD. We hypothesize that APP exerts protective effects mostly via APPsa in various acute conditions of brain injury. However, chronic and severe injuries may lead to a maladaptation of APP expression and processing, followed by diminished secretion of APPs $\alpha$ , excessive production of A $\beta$ , and development of AD (Fig. 6). Restoring the balance between APP metabolites toward APPs $\alpha$  is being increasingly discussed as a potential therapeutic strategy in AD (Haass and Willem 2019; Habib and others 2017; Hefter and Draguhn 2017; Mockett and others 2017).

Currently available treatment options for AD are purely symptomatic and can only slightly delay the progression of dementia (Knight and others 2018). The recent failure of large clinical anti-amyloid trials (van Dyck 2018) is a major challenge for presently developed therapeutic approaches. While AD is likely diagnosed far too late for any complete remission, there are no means of prediction or prevention of the disease. We are in urgent need of new interventions at the synaptic, cellular, and network levels, especially at early stages of the disease. Therefore, it seems imperative to tackle the issues described above and advance our understanding of function and malfunction of APP family proteins.

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#### ORCID iD

Dimitri Hefter (D) https://orcid.org/0000-0003-2861-7865

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