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Synapse Formation and Function Is Modulated by the Amyloid Precursor Protein

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The amyloid precursor protein (APP) is critical in the pathogenesis of Alzheimer's disease. The question of its normal biological function in neurons, in which it is predominantly located at synapses, is still unclear. Using autaptic cultures of hippocampal neurons, we demonstrate that hippocampal neurons lacking APP show significantly enhanced amplitudes of evoked AMPA- and NMDA-receptor-mediated EPSCs. The size of the readily releasable synaptic vesicle pool was also increased in neurons lacking APP, whereas the release probability was not affected. In addition, the analysis of spontaneous miniature synaptic currents revealed an augmented frequency in neurons lacking APP, whereas the amplitude of miniature synaptic currents was not found to be altered. Together, these findings strongly indicate that lack of APP increases the number of functional synapses. This hypothesis is further supported by morphometric immuno-histochemical analysis revealing an increase of synaptophysin-positive puncta per cultured APP knock-out neuron. In conclusion, lack of APP affects synapse formation and transmission in cultured hippocampal neurons.

Key words: amyloid precursor protein; Alzheimer's disease; synaptic transmission; amyloid; synaptic vesicle; patch clamp

Introduction

Alzheimer's disease (AD) is the most common form of dementia in the elderly. The pathology of AD is characterized by a progressive neuronal dysfunction, reactive gliosis, and the deposition of amyloid- β (A β) plaques in the brain. These mainly consist of A β peptides 1–40 and 1–42, which are derived from proteolysis of the amyloid precursor protein (APP), a type I transmembrane protein. The duration and severity of cognitive impairment in AD patients is closely associated with synaptic loss (Terry et al., 1991), leading to the notion that synaptic dysfunction is a critical element in the pathogenesis of AD (Selkoe, 2002). Given the pivotal role of APP in AD pathogenesis, it is therefore essential to understand its physiological function, particularly its possible involvement in synaptic transmission.

The proteolytic processing pathways leading to the formation of A β from APP have been well characterized (Selkoe, 2000). APP is delivered to the surface membrane in which it is subjected to proteolytic processing by α -secretase. APP molecules that fail to be cleaved by α -secretase can be internalized into endocytic compartments and subsequently cleaved by β -secretase and γ -secretase to generate A β . A fraction of A β peptides is also generated in the Golgi apparatus and, to a lesser extent, the endoplasmic reticulum. A β peptides generated in the Golgi and in recycling compartments are secreted into the extracellular space

(Greenfield et al., 1999). A β peptides are generated only by the cleavage of APP and not by the cleavage of other APP protein family members such as Apl-1 in Caenorhabditis elegans (Daigle and Li, 1993), Appl in Drosophila (Rosen et al., 1989), APP-like protein 1 (APLP1) (Wasco et al., 1992) and APLP2 (Wasco et al., 1993; Slunt et al., 1994) in mammals, which all lack the A β domain. It follows that the function of the A β domain has to be different from the function of other domains of the APP family member proteins (Herms et al., 2004). The A β sequence is the least conserved part between the human and mouse APP sequences. Mouse APP (when compared with human) is processed poorly by β -secretase, resulting in approximately threefold lower amounts of A β peptide. This suggests that A β peptides may be less critical in mouse than in human brain (De Strooper et al., 1995). This is one possible explanation why mice lacking APP only have subtle phenotypes, including impairment in behavior and long-term potentiation in old age (Zheng et al., 1995; Dawson et al., 1999; Phinney et al., 1999; Seabrook et al., 1999). Recently, however, convincing evidence was presented showing that APP processing has a physiological role in synaptic transmission in mice (Kamenetz et al., 2003). Neuronal activity was found to positively modulate the secretion of A β peptides. In turn, A β was found to suppress excitatory synaptic transmission. It was suggested that A β generation may be part of a negative feedback mechanism that controls neuronal excitability. If this is true, then lack of APP and consequently a total lack in the production of A β peptides should have a physiological effect on synaptic transmission as well. To validate this hypothesis, we studied synaptic transmission in autaptic hippocampal cultures from APP knock-out (KO) mice. This preparation allows a detailed analysis of presynaptic and postsynaptic parameters of synaptic transmission.

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Materials and Methods

APP knock-out mice. The APP knock-out mice used in this study were described originally by Li et al. (1996). The wild-type (WT) mice used in this study are descendants of an F1 generation mouse produced by interbreeding the original parental strain (129Ola and C57BL/6 mice) used to generate the APP knock-out mice initially. Littermates were generated by crossing heterozygous C57BL/6J × 129/Ola APP +/- mice. For genotyping from tail biopsies, a three-primer PCR was used amplifying for the WT allele a 650 bp fragment with primers UM44 (5'-GAGACGAGGACGCT-CAGTCCTAGGG-3') and UM42 (5'-ATCACCTGGTTCTAATCAGAG-GCCC-3') flanking exon 17 and, for the mutant allele, a 430 bp fragment with primers UM42 located in intron 17 and P3-hygro (5'-CGAGATCAG-CAGCCTCTGTTCCACA-3') derived from the phosphoglycerate kinase–hygro cassette, respectively.

Cell culture. Microisland cultures of mouse hippocampal neurons were prepared and maintained as described previously (Bekkers and Stevens, 1991; Rosenmund et al., 1993, 1995). In brief, astrocyte feeder wells enriched in type 1 astrocytes (>95%) were first prepared from postnatal day 1 mouse hippocampal tissue. After dissection and removal of the chorioid plexus and meninges, the material was dissociated and trypsinized. Cells were grown in culture medium consisting of DMEM with 4.5 g/liter glucose (Pan, Aidenbach, Germany), 5% fetal bovine serum (Pan), 100 IU/ml penicillin/streptomycin (Invitrogen, Karlsruhe, Germany), N2 supplement (Invitrogen), Glutamax (Invitrogen), and Mito + serum extender (BD Biosciences, Bedford, MA) at their indicated concentrations. Cells were kept at 37°C in a 10% CO₂-enriched atmosphere. Once a monolayer was achieved, cells were shaken for 12 h so that only astrocytes remained attached to the flask bottom. Astrocytes were trypsinized and replated onto microdot-coated glass coverslips. Hippocampi were dissected from newborn mice pups and digested in DMEM solution containing 0.2 mg/ml cysteine, 100 mm CaCl₂, 50 mm EDTA, and 10 U/ml papain (Roche, Mannheim, Germany) at 37°C. Digestion was stopped after 60 min, and tissue was gently triturated using fire-polished Pasteur pipettes. Cell suspensions, diluted to $5-8 \times 10^3$ cells/ml, were then added to the prepared astrocyte feeder wells (see above). To prevent excessive proliferation of astrocytes, the anti-mitotic agent 5-fluoro-2'-deoxyuridine (8.1 mm) was applied to the cultures on the day after neuronal plating. APP knock-out and wild-type hippocampal neurons were cultured on astrocytic feeder layers prepared from wild-type mice. We did not use APP knock-out astrocytes (which would have had the advantage of an exclusion of astrocytic-derived A β) because we did not know whether the trophic function of astrocytes in hippocampal neuronal cultures would suffer from lack of APP expression in astrocytes. For the γ -secretase inhibitor experiment, 1 μ M N-[N.(3,5difluorophenacetyl-L-alanyl)]-S-phenylglycine t-butyl ester (DAPT) (Calbiochem Merck, Darmstadt, Germany) was added to the neuronal cultures on the day of neuronal plating. Medium was partly exchanged every week and new DAPT was added.

For the A β incubation studies, freshly diluted monomeric A β_{42} (10 nm; Bachem, Bubendorf, Switzerland) was added to the autaptic cultures, and electrophysiological measurements were performed after 48 h.

For the medium exchange experiment, APP knock-out medium was replaced by conditioned wild-type medium 1 week after neuronal plating and the following week.

Electrophysiology. For each experiment, approximately equal numbers of cells from the respective genotypes were measured in parallel and blindly on the same day *in vitro* [18–20 d *in vitro* (DIV)]. Whole-cell recordings were performed using an EPC 9 amplifier (HEKA Elektronik, Lambrecht/Pfalz, Germany). The standard extracellular medium contained the following: 140 mm NaCl, 2.4 mm KCl, 10 mm HEPES, 10 mm glucose, 4 mm CaCl₂, 4 mm MgCl₂, and 15 μ m bicuculline, 300 mOsm, pH 7.3. NMDA EPSCs were measured at 4 mm external Ca ²⁺ (except for 2.7 mm in the experiment with MK-801 (+)-5-methyl-10,11-dihydro-5*H*-dibenzo [a,d] cyclohepten-5,10-imine hydrogen maleate) in the presence of 5 μ m 6-cyano-7-nitroquinoxaline-2,3-dione (CNQX) and 10 μ m glycine and in the absence of external Mg ²⁺. Internal pipette recording solution consisted of the following: 10 mm NaCl, 125 mm K-gluconate, 1 mm EGTA, 4.6 mm MgCl₂, 4 mm Na₂ATP, 15 mm creatine

phosphate, and 20 U/ml phosphocreatine kinase, 300 mOsm, pH 7.3. Data were filtered at 2 kHz and acquired at 10 kHz using Pulse 8.5 software (HEKA Elektronik). Patch electrodes had resistances of 3-4 $M\Omega$. Series resistance and capacity of the electrodes were compensated; only recordings with access resistance below 15 M Ω and with an initial membrane potential less than -50 mV (measured in the current-clamp mode shortly after cell access was obtained) were included in the analysis. Access resistance and cell capacities (5-25 pF) were compensated. Cells were voltage clamped at -70 mV, except for 1 ms depolarizations to 0 mV to elicit an action potential-like stimulus. The rise time constant (10-90%) of the synaptic currents was determined by fitting a linear function to the activating current, and the decay time constant was determined by fitting a single exponential function to the current decay. Detection of miniature EPSCs (mEPSCs) was usually performed for at least 60 s in presence of 200 nm tetrodotoxin (TTX) (Biotrend, Köln, Germany). Data were analyzed with a template detection program (Axograph 4.1; Molecular Devices, Palo Alto, CA). Threshold for detection was set to 3.5 times the baseline SD. Captured mEPSCs of individual cells were averaged to determine mean amplitude and charge per cell. Application of hypertonic solution for pool size determination was made by pulsed application of 500 mm sucrose to the extracellular solution for 4 s using a fast perfusion system (Warner Instruments, Hamden, CT). Evoked responses and responses to hypertonic sucrose solutions were always recorded successively from the same cell. The readily releasable vesicle pool (RRP) per cell was calculated by dividing the charge of the transient sucrose response with the mean mEPSC charge of the same cell, if available. To further determine the mean vesicular release probability $(P_{\rm vr})$, the number of vesicles released by a single action potential-like stimulus was divided by the number of vesicles in the RRP. Recordings and data analysis were performed blindly on each sample.

Immunocytochemistry. Hippocampal cultures of 18-20 DIV were used for immunocytochemistry. For fluorescent microscopic determinations of synaptic density in neurons, cultures were fixed for 10 min with 4% paraformaldehyde, blocked, and permeabilized with PBS containing 2% BSA, 10% goat serum (DakoCytomation, Hamburg, Germany), and 0.3% Triton X-100 (Calbiochem, La Jolla, CA) for 30 min. Cells were incubated with rabbit polyclonal anti-microtubule-associated protein 2 (MAP-2) (1:100; Chemicon, Hampshire, UK) for 60 min, washed with PBS, and incubated with FITC-conjugated goat anti-rabbit IgG (1:100) for another 60 min. After another washing step, fluorescent-labeled mouse monoclonal anti-synaptophysin (1:1000; Synaptic Systems, Göttingen, Germany) was added, incubated for 60 min, and finally fixed in fluorescent mounting medium (DakoCytomation). Images were acquired on an Axiovert 200M microscope (Zeiss, Oberkochen, Germany) with a 63× LD Achroplan objective and Axiovision 4.5 software (Zeiss). For quantification of the total dendritic length and total amount of synapses, analySIS software (Soft Imaging System, Münster, Germany) was used. The summary graph represents four independent experiments using different culture preparations from APP knock-out animals and littermate controls.

All chemicals were obtained from Sigma-Aldrich (Deisenhofen, Germany), unless otherwise noted.

Immunohistochemistry. Six hemispheres (three of each genotype) of 20-d-old and 11-month-old APP knock-out animals and littermate controls were removed and fixed in 4% paraformaldehyde overnight. Coronal 50 µm free-floating sections were prepared with a VT 1000S vibratome (Leica, Wetzlar, Germany) and immunostained with antisynaptophysin and indocarbo-cyanine 3 (Cy3)-conjugated streptavidin. Briefly, sections were placed in PBS, blocked with antibody diluent (Ventana Medical Systems, Illkirch, France), and then incubated overnight at 4°C with the polyclonal antibody against synaptophysin (1:300; Chemicon). This step was followed by incubation in biotinylated swine antirabbit IgG (1:150; DakoCytomation), and finally the sections were fluorescence labeled with Cy3-conjugated streptavidin (1:25; Dianova, Hamburg, Germany). The sections were mounted on precoated glass slides (Super-frost-plus; Menzel, Braunschweig, Germany) and covered (Coverquick; Labonord, Templemars, France). All sections were processed simultaneously under the same conditions. The immunofluorescent labeling protocol was repeated three times to assess the reproducibility of results. Immunolabeled brain sections were viewed with a Zeiss $40\times/0.8$ IR Achroplan objective on a Zeiss Axioskop 2FS MOT LSM510 confocal microscope.

Cy3 was excited with a helium/neon laser at 543 nm, and the emitted fluorescence was detected with a long-pass, 560 nm emission filter. Frames of $230 \times 230~\mu \text{m}$ size were scanned with $0.22~\mu \text{m/pixel}$, and the pinhole diameter was set to an area unit of 1. The z distance of each z stack ranged from 10 to 15 μm and consisted of 20–30, 0.5 μm spaced images.

Synaptophysin levels were assessed in the stratum radiatum and stratum moleculare of the hippocampus, in three coronal sections per animal. Two serial optical z stacks (one for each marker) of the synaptophysin-labeled structures were collected from each section. All series of z stacks from each section were maximum intensity projected and acquired under standard conditions maintaining the same gain, aperture, and black level settings. This method of analysis was evaluated previously by Masliah et al. (1990a,b, 1991). The digitized images were then analyzed regarding the signal intensity and density (covered area of immunohistochemical-positive objects) of synaptophysin-labeled structures (PictureAnalyzer Software; Bjarne Krebs, Zentrum für Neuropathologie und Prion Forschung, Munich, Germany). A threshold was set so that all objects with an intensity value below the threshold were set equal to the background value. For each mouse, the values obtained for the stratum radiatum and stratum moleculare of the hippocampus were averaged.

Western blot. For measurement of the expression of synaptophysin in hippocampus, coronal brain sections were collected on glass slides and Nissl stained. The hippocampal region was then separately lysed and subjected to SDS-PAGE as reported previously (Krebs et al., 2006). After electrotransfer of proteins to the polyvinylidene fluoride membrane, immunoblotting was followed by the incubation with primary antibody against synaptophysin (1:5000; Chemicon) and β -actin (1:5000, actin I-19; Santa Cruz Biotechnology, Heidelberg, Germany) overnight. The visualization was performed with a secondary antibody coupled to alkaline phosphatase (polyclonal goat anti-mouse and anti-rabbit Ig/alkaline phosphatase; DakoCytomation) and chromogene nitroblue tetrazolium salt with 5-bromo-4-chloro-3-indolylphosphate. Western blots were scanned with a high-resolution flatbed scanner, and the density of bands was determined using Total Lab version 2.01 software (Nonlinear Dynamics, Newcastle-upon-Tyne, UK).

Statistical analysis. All error bars indicate the SEM. All results are reported as \pm SEM. A value < 0.05 was considered as statistically significant (Student's t test).

Results

Evoked neurotransmitter release in wild-type and APP knock-out neurons

To determine whether loss of APP leads to alterations in neurotransmitter release, patch-clamp recordings were made from cultured primary hippocampal neurons. Microisland cultures were used to induce the formation of autaptic synaptic connections on individual neurons (Bekkers and Stevens, 1991). Synaptic transmission could then be evoked by briefly depolarizing a voltageclamped neuron, while using the same patch pipette to measure the resulting EPSCs. The advantage of this method over analysis of synaptic transmission in conventional hippocampal neuronal cultures or brain slice preparations is that this approach results in the stimulation of all synapses of a single neuron. Thus, the evocation of the postsynaptic current is not dependent on the stimulus intensity at all. Therefore, the synaptic response does not need to be normalized to stimulus intensity, giving a more accurate measurement of the synaptic strength of single neurons. Using this approach, we concentrated our analysis on glutamatergic neurons because they are much more abundant than inhibitory GABAergic cells in autaptic cultures. We thus recorded AMPAand NMDA-receptor-mediated EPSCs from 24-36 WT and 23-33 APP KO neurons from three to five cultures. All analyses were performed simultaneously on knock-out and control cul-

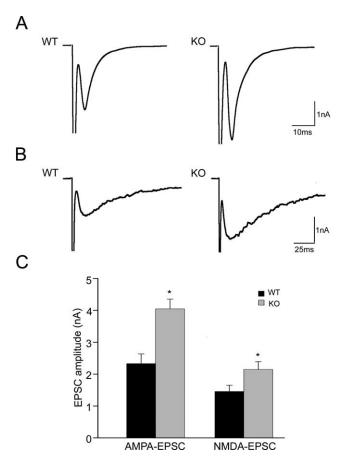


Figure 1. Evoked neurotransmitter release in WT and APP KO neurons. *A*, Representative AMPA EPSC traces from wild-type and APP knock-out neurons. *B*, Representative NMDA EPSC traces from wild-type and APP knock-out neurons. *C*, Mean AMPA and NMDA EPSC amplitudes in neurons from wild-type (n = 24 and n = 35, respectively) and APP knock-out (n = 23 and n = 33, respectively) mice (*p < 0.05).

tures derived from littermates to avoid culture-specific artifacts. No differences were observed in the resting membrane potential between wild-type (59 \pm 3 mV; n=11) and APP knock-out (60 \pm 4 mV; n=11) autaptic hippocampal neurons at 18–20 d in culture.

We then tested the action potential-evoked neurotransmitter release. Synaptic responses were evoked by a brief somatic depolarization (1 ms depolarization from -70 mV holding potential to 0 mV) and measured as peak inward currents a few ms after action potential induction (Fig. 1A,B). The amplitude of the AMPA-mediated EPSCs was enhanced ~1.7-fold in APP knockout cells compared with control littermate neurons (2.33 \pm 0.34 nA for WT and 4.05 \pm 0.26 nA for KO; p < 0.05) (Fig. 1C). Similarly, NMDA-receptor-mediated EPSCs, analyzed by blocking AMPA receptors using CNQX (5 μ M), revealed a 47% increased amplitude in APP knock-out neurons (1.46 \pm 0.19 nA for WT and 2.14 \pm 0.24 nA for KO; p < 0.05) (Fig. 1C). No significant differences were observed in the kinetics (rise time and decay time constant τ) of both AMPA- and NMDA-receptormediated EPSCs between APP knock-out neurons and control cells (Table 1).

Thus, the strength of excitatory synaptic transmission was found to be enhanced in autaptic hippocampal neurons lacking APP. Factors possibly affecting this change include an increase in postsynaptic responsiveness, an alteration in presynaptic release parameters, and a higher number of synapses.

Table 1. Summary of evoked EPSC kinetics

	Rise time (ms)	au (ms)
AMPA EPSC		
WT	1.88 ± 0.14	5.87 ± 0.38
КО	2.30 ± 0.18	6.63 ± 0.32
NMDA EPSC		
WT	7.94 ± 0.71	69.76 ± 5.11
КО	7.33 ± 0.63	58.03 ± 5.40

Data are expressed as mean \pm SEM; n = 20-31 cells per genotype.

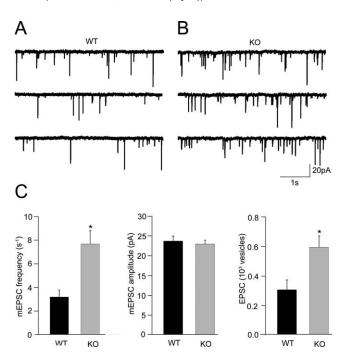


Figure 2. Spontaneous neurotransmitter release in WT and APP KO neurons. **A, B,** Representative mEPSC traces from wild-type and APP knock-out neurons in the presence of 200 nm TTX. **C,** Left, mEPSC frequency in neurons from wild-type (n=25) and APP knock-out (n=23) mice. Middle, mEPSC amplitude in neurons from wild-type (n=25) and APP knock-out (n=23) mice. Right, Estimated EPSC vesicle number per evoked EPSC in wild-type (n=10) and APP knock-out (n=11) neurons (*p<0.05).

Spontaneous neurotransmitter release in wild-type and APP knock-out neurons

As an indicator of postsynaptic responsiveness, we next examined spontaneous mEPSCs (Fig. 2A, B). We found that the frequency of mEPSCs was \sim 2.4-fold higher in APP knock-out neurons than in controls ($3.2 \pm 0.6 \, \text{s}^{-1}$ for WT and $7.6 \pm 1.1 \, \text{s}^{-1}$ for KO; p < 0.05) (Fig. 2C). The mean mEPSC amplitude, however, was not affected ($23.7 \pm 1.2 \, \text{pA}$ for WT and $22.9 \pm 1.0 \, \text{pA}$ for KO; p > 0.05) (Fig. 2C), indicating that the responsiveness and number of postsynaptic receptors is not altered by a lack of APP. In addition, we approximated the number of vesicles released by an action potential-like stimulus by dividing the charge of evoked responses by that of a single mEPSC. For those neurons in which both EPSCs and mEPSCs were measured, the quantity of vesicles released during an EPSC was determined and found to be almost twice as high as in neurons lacking APP ($306 \pm 67 \, \text{for WT}$ and $593 \pm 78 \, \text{for KO}$; p < 0.05) (Fig. 2C).

Readily releasable vesicle pool and release probability in wild-type and APP knock-out neurons

Presynaptic release is thought to depend on the size of the RRP and release probability (P_r) . We therefore performed a detailed

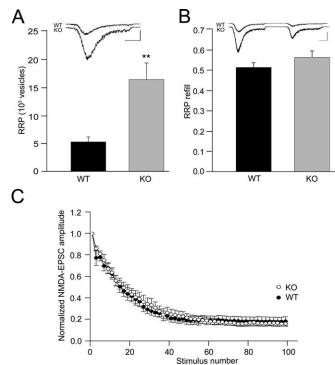


Figure 3. Estimated RRP, RRP refilling, and release probability in WT and APP KO neurons. *A*, Estimated RRP vesicles for wild-type (n=10) and APP knock-out (n=11) mice. Inset shows representative traces. Calibration: 1 s, 0.5 nA. ** p<0.005. *B*, Average ratio of the second hypertonic sucrose response to the initial sucrose response measured 4 s before wild-type (n=14) and APP knock-out (n=18) mice. Inset shows sample traces. Calibration: 1 s, 0.5 nA). *C*, Analysis of MK-801 blocking rate in wild-type (n=13) and APP knock-out (n=10) mice. The average NMDA EPSC amplitude normalized to the first response in the presence of MK-801 (5 μ M) is plotted against evoked EPSC stimulus number.

analysis of the sizes of RRPs in neurons of APP knock-out mice and their littermate controls. The fact that the mEPSC rates were significantly different between the two genotypes already suggested that the size of the RRP pool is changed in the absence of APP. The sizes of RRPs were quantified by measuring the response of cells lacking APP and their wild-type controls to pulsed application of 500 mm hypertonic sucrose solution for 3–4 s. This treatment induces the release of the entire pool of readily releasable vesicles, which in turn leads to a transient inward current followed by a steady current component (Bekkers and Stevens, 1991; Rosenmund and Stevens, 1996). We calculated the amount of RRP vesicles by dividing the total charge of the transient current component after application of hypertonic sucrose by the charge of a single mEPSC measured from the same cell. Neurons lacking APP showed significantly more releasable vesicles in their RRP compared with control cells $(5.36 \pm 0.87 \times 10^{3})$ for WT and $16.35 \pm 2.94 \times 10^3$ for KO; p < 0.005) (Fig. 3A). No differences were observed in the basal rate of RRP refilling, which was estimated by averaging the ratio of the charge of a second hypertonic sucrose response over the initial sucrose response measured 3.5 s before $(0.52 \pm 0.02 \text{ for WT and } 0.57 \pm 0.03 \text{ for KO})$ (Fig. 3B).

To assess whether the efficiency of the action potential-induced vesicle release is influenced by lack of APP, we analyzed $P_{\rm vr}$. We determined $P_{\rm vr}$ by dividing the number of vesicles released during an action potential by the number of vesicles in the RRP. We found that $P_{\rm vr}$ did not vary significantly between neurons from control animals (5.4 \pm 0.5%; n=10) and from animals lacking APP (4.2 \pm 0.5%; n=11; p>0.05). The $P_{\rm vr}$ values obtained are similar to the values obtained in other studies on

autaptic hippocampal neurons (Reim et al., 2001). The synaptic release probability $P_{\rm r}$ depends on the number of fusioncompetent vesicles per synapse and $P_{\rm vr.}$ Observing a significant difference only in the RRP size but not in $P_{\rm vr}$ between the two genotypes, we wanted to rule out a change in synaptic release probability P_r . We estimated P_r by analyzing the progressive block of NMDA-receptor-mediated synaptic currents in the presence of MK-801, which reflects the release probability across all synapses of a given neuron (Rosenmund et al., 1993). APP knock-out and control cells showed the typical amplitude decay in the presence of MK-801 (Fig. 3C). In summary, these results indicate that presynaptic vesicles seem to release efficiently and that the increase in synaptic responses in APP knock-out autapses can be attributed to the increase in RRP.

Prolonged repetitive and highfrequency stimulation in wild-type and APP knock-out neurons

In additional electrophysiological experiments, we analyzed the response of wild-type and APP knock-out neurons to prolonged repetitive and high-frequency stimulation (Fig. 4A, C).

Repetitive stimulation gives insights into the nature of the release probability, $P_{\rm r}$, and the size of the RRP in the central synapse (Schneggenburger et al., 1999). For prolonged repetitive stimulation, we used a 10 Hz protocol and, for high-frequency stimulation, a 50 Hz protocol at 4 mm [Ca²⁺]_{out}. Both in control and in APP knock-out neurons, the relative EPSC amplitude decreases gradually during repetitive stimulation at 10 Hz (Fig. 4*B*). In cells lacking APP, the decay of the relative EPSC amplitude is slightly reduced com-

pared with control cells, without however reaching a significant level. Similar properties were observed in evoked responses elicited by high-frequency stimulation at 50 Hz (Fig. 4D).

γ -Secretase inhibitor enhances evoked and spontaneous neurotransmitter release in WT neurons

To determine whether the increase in EPSC amplitude of APP knock-out neurons is caused by loss of γ -secretase cleavage products, we blocked γ -secretase activity in wild-type neurons by addition of 1 μ M DAPT. DAPT-treated wild-type neurons showed significantly higher AMPA EPSC responses compared with untreated neurons (2.76 \pm 0.30 nA for control WT and 3.66 \pm 0.30 nA for DAPT-treated WT neurons; p< 0.05) (Fig. 5A). The frequency of spontaneous neurotransmitter release was also found to be increased in DAPT-treated wild-type neurons (4.0 \pm 1.1 s $^{-1}$ for control WT and 10.2 \pm 1.8 s $^{-1}$ for DAPT-treated WT neurons; p< 0.05) (Fig. 5B). Both results indicate that the APP knock-out phenotype can be mimicked by adding DAPT to wild-type neurons.

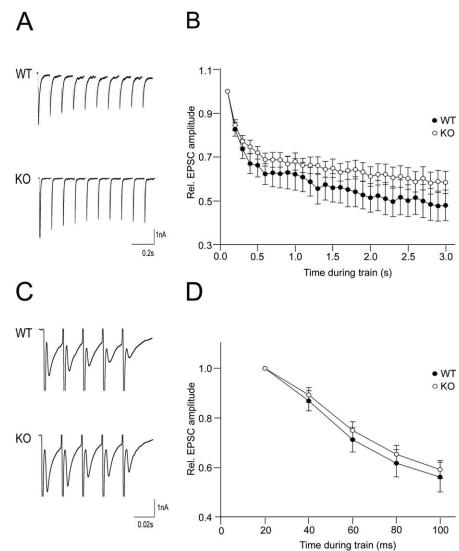
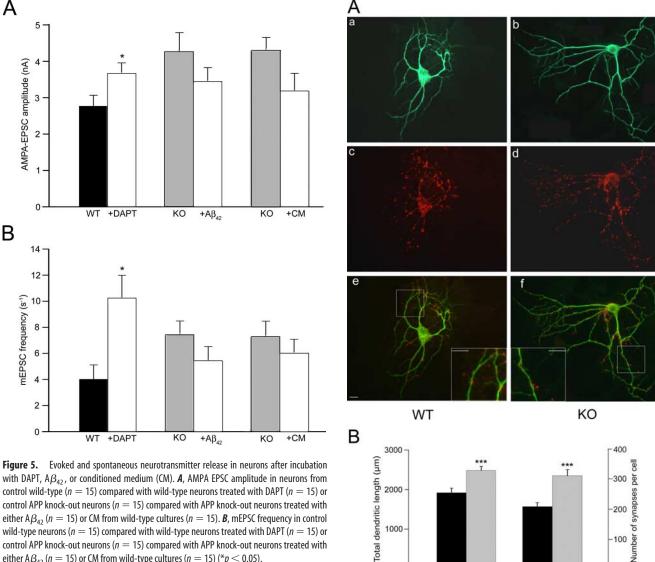


Figure 4. Prolonged repetitive and high-frequency stimulation of WT and APP KO neurons. *A*, Sample recordings of the first 10 responses in a 10 Hz train. *B*, Average EPSC amplitudes of wild-type (n=19) or APP knock-out (n=16) neurons during 10 Hz train normalized to the size of first response. *C*, Sample recordings of five responses in a 50 Hz train. *D*, Average EPSC amplitudes of wild-type (n=22) and APP knock-out (n=23) cells during 50 Hz train normalized to the size of first response.

Addition of $\Lambda\beta_{42}$ or conditioned medium slightly decreases evoked and spontaneous neurotransmitter release in APP knock-out neurons

If the DAPT effect of mimicking the APP knock-out phenotype is mediated by a decreased production and extracellular release of $A\beta$ peptides, then monomeric $A\beta_{42}$ or conditioned medium of wild-type cells should negatively affect synaptic function in APP knock-out cells. Indeed, APP knock-out neurons treated with $A\beta_{42}$ (10 nm) showed a slight but not significant reduction in AMPA EPSC responses compared with untreated APP knock-out neurons (4.25 \pm 0.53 nA for control KO and 3.43 \pm 0.40 nA for Aβ-treated KO neurons; p > 0.05) (Fig. 5A). The frequency of spontaneous neurotransmitter release was also slightly lower in APP knock-out neurons treated with A β_{42} (7.5 \pm 1.0 s⁻¹ for control KO and 5.4 \pm 1.1 s $^{-1}$ for A β_{42} -treated KO cells; p > 0.05) (Fig. 5B). The incubation of APP knock-out neurons with conditioned wild-type medium revealed similar results. Both the amplitude of evoked EPSCs (4.29 \pm 0.36 nA for control KO and 3.18 ± 0.49 nA for KO neurons treated with conditioned medium; p > 0.05) (Fig. 5A) and the frequency of the spontaneous



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either A β_{A2} (n=15) or CM from wild-type cultures (n=15). **B**, mEPSC frequency in control wild-type neurons (n = 15) compared with wild-type neurons treated with DAPT (n = 15) or control APP knock-out neurons (n = 15) compared with APP knock-out neurons treated with either A β_{42} (n=15) or CM from wild-type cultures (n=15) (*p < 0.05).

neurotransmitter release were reduced, but not to statistically significant levels, in APP knock-out cells treated with conditioned medium (7.3 \pm 1.2 s⁻¹ for control KO and 6.0 \pm 1.1 s⁻¹ for KO cells treated with conditioned medium; p > 0.05) (Fig. 5B). In summary, extracellularly applied A β or conditioned medium from wild-type neurons have the tendency to reduce both the amplitude and the frequency of excitatory synaptic currents in autaptic hippocampal cultures of APP knock-out mice, without reaching significant values other than the effect of DAPT on wild-type neurons.

Increase in synapse formation in APP knock-out neurons

Based on the increase in evoked EPSCs and larger RRP in APP knock-out autapses, we next turned to assessing whether the absence of APP alters the actual number of synapses. Therefore, we performed immunofluorescence labeling using antibodies against MAP-2 and synaptophysin, a well characterized presynaptic marker (Fig. 6A) (Augustin et al., 1999). We first evaluated total dendritic length and found a 29% increase in dendritic length in APP knock-out neurons (1929 \pm 113 μ m for WT and 2486 \pm 101 μ m for KO; p < 0.0005) (Fig. 5B). We subsequently determined the total number of synapses per cell and observed significantly more synapses in cells lacking APP (209 \pm 13 for WT

Structural analysis of synapses in microisland cultures from WT and APP KO neurons. A, Immunostains of autaptic neurons. Aa, Ab, Staining for the dendritic marker MAP-2 to assess the size of wild-type and APP knock-out neurons. Ac, Ad, Staining of the same culture for synaptophysin to assess the number of synapses. Ae, Af, Merged picture of MAP-2 and synaptophysin staining. Scale bar, 25 μ m. **B**, Left, Total dendritic length of wild-type (n=55) and APP knock-out (n = 62) neurons (left). Right, Average number of synapses of wild-type (n = 62) 55) and APP knock-out (n = 62) neurons (***p < 0.0005).

ко

100

KO

and 312 \pm 20 for KO; p < 0.0005) (Fig. 6*B*). However, the number of synaptophysin-positive puncta per 10 µm dendrite was not affected by loss of APP (1.19 \pm 0.07 for WT, n = 55; 1.25 \pm 0.06 for KO, n = 62; p > 0.05). Together, these results suggest that, in the absence of APP, neurons form more dendrites and consequently exhibit an enhanced formation of synapses. This can explain the previous results showing an increase in evoked EPSC amplitude and the size of the RRP in APP knock-out neurons as well as the enhanced mEPSC frequency.

Staining intensity and expression of synaptophysin in hippocampus of wild-type and APP knock-out mice

To demonstrate that the increase in synapse formation in cultured APP knock-out neurons is also present in vivo, we performed immunohistochemistry on floating sections of young (20

d) and old (11 months) wild-type and APP knock-out animals. We then analyzed synaptophysin levels in the stratum radiatum and the stratum moleculare of the hippocampus by confocal microscopy (Fig. 7*A*). The average staining intensity of synaptophysin-positive presynaptic terminals in the stratum radiatum was 17% higher in 20-d-old mice lacking APP compared with wild-type mice (77.04 \pm 3.05% for WT and 90.13 \pm 1.95% for KO; p < 0.005) (Fig. 7*B*), whereas no significant difference could be observed in 11-month-old mice $(85.77 \pm 2.19\% \text{ for WT and } 88.50 \pm$ 3.44% for KO; p > 0.05) (Fig. 7*B*). Evaluation of the average synaptophysin-staining intensity in the stratum moleculare showed an enhanced staining intensity in 20-d-old APP knock-out mice (75.87 ± 3.66% for WT and $92.81 \pm 1.00\%$ for KO; p < 0.005) (Fig. 7C). No difference in staining intensity in the stratum moleculare could be found in 11-month-old animals (69.73 \pm 4.58% for WT and 72.90 \pm 4.97% for KO; p > 0.05) (Fig. 7C). Altogether, these results are consistent with an enhanced synapse formation in young APP knock-out hippocampal neurons.

To corroborate the histological difference of synaptophysin expression in wild-type and APP knock-out mice, we performed Western blots from the hippocampal region. Figure 7D shows a typical blot of the hippocampus of a wild-type (left lane) and an APP knock-out (right lane) mouse. Synaptophysin (SYN), detected at 38 kDa (bottom band), gave a stronger signal in the sample of the APP knock-out mouse. The equal loading of tissue was controlled by β -actin (42 kDa; top band) staining. The relative expression

of synaptophysin compared with β -actin was determined by densitometric measurements and showed significantly higher levels in APP knock-out samples (0.38 \pm 0.04 for WT, n=3; 0.44 \pm 0.04 for KO, n=3; p<0.05).

Discussion

Our study has primarily shown that excitatory synaptic transmission in cultured hippocampal neurons lacking APP is strongly enhanced. Both the amplitude of evoked excitatory synaptic currents as well as the frequency of spontaneous synaptic currents were found to be increased. In line with these findings, we observed a significantly enhanced pool size of synaptic vesicles per neuron, whereas the synaptic and vesicular release probability remained unchanged. Quantitative immunohistochemical analysis of cultured autaptic neurons and brain slices from 3-weekold mice further supported the notion that the above electrophysiological observations are most likely attributable to a physical increase in the number of functional synapses per neuron. This conclusion is very much in line with morphometric studies on cultured neurons in APP knock-out mice that noted an enhanced neuritic outgrowth (Young and Selkoe, 2005). An enhanced neuritic growth and consequently an increased num-

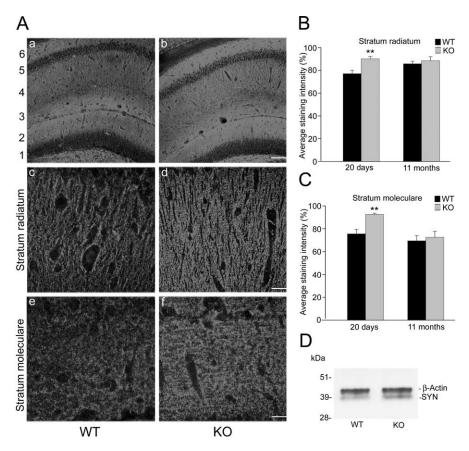


Figure 7. Synaptophysin staining of sections derived from wild-type and APP knock-out mice. **A**, Synaptophysin-positive presynaptic terminals in mouse hippocampus. **Aa**, **Ab**, Low-magnification view of wild-type and APP knock-out hippocampus from 20-d-old mice: 1, stratum granulosum; 2, stratum moleculare; 3, stratum lacunosum; 4, stratum radiatum; 5, stratum pyramidale; 6, stratum oriens. Scale bar, 100 μ m. **Ac**, **Ad**, Synaptophysin staining of the stratum radiatum from 20-d-old wild-type and APP knock-out mice. **Ae**, **Af**, Synaptophysin-positive presynaptic terminals in the stratum moleculare of 20-d-old wild-type and APP knock-out mice. Scale bar, 25 μ m. **B**, Average staining intensity in the stratum radiatum of wild-type (n=18 areas, 9 sections) and APP knock-out (n=18 areas, 9 sections) mice at the age of 20 d and 11 months, respectively. **C**, Average staining intensity in the stratum moleculare of wild-type (n=18 areas, 9 sections) and APP knock-out (n=18 areas, 9 sections) mice at the age of 20 d and 11 months, respectively (**p<0.005). **D**, Representative Western blot analysis of synaptophysin (SYN; bottom band) and β -actin (top band) expression in hippocampal brain sections of wild-type (left lane) and APP knock-out (right lane) mice.

ber of synapses could also explain the enhanced glutamate sensitivity and toxicity in mice lacking APP (Steinbach et al., 1998).

The finding of enhanced synapse formation did not extend to hippocampal slice preparations derived from 11-month-old animals. In line with previous studies on adult APP knock-out brains (Seabrook et al., 1999; Heber et al., 2000), we observed no significant enhancement of the synaptophysin staining in aged mice. Indeed, previous electrophysiological studies performed on slice preparations of adult APP knock-out mice using extracellular stimulation electrodes to induce synaptic vesicle release revealed no significant alterations in the synaptic strength in adult APP knock-out mice (Dawson et al., 1999; Phinney et al., 1999; Seabrook et al., 1999).

What could be the reason why loss of APP significantly affects only the synapse number in young age but not in adult animals? Basically, two mechanisms have to be considered: either loss of APP accelerates synapse formation during development, or an enhanced synapse formation in young APP knock-out mice is compensated by accelerated synapse elimination during later development. We cannot exclude the first alternative, but it does not seem to be very likely. Synapse formation in autaptic cultures of wild-type neurons reaches its plateau during the third week in

culture (Zhang and Benson, 2001), and we studied the neurons at the end of the third week. Rather, we suggest that the physiological mechanisms that modulate synaptic density during late brain development, namely activity-dependent synaptic elimination, convert the APP knock-out phenotype to the wild-type phenotype via those adaptive mechanisms (Cohen-Cory, 2002).

How can a lack of APP enhance the number of synapses per cultured neuron? Interestingly, a recent electrophysiological study on cultured neurons of presenilin-1 (PS1) knock-out mice revealed findings very similar to ours (Parent et al., 2005). Here an enhanced mEPSC frequency as well as enhanced synaptic formation were observed. Because PS1 is part of the γ -secretase complex, critically involved in APP cleavage and A β generation, it is tempting to speculate that the electrophysiological observations we obtained in APP knock-out hippocampal neurons may actually be caused by a lack of γ -secretase-derived APP cleavage products, namely $A\beta$ or the APP intracellular domain (AICD). Here we show that wild-type neurons cultured in the presence of the γ -secretase blocker DAPT indeed reveal significant increases in both the amplitude of evoked EPSCs and the frequency of spontaneous neurotransmitter release. Thus, the APP knock-out phenotype can be mimicked by adding DAPT to cultured wildtype neurons. Whether this is mediated by AICD or intracellular or extracellular A β cannot be answered from our results with certainty. The involvement of AICD, however, does not very likely account for the phenotype we observed in hippocampal neurons lacking APP because APP knock-out mice do not lack AICD completely. There are two proteins expressed in APP knock-out mice that are very similar to APP, APLP1 and APLP2 (Zheng et al., 1995). These proteins are cleaved by γ -secretase as well and give rise to AICD; however, these proteins lack the A β domain. Therefore, we suppose that it is unlikely that a reduced amount of AICD is responsible for the alterations we observed in neurons lacking APP.

It is more likely that the effect is mediated by A β peptides. To study this, we applied synthetic monomeric $A\beta_{42}$ at a low concentration (10 nm) as well as conditioned medium derived from wild-type autaptic cultures onto APP knock-out autaptic cultures. Indeed, we observed a depression of the evoked EPSC amplitude and the mEPSC frequency by extracellular A β but not reaching significant levels as obtained by applying DAPT to wildtype cultures. This may have several reasons. First, DAPT may block the production of A β close to the cell membrane, the location in which A β may bind to certain receptors at the cell membrane involved in synapse formation. When A β is applied to the cell culture medium, it gets highly diluted and its concentration at the synaptic plasma membrane becomes much lower. Second, we applied monomeric A β only for 2 d and in low, physiological concentrations [10 nm (Haass et al., 1992; Seubert et al., 1992; Shoji et al., 1992)] to avoid being confronted with effects attributable to A β oligomerization or aggregation. Those explanations may also hold true for the conditioned medium. A β released from wild-type cells becomes highly diluted within the medium and may also oligomerize or aggregate to some extent, which further reduces the concentration of A β monomers.

Together, therefore, our results strongly indicate that $A\beta$ has an important physiological role in modulating synaptic formation and function. Support for this novel idea is increasing. For instance, it was shown recently that APP overexpression in hippocampal slice preparations depressed both AMPA- and NMDA-receptor-mediated synaptic currents and reduced the miniature EPSC frequency (Kamenetz et al., 2003). Moreover, the expression of APP mutant constructs that do not give rise to $A\beta$ was not

found to induce these phenomena, which again implies that A β generation is indeed critical. There was no indication that AICD is involved because the expression of APP/APLP2 fusion constructs that lack the A β domain was not found to affect synaptic transmission. It may be the function of $A\beta$ to reduce excitatory synaptic transmission via a modulation of the number of functional synapses, thus limiting neuronal excitability. This would be beneficial because it reduces both glutamate excitotoxicity and cellular energy demands. Whether it is primarily the A β monomers, $A\beta$ oligomers, or $A\beta$ fibrils that are involved physiologically is very difficult to assess. Judging from our results based on the comparison of APP knock-out mouse and wild-type mouse cultured neurons of very low density (<10,000 cells per well), conditions in which appreciable oligomer formation is highly unlikely, the effect of $A\beta$ on synaptic formation is most likely mediated by $A\beta$ monomers.

The mechanisms by which $A\beta$ modulates neurite outgrowth and the number of functional synapses is still unknown. One possible mechanism may be related to the observation that the unaggregated, nonpathological form of $A\beta$ modulates the function of calcium and potassium channels within the cell membrane (Price et al., 1998; Ramsden et al., 2001, 2002). This could in turn influence neurite outgrowth and synapse formation. The motility of dendritic filopodia, for example, is modulated by depolarization-induced calcium influx (Jontes et al., 2000).

How can the proposed physiological function of monomeric $A\beta$ in regulating synaptic formation be related to Alzheimer's disease? It is well known that APP is highly expressed under all kinds of neuronal injury. Under those conditions, A β may reduce excitatory synaptic transmission via a modulation of the number of functional excitatory synapses, thus limiting excitotoxicity. Chronically enhanced levels of $A\beta$, however, attributable to repetitive mild neuronal injuries of any kind including hypoxia or trauma, may either directly alter synaptic transmission via a gainof-function mechanism or lead to the formation of toxic A β aggregates or oligomers. These A β aggregates or soluble oligomers may further alter synaptic transmission and cellular excitability by altering the function of voltage-gated calcium channels (Price et al., 1998; Ramsden et al., 2002), certain types of potassium channels (Colom et al., 1998; Yu et al., 1998), AMPA receptors (Chang et al., 2006), NMDA receptors (Snyder et al., 2005), and the α 7 nicotinic receptor (Wang et al., 2000; Dineley et al., 2001) or by the formation of calcium-permeable membrane pores (Arispe et al., 1993; Hirakura et al., 2000; Kawahara et al., 2000; Kourie et al., 2001; Kayed et al., 2004; Demuro et al., 2005). Alternatively, as suggested by Kamenetz et al. (2003), neurons may lose their sensitivity to A β -induced synaptic depression during aging, for example, via the modulation of AMPA-receptormediated synaptic currents (Kamenetz et al., 2003; Almeida et al., 2005), which may cause a persistently elevated neuronal activity known to be toxic.

The concept of a physiological function of A β , previously proposed also by Plant et al. (2003) and Ramsden et al. (2002), may hence have an important impact on therapeutic approaches that aim at reducing A β production in AD patients.

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