



Published in final edited form as:

Biol Psychiatry. 2019 August 01; 86(3): 208–220. doi:10.1016/j.biopsych.2019.04.026.

A Novel Apolipoprotein E Antagonist Functionally Blocks Apolipoprotein E Interaction With N-terminal Amyloid Precursor Protein, Reduces β -Amyloid-Associated Pathology, and Improves Cognition

Darrell Sawmiller^{*,1}, Ahsan Habib^{*,1}, Huayan Hou^{*,1}, Takashi Mori², Anran Fan¹, Jun Tian¹, Jin Zeng¹, Brian Giunta³, Paul R. Sanberg⁴, Mark P. Mattson^{5,6}, and Jun Tan^{1,#}

¹Rashid Laboratory for Developmental Neurobiology, Silver Child Development Center, Department of Psychiatry & Behavioral Neurosciences, Morsani College of Medicine, University of South Florida, Tampa, Florida 33613, USA.

²Departments of Biomedical Sciences and Pathology, Saitama Medical Center and Saitama Medical University, Kawagoe, Saitama 350-8550, Japan.

³Neuroimmunology Laboratory, Department of Psychiatry & Behavioral Neurosciences, Morsani College of Medicine, University of South Florida, Tampa, Florida 33613, USA.

⁴Department of Neurosurgery & Brain Repair, Center for Aging and Brain Repair, Morsani College of Medicine, University of South Florida, Tampa, Florida 33613, USA.

⁵Laboratory of Neurosciences, National Institute on Aging Intramural Research Program, Baltimore, Maryland 21224, USA.

⁶Department of Neuroscience, Johns Hopkins University School of Medicine, Baltimore, Maryland 21224, USA.

Abstract

[#]Correspondence and requests for materials should be addressed to Jun Tan, Department of Psychiatry & Behavioral Neurosciences, Morsani College of Medicine, University of South Florida, Tampa, Florida, jtan@health.usf.edu.

^{*}These authors contributed equally to this work.

AUTHOR CONTRIBUTIONS

D.S. and J.T. designed all experiments; D.S., H.H., A.H., A. F., J.Tian, and J.Z. carried out all experiments except for IHC and IF analyses (A.H. and T.M.); D.S., H.H., and J.T. conducted statistical analysis of the data; J.T. supervised the project; D.S. and J.T. wrote the manuscript and M.P.M., T.M., A.H. B.G., and P.R.S. critically edited the manuscript.

FINANCIAL DISCLOSURE

J.T., D.S., H.H., and A.H. are inventors on a patent application submitted by University of South Florida. All other authors report no biomedical financial interests or potential conflicts of interest.

DATA AVAILABILITY

The authors declare that all data supporting the findings of this study are available within the article and its supplementary files and from the corresponding author upon reasonable request.

Publisher's Disclaimer: This is a PDF file of an unedited manuscript that has been accepted for publication. As a service to our customers we are providing this early version of the manuscript. The manuscript will undergo copyediting, typesetting, and review of the resulting proof before it is published in its final citable form. Please note that during the production process errors may be discovered which could affect the content, and all legal disclaimers that apply to the journal pertain.

BACKGROUND—The E4 isoform of apolipoprotein E (apoE4) is a major genetic risk factor for the development of sporadic Alzheimer’s disease (AD) and its modification has been an intense focus for treatment of AD in recent years.

MEHTHODS—We investigated the binding of apoE, a peptide corresponding to its low density lipoprotein receptor (LDRL) binding domain (aa 133–152, ApoEp) and modified ApoEp to amyloid precursor protein (APP) and their effects on A β production in cultured cells. Having discovered a peptide which blocks the interaction of apoE with N-terminal APP, we investigated the effects of this peptide and ApoEp on AD-like pathology and behavioral impairment in 3XTg and 5XFAD transgenic mice.

RESULTS—ApoE and ApoEp, but not truncated apoE lacking the LDLR binding domain, physically interacted with N-terminal APP and thereby mediated A β production. Interestingly, the addition of six lysine residues to the N-terminal ApoEp (6KApoEp) directly inhibited apoE binding to N-terminal APP and markedly limited apoE- and ApoEp-mediated A β generation, presumably through decreasing APP cellular membrane trafficking and p44/42 mitogen-activated protein kinase phosphorylation. Moreover, while promoting apoE interaction with APP by ApoEp exacerbated A β and tau brain pathologies in 3XTg-AD mice, disrupting this interaction by 6KApoEp ameliorated cerebral A β and tau pathologies, neuronal apoptosis, synaptic loss, and hippocampal-dependent learning and memory impairment in 5XFAD mice without altering cholesterol, LDLR, and apoE expression levels.

CONCLUSIONS—These data suggest that disrupting apoE interaction with N-terminal APP may be a novel disease-modifying therapeutic strategy for AD.

Keywords

acetylated and phosphorylated tau; Alzheimer’s disease; Alzheimer’s mouse model; amyloidogenesis; amyloid precursor protein; apolipoprotein E; low density lipoprotein receptor binding domain

INTRODUCTION

Afflicting as many as 47 million people worldwide, Alzheimer’s disease (AD) is characterized by the accumulation of amyloid- β (A β) plaques and neurofibrillary tangles (NFTs) in the brain (1–4). A β is produced *via* β - and γ -secretase-mediated proteolysis of amyloid precursor protein (APP) (5,6), a type I transmembrane protein, which can then aggregate to form plaques. According to the amyloid hypothesis, overproduction of A β triggers neuronal apoptosis (7,8), inflammation (9, 10), oxidative stress (11,12), and tau phosphorylation and aggregation in intracellular neurofibrillary tangles (NFTs) (13). While inherited early-onset familial AD (FAD) results from mutations in *APP* or *presenilin* (*PS*) genes, in the more common late-onset sporadic AD (SAD) excess A β generation is enhanced by age-related factors, metabolic dysfunction, cardiovascular disease, and brain injury (14–16). In addition, the apoE4 isoform of apolipoprotein E (apoE4) has been found to be the major genetic risk factor for the development of SAD (17). Recently, since several therapeutic approaches targeting A β have failed, the relevance of A β in the AD pathogenesis has been questioned.

ApoE, a major component of chylomicron remnants and very-low, intermediate, and high-density peripheral- and brain-derived lipoproteins, plays an important role in receptor-mediated cholesterol endocytosis (18,19). The cause of the increased risk for AD associated with the presence of apoE4 may involve enhanced formation and reduced clearance of A β (20,21), the formation of neurotoxic apoE4 peptide fragments (22,23), abnormal tau phosphorylation, neuroinflammation, and neurodegeneration (24–27). Early studies utilizing yeast two-hybrid and immunoprecipitation suggested that the N-terminal APP can directly bind to apoE, thereby enhancing intracellular APP endocytosis and reducing sAPP α production (28). More recent study indicates that both glia-derived and recombinant apoE stimulates AP production in human neurons with a rank order of potency of apoE4 > apoE3 > apoE2, mediated by activation of a non-canonical mitogen activated protein kinase (MAPK) p44/p42 and enhanced transcription/translation of APP (29). These studies suggest that apoE directly binds to the N-terminal region of APP, thereby enhancing APP endocytosis and directing its processing from sAPP α to Ap. Recently, we further explored the binding of apoE to the N-terminal APP in A β production and generated a novel peptide antagonist of this interaction, which reduced A β production and pathology in AD mouse models. Taken together, our results suggest that disruption of apoE interaction with the N-terminal APP may be a novel disease-modifying therapeutic strategy for AD.

METHODS AND MATERIALS

Cell culture

CHO cells engineered to express human wild-type APP (CHO/APPwt) or Swedish mutant APP (CHO/APPswe) were cultured in 96 or 24 well plates at 4×10^4 or 2×10^5 cells/well, respectively, in Dulbecco's modified Eagle's medium (DMEM) with fetal bovine serum (FBS, 10%), 1 mM sodium pyruvate, and 100 U/mL of penicillin/streptomycin. In addition, SH-SY5Y cells transfected with APPswe and wild-type SH-SY5Y cells were cultured as previously described (30). Primary hN2TM human neurons were cultured in hN2TM human neuron culture media and primary HCN2 human neurons were cultured in DMEM with 4 mM L-glutamine adjusted to contain 1.5 g/L sodium bicarbonate, 4.5 g/L glucose, and FBS (10%).

Enzyme-linked immunosorbent assay, Western blotting, and immunoprecipitation

A β _{1–40,42} and sAPP α from cell cultures and brain homogenates were detected by A β and sAPP α enzyme-linked immunosorbent assay (ELISA) kits (IBL-America), strictly following the manufacturer's instructions (31,32). Western blotting (WB) analyses were performed as previously described (31,32). Immunoprecipitation (IP) was performed by first incubating conditioned media or cell lysates with appropriate antibodies and Protein-A/G Mag Sepharose beads (GE Healthcare Life Sciences, Pittsburgh, PA) overnight with gentle rocking at 4°C, followed by three washes with binding buffer (50 mM Tris, 150 mM NaCl, pH 7.5) and analysis by WB. The effect of apoE4 and 6KApoEp on cell surface expression of APP, LDLR, and LRP1 was determined by biotinylation and avidin precipitation as described previously (33).

Real-Time PCR

Total RNA was extracted from CHO/APPwt cells after treatment with 6KApoEp using RNeasy Plus Mini Kit (Qiagen). The purity and concentration of RNA was quantified using Nanodrop 2000c spectrophotometer (Thermo Fisher Scientific). The quantification of target RNAs was performed in a total volume of 50 μ L by real-time one-step RT-qPCR reactions (10 ng of RNA, 250 mM forward and reverse primers) using SYBR Green I in a IQ5 multi-color real-time PCR detection system (Bio-Rad), according to manufacturer's instructions. RNA primers were designed to selectively amplify and quantify human APP and LDLR and CHO cell β -actin (IDT) as indicated.

Cholesterol quantitation

For determination of CHO/APPwt cell associated cholesterol levels, cells were washed three times with ice-cold PBS, lysed by sonication in chloroform:isopropanol:IGEPAL CA-630 (7:11:0.1), and centrifuged. The organic phase was air dried at 50°C to remove chloroform and diluted 10-fold in the cholesterol assay buffer for the cholesterol determination by fluorometric cholesterol quantitation kit according to manufacturer's instructions (MilliporeSigma).

Mice

All mice were housed and maintained in the Morsani College of Medicine Animal Facility at the University of South Florida (USF), and all experiments were conducted in compliance with protocols approved by the USF Institutional Animal Care and Use Committee. 5XFAD mice at 6 weeks of age ($n =$ ten, five female/five male) were intraperitoneally (i.p.) treated with 6KApoEp (250 μ g/kg in 50 μ L PBS) or PBS (50 μ L) daily for 12 weeks. 3XTg-AD mice at 9 months of age ($n =$ six, three female/three male) were i.p. treated with ApoEp (250 μ g/kg in 50 μ L PBS) or PBS (50 μ L) daily for 12 weeks. After 11-week treatment, 5XFAD mice were subjected to the Y Maze, fear conditioning, novel object recognition and open field behavioral testing as described previously (34,35).

After treatment and behavioral testing, all mice were anesthetized with 2–4 % isoflurane (Millipore Sigma), followed by collection of blood, euthanization by bilateral thoracotomy, transcardial perfusion with physiological saline containing heparin (10 U/mL, Millipore Sigma), and isolation of the brain for biochemical, immunohistochemical (IHC), and immunofluorescence (IF) analyses. Briefly, one hemisphere was frozen immediately in liquid nitrogen and stored at -80°C , followed by sonication in RIPA buffer (Cell Signaling Technology) containing protease inhibitor and phosphatase inhibitor cocktail (Thermo Fisher Scientific), centrifugation, and WB analysis of the supernatant. The other hemisphere was placed in 4% paraformaldehyde in PBS for cryostat sectioning. The 25- μ m free-floating coronal sections were collected and stored in PBS with 100 mM sodium azide at 4°C for IHC and IF analyses.

IHC and IF

Brain sections from 5XFAD and 3XTg-AD mice were stained with biotin anti- $A\beta_{17-24}$ monoclonal antibody (4G8), VECTASTAIN *Elite* ABC Kit (Vector Laboratories, Burlingame, CA) and diaminobenzidine substrate, followed by quantitative image analysis

of A β burden, as described previously (31,32). In addition, brain coronal sections were analyzed by IHC staining with anti-acetylated tau (K¹⁷⁴, K²⁷⁴) and anti-phosphorylated tau (Thr²³¹, Thr¹⁸¹, Thr⁴⁰⁴, and Ser²⁰²/Thr²⁰⁵) antibodies and IF staining with anti- β -tubulin III, anti-NeuN, anti-cleaved caspase-3, anti-synapsin I, anti-apoE, and anti-N-terminal APP antibodies. For thioflavin S staining, free floating brain tissues sections mounted on slides were washed in ddH₂O, stained in 1% filtered thioflavin S for 5 min and differentiated in 70% alcohol. For all cell culture and brain tissue staining analyses, images were taken by a BX60 microscope with an attached CCD camera system (DP-72, Olympus, Tokyo, Japan), or using an Olympus FV1000 laser scanning confocal microscope.

Statistical analysis

All data were normally distributed. Therefore, in instances of single mean comparisons, Levene's test for equality of variances followed by *t*-test for independent samples was used to assess significance. In instances of multiple mean comparisons, analysis of variance was used, followed by *post-hoc* comparison using Bonferonni's method. Alpha levels were set at 0.05 for all analyses. The statistical package for the social sciences release 23.0 (IBM SPSS) was used for all data analyses.

RESULTS

6KApoEp treatment markedly suppresses human recombinant and lipidated apoE-induced A β production

Previous findings suggest that apoE4 might exacerbate AD pathology, in part, by enhancing APP amyloidogenic processing (29). In addition, the N-terminal region of apoE (residues 133–152) is known to contain the apoE receptor binding domain, while structural modifications of apoE are known to mediate differential interaction of apoE isoforms with its receptor (21). In order to determine the specific region of apoE mediating A β production and further investigate the effects of structural modifications of apoE, we focused on the apoE LDLR binding domain and proximal structural modifications. As expected, an apoE peptide (ApoEp), consisting only of the LDLR binding domain of apoE (residues 133–152), markedly increased A β production in a concentration-dependent manner in CHO cells engineered to stably overexpress human wild-type APP (CHO/APPwt cells, Figure 1A, B and Supplement Figure 1A). Interestingly, while N-terminal addition of Flag-tag greatly enhanced the efficacy of ApoEp to increase A β _{1–40, 42} levels, the addition of three lysine residues (3K) terminated this amyloidogenic effect (Figure 1A).

To test the hypothesis that the addition of N-terminal lysine residues might convert ApoEp to an apoE antagonist, we examined the effects of ApoEp containing 3, 6, 7, 8, or 9 lysines on apoE4-induced A β production. 3KApoEp moderately while 6 to 9KApoEp markedly and maximally reduced both basal and apoE4-induced A β production (Figure 1C and Supplement Figure 1B–D). In addition, 6KApoEp reduced apoE4-induced A β production in a concentration-dependent fashion, starting at 40 nM and with an IC₅₀ of approximately 0.32 to 0.63 μ M (Figure 1D, E and Supplement Figure 1C). This reduction was observed regardless of whether the cells were co-treated with 6KApoEp together with apoE4, pre-treated with 6KApoEp, or treated with apoE4 pre-incubated with 6KApoEp (Supplement

Figure 3). 6KApoEp also reduced A β production elicited by HDL-lipidated apoE4, human plasma-derived apoE3, and human astrocyte media-derived apoE in CHO/APPwt and SH-SY5Y human neuroblastoma cells as well as hN2TM and HCN2 human neurons (Figure 1F–H), confirming that 6KApoEp reduces natural apoE-mediated A β production.

6KApoEp reduces physical association of apoE with N-terminal APP

Previously, apoE was found to bind to the N-terminal APP, upstream of the A β region, and thereby enhances APP endocytosis and reduces sAPP α production (28). In order to confirm this physical association of apoE with N-terminal APP, CHO/APPwt cells were treated with ApoEp, 6KApoEp, 6K, apoE3, or apoE4 followed by immunoprecipitation (IP of apoE with anti-LDLR binding domain antibody (6H3B5). Total secreted APP in conditioned media and full length APP in cell lysates were then determined by WB analysis using anti-N-terminal (22C11) and anti-C-terminal APP antibodies (pC-APP), respectively. Alternatively, N-terminal APP was immunoprecipitated with anti-N-terminal APP41/66 antibody (pN-APP) followed by analysis of apoE, 6KApoEp, and ApoEp using anti-human apoE (mApoE) and 6H3B5 antibodies. ApoE, ApoEp, and 6KApoEp were co-immunoprecipitated with secreted APP in conditioned media and full length APP in cell lysates (Figure 2A, B and Supplement Figure 2A–C). In addition, ApoEp and 6KApoEp were co-localized with N-terminal APP in cultured CHO/APPwt cells, as determined by immunofluorescence (IF) staining with 6H3B5 and pN-APP antibodies (Figure 2C), respectively, confirming that N-terminal APP physically associates with apoE in vitro. ApoE was also co-immunoprecipitated with N-terminal APP in homogenates prepared from brains of TgAPPwt mice, AD patients, and age-match controls, confirming that the physical association of apoE with N-terminal APP occurs in vivo (Figure 2D, E). Most interestingly, apoE was colocalized with cell surface APP more in brains from aged compared with young TgAPPwt mice, suggesting an age-associated increase in apoE-N-terminal APP interaction (Figure 2F). However, WB analysis showed no significant differences in total apoE and APP levels between young and aged TgAPPwt mouse brain homogenates (data not shown).

As in vitro confirmation of physical association of apoE with N-terminal APP, human recombinant sAPP β (without A β domain), sAPP α , CHO/APPwt conditioned media-derived sAPP α (cmsAPP α), or APP N-terminal peptide (aa 31–60, APP31/60) was incubated with 6KApoEp, 6K, apoE3, or apoE4 followed by IP of apoE LDLR binding domain with 6H3B5 and WB analysis of sAPP α / β and APP31/60 (Figure 2G, H). Human recombinant sAPP α protein was also incubated with apoE4 in the absence or presence of 6KApoEp or 6K followed by IP with pN-APP and WB analysis of apoE and sAPP α (Figure 2I, J). Overall, sAPP α / β and APP31/60 were co-immunoprecipitated with 6KApoEp and apoE in vitro, confirming that 6KApoEp and apoE physically associates with N-terminal APP, and 6KApoEp reduced this association.

6KApoEp reduces A β production by blocking apoE interaction with N-terminal region of APP

Since 6KApoEp reduced apoE- and ApoEp-mediated A β production (Figure 1) as well as the physical association of apoE with N-terminal APP (Figure 2 and Supplement Figure 2), we wished to further investigate if apoE LDLR binding domain interaction with N-terminal

APP mediates A β production. Conditioned media collected from CHO/APPwt cells transiently coexpressing human wild-type apoE4 (CHO/APPwt/ApoEwt cells), truncated apoE4 lacking the LDLR binding domain (CHO/APPwt/ApoEtrun cells), or control vector (CHO/APPwt/Ctrl cells) were immunoprecipitated with anti-N-terminal APP antibody (pN-APP) followed by WB analysis of apoE and sAPP α . ApoEwt but not ApoEtrun was co-immunoprecipitated with sAPP α , confirming interaction of apoE with N-terminal APP *via* its LDLR binding domain (Figure 3A). As confirmation, ApoEwt but not ApoEtrun was co-localized with N-terminal APP, as determined by IF staining (Figure 3C). In addition, CHO/APPwt/ApoEwt cells produced markedly more A β than CHO/APPwt/ApoEtrun or CHO/APPwt/Ctrl cells. 6KApoEp reduced A β production in CHO/APPwt/ApoEwt cells and ApoEp enhanced A β production in CHO/APPwt/ApoEtrun cells, while conditioned media collected from CHO/APPwt/ApoEwt but not CHO/APPwt/ApoEtrun cells increased A β production in CHO/APPwt cells (Figure 3B). Thus, the apoE binding domain is required for apoE association with N-terminal APP and apoE-mediated A β production.

In order to confirm that apoE binding domain mediates A β production by interaction with the N-terminal region of APP, CHO/APPwt cells were treated with apoE3, apoE4, or ApoEp in the absence or presence of N-terminal APP (22C11) or apoE LDLR binding domain antibodies (6H3B5). Antibodies 22C11 and 6H3B5 reduced apoE3, apoE4, and ApoEp-mediated A β production in cultured cells in a dose-dependent fashion (Figure 4A and Supplement Figure 4). In addition, CHO/APPwt cells and CHO cells expressing truncated APP lacking the N-terminal E1 region (CHO/APPdE1 cells) were treated with 6KApoEp, ApoEp, apoE3, or apoE4 followed by analysis of A β production. CHO/APPdE1 cells produced markedly less A β production compared with CHO/APPwt cells after treatment with ApoEp, apoE3, or apoE4 (Figure 4B, *lower panel*). In contrast, CHO/APPwt and CHO/APPdE1 cells elicited similar spontaneous A β production in the absence of apoE and 6KApoEp reduced A β production elicited by CHO/APPwt but not CHO/APPdE1 cells. Therefore, apoE and ApoEp increases and 6KApoEp reduces A β production by binding to the N-terminal E1 region of APP, but this region of APP is not required for spontaneous A β release in the absence of apoE.

6KApoEp inhibits cell surface APP trafficking and p44/42 MAPK phosphorylation

Under physiological conditions, APP is known to be synthesized in the endoplasmic reticulum and trafficked *via* the trans-Golgi network to the plasma membrane where approximately 90% of it is cleaved by members of a disintegrin and metalloproteinase domain-containing protein (ADAM) (α -secretase), yielding a membrane-bound α -C-terminal APP fragment (α -CTF) and secreted sAPP α (36). The remaining unprocessed APP (~10%) can be cleaved at the plasma membrane or further trafficked back into the cell by endocytosis, followed by cleavage by β -site APP converting enzyme 1 (β -secretase, BACE1), yielding β -C-terminal fragment (β -CTF) and sAPP β , and by γ -secretase, ultimately generating A β peptides which can then be secreted from the cell (37–39). Since α -secretase cuts APP within the A β region, increasing A β generation *via* the endocytic pathway precludes sAPP α production. We hypothesized that decreased A β generation by 6KApoEp might result from decreased membrane APP trafficking and subsequent amyloidogenic processing. To test this hypothesis, we investigated the effects of apoE4 and

6KApoEp on sAPP α and A β production into the media as well as β -CTF and total APP levels in the plasma membrane.

CHO/APPwt cells treated with apoE4 markedly increased A β and β -CTF levels, while reducing sAPP α levels, presumably by enhancing APP endocytosis (Figure 5A–C). Indeed, apoE4 enhanced A β and β -CTF levels much more in CHO/APPwt cells compared with CHO cells overexpressing APP with the Swedish mutation (CHO/APP^{sw} cells), which is a better substrate for BACE1 and directly processed to A β prior to its trafficking to the cell surface (Supplement Figure 8). In contrast, 6KApoEp reduced both basal and more profoundly apoE4-induced production of A β , β -CTF, and sAPP α , without altering total APP levels, suggesting that 6KApoEp inhibits an early stage of APP processing, such as the initial trafficking of APP to the plasma membrane. As further confirmation, 6KApoEp reduced both basal and apoE4-mediated cell surface APP levels, as determined by WB, flow cytometry, and confocal microscopy, regardless of whether the cells were co-treated with 6KApoEp together with apoE4, pre-treated with 6KApoEp, or treated with apoE4 pre-incubated with 6KApoEp (Figure 5D–G and Supplement Figure 3B). 6KApoEp also reduced A β and β -CTF levels more in CHO/APPwt than in CHO/APP^{sw} cells, consistent with reduction of APP trafficking to the cell surface (Supplement Figure 8). Notably, apoE4 and 6KApoEp, in the absence or presence of β - or γ -secretase inhibitors, did not alter cell surface protein levels of LDLR or LRP1 as well as mRNA levels of APP or LDLR, suggesting that apoE and 6KApoEp do not alter APP, LDLR and LRP1 expression or recycling (Figure 5D and Supplement Figures 9, 10).

Previously, apoE-mediated APP transcription/translation and A β production were found to be mediated by activation of a non-canonical p44/42 MAPK (29). In order to determine if apoE- and ApoEp-mediated APP trafficking and processing might also be mediated by this signaling pathway, we determined the effects of apoE, ApoEp, and 6KApoEp on p44/42 and p38 MAPK phosphorylation. While ApoEp, apoE3, and apoE4 activated both p44/42 and p38 MAPK phosphorylation, 6KApoEp only activated p38 MAPK phosphorylation and inhibited apoE-induced p44/42 phosphorylation (Figure 5H and Supplement Figure 5). Taken together, our findings suggest that ApoEp, apoE4, and 6KApoEp may have different effects on APP processing, with ApoEp and apoE4 enhancing and 6KApoEp reducing cell surface APP trafficking, APP endocytosis, and amyloidogenic processing, potentially mediated by differential activation of MAPK pathways.

6KApoEp reduces cerebral β -amyloid and tau pathologies, and memory impairment in AD mouse models

Since 6KApoEp inhibits apoE-APP receptor mediated A β generation, we examined if this apoE antagonist could reduce AD-like pathology in 5XFAD mice, known to develop extensive and aggressive β -amyloid neuropathology. In addition, we examined if mimicking the function of apoE by treatment with ApoEp could accelerate AD-like pathologies in 3XTg-AD mice, where A β seeding might play role in accelerating the progression of tau neuropathology. These mice were treated with ApoEp or 6KApoEp by intraperitoneal injection for 12 weeks and then euthanized, followed by analysis of cerebral A β and tau pathologies. Peripheral treatment of 3XTg-AD mice with ApoEp increased β -amyloid

plaques, as visualized by immunohistochemistry (IHC) with antibody 4G8, in comparison with mice treated with PBS as control (Figure 6A–C). Correspondingly, ApoEp increased levels of soluble A β _{1–40}, ₄₂ and β -CTF, as determined by ELISA and WB analysis (Figure 6D, E), as well as levels of acetylated and phosphorylated tau, as evidenced by IHC and WB analyses (Figure 6F–K and Supplement Figure 6). In contrast, 6KApoEp treatment reduced β -amyloid plaques, A β , and β -CTF levels (Figure 7A–E and Supplement Figure 7) as well as phosphorylated and acetylated tau in 5XFAD mice (Figure 7F–K). In addition, 6KApoEp treatment enhanced synaptogenesis (presynaptic synapsin I and synaptophysin, and postsynaptic PSD95) and reduced neuronal apoptosis (cleaved caspase-3) (Figure 8).

In addition to determination of A β and tau pathologies, the effect of 6KApoEp on AD-like hippocampus-dependent learning and memory impairment was determined in 5XFAD mice utilizing the Y maze, fear conditioning, and novel object recognition tests. Untreated 5XFAD mice exhibited learning and memory impairment compared with non-transgenic control mice, as determined by reduced spontaneous alternation in the Y maze test, which was reversed by 6KApoEp treatment (Figure 9A). 6KApoEp treatment also reduced alternate arm entry returns in both 5XFAD and non-transgenic control mice (Figure 9B). Likewise, learning impairment in 5XFAD mice was exhibited by reduced freezing times during contextual and cued testing after fear conditioning (Figure 9E, F) and reduced ability to discriminate between familiar and novel objects (Figure 9G), which were all reversed upon treatment with 6KApoEp. Notably, 5XFAD mice exhibited some hyperactivity, as shown by enhanced total arm entries in the Y maze, which was reversed by 6KApoEp treatment (Figure 9C). However, 5XFAD and non-transgenic mice, whether untreated or treated with 6KApoEp, exhibited similar levels of anxiety since they spent similar amounts of time in central and peripheral zones of the open field (Figure 9H, I). Overall, these results confirm that while apoE accelerates A β and tau pathologies, antagonizing the effect of apoE by 6KApoEp reduces AD-like pathology, learning impairment, and hyperactivity in an AD mouse model.

DISCUSSION

In the present study, we explored the interaction of apoE with the N-terminal region of APP as a novel therapeutic target for AD. Based on previous studies that suggest that this interaction may enhance A β production (28,29), we initially focused on the apoE LDLR binding domain (residues 133–152, ApoEp). Like apoE, we found that ApoEp also interacts with N-terminal APP and dose-dependently increases A β production (Figures 1, 2). Antibodies against the N-terminal region of APP or the LDLR binding domain of apoE and truncation of the N-terminal domain of APP reduced apoE-mediated A β production (Figures 3, 4). Thus, the apoE LDLR binding domain increases A β production by interacting with the N-terminal APP. Moreover, 6KApoEp, addition of lysine residues to the N-terminal apoE inhibited apoE- and ApoEp- mediated A β production, presumably through decreasing apoE-N-terminal APP interaction, APP cellular membrane trafficking and p44/42 MAPK phosphorylation (Figure 5 and Figure 10).

Based on our findings *in vitro*, we further investigated the potential of 6KApoEp, as a therapeutic agent for the treatment of AD-like pathology in AD mouse models. While

ApoE enhanced A β_{1-40} , 42 levels and β -amyloid plaque as well as acetylated and phosphorylated tau in 3XTg-AD mice (Figure 6), 6KApoE reduced AD-like pathology in 5XFAD mice, reducing β -amyloid plaques, acetylated and phosphorylated tau, apoptosis, and neuroinflammation, while enhancing synaptogenesis and improving hippocampus-dependent learning and memory functions (Figures 7–9 and Supplement Figure 11). In addition, 6KApoE presumably elicited its therapeutic effects without altering cholesterol homeostasis (Supplement Figure 12). Taken together, these findings point to the potential of 6KApoE as a viable therapeutic agent for the treatment of AD pathology and behavioral impairment.

Our study suggests that 6KApoE reduces AD-like pathology by interfering the physical interaction of apoE with N-terminal APP. Based on our findings, we suggest that apoE might have a dual function in the brain, (1) mediating cholesterol transport into the neuron and thereby promoting neuronal proliferation, differentiation, and health (18,19) and (2) binding to N-terminal APP and thereby promoting APP amyloidogenic proteolysis and resultant AD-like pathology (28,29). While the cholesterol transporting role of apoE may function well in the young and healthy brain, the APP proteolytic role of apoE might be expected to be a function of aging and disease. This hypothesis is also based on recent findings that APP has a receptor function like other type 1 transmembrane receptors, which either mediates axon guidance, synaptogenesis, and growth factor signaling or AD pathogenesis, depending on the environment (40–44). For example, over-stimulation of APP by apoE might lead to over-activation of G α protein and APP intracellular domain (AICD), which can be pathogenic. In addition, other factors such as apoE lipidation, glycosylation, and oxidation could play a role in determining how apoE functions. Clearly, the interaction of apoE with N-terminal APP as a therapeutic target for AD should be further explored.

Lastly, our novel apoE antagonist 6KApoE might be expected to be particularly beneficial for AD patients who are apoE4 carriers as well as those carrying apoE3. ApoE3 and apoE4 have a much stronger binding affinity to its receptors compared with apoE2 (45) and individuals carrying the lower affinity apoE2 are protected against AD and have much less accumulation of β -amyloid pathology in the brain as they age (46,47). Clinically, older apoE2 carriers display superior verbal learning abilities, improved recall memory, faster processing of information, and better test performance (47). 6KApoE may specifically counteract the adverse effects of apoE4 by dampening its binding to receptors. A better understanding for the role of apoE isoforms in neuroplasticity and AD as well as their interaction with N-terminal APP and molecular mechanisms may reveal novel approaches for extending brain health span.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

ACKNOWLEDGEMENTS

This work was supported by the NIH (R01AG050253, R01AT007411, and R21AG049477) to Dr. Jun Tan. This work was supported in part by the National Institute on Aging Intramural Research Program. We would like to thank Dr. Li Gan (Gladstone Institutes) for kindly providing us with monoclonal antibodies against acetylated tau

[lysine¹⁷⁴ (K¹⁷⁴) and lysine²⁷⁴ (K²⁷⁴)]. We would like to thank Dr. Song Li for his assistance in the depiction of Figure 10, Drs. Yang Xiang and Song Li for their critical discussion, and Dr. Jared Ehrhart for his technical supports in IHC and IF image analyses. Finally, we will specifically thank Dr. Gobinda Sarkar for his critical discussion in designing apoE LDLR binding domain peptides.

REFERENCES

1. Glenner GG, Wong CW (1984): Alzheimer's disease: initial report of the purification and characterization of a novel cerebrovascular amyloid protein. *Biochem Biophys Res Commun* 120:885–890. [PubMed: 6375662]
2. Grundke-Iqbal I, Iqbal K, Tung YC, Quinlan M, Wisniewski HM, Binder LI (1986): Abnormal phosphorylation of the microtubule-associated protein tau (tau) in Alzheimer cytoskeletal pathology. *Proc Natl Acad Sci U S A* 83:4913–4917. [PubMed: 3088567]
3. Selkoe DJ (2001): Alzheimer's disease: genes, proteins, and therapy. *Physiol Rev* 81:741–766. [PubMed: 11274343]
4. Hardy J, Selkoe DJ (2002): The amyloid hypothesis of Alzheimer's disease: progress and problems on the road to therapeutics. *Science* 297:353–356. [PubMed: 12130773]
5. Schenk DB, Rydel RE, May P, Little S, Panetta J, Lieberburg I, et al. (1995): Therapeutic approaches related to amyloid- β peptide and Alzheimer's disease. *J Med Chem* 38:4141–4154. [PubMed: 7473539]
6. Selkoe DJ, Yamazaki T, Citron M, Podlisny MB, Koo EH, Teplow DB, et al. (1996): The role of APP processing and trafficking pathways in the formation of amyloid β -protein. *Ann N Y Acad Sci* 777:57–64. [PubMed: 8624127]
7. LaFerla FM, Tinkle BT, Bieberich CJ, Haudenschild CC, Jay G (1995): The Alzheimer's A β peptide induces neurodegeneration and apoptotic cell death in transgenic mice. *Nat Genet* 9:21–30. [PubMed: 7704018]
8. Loo DT, Copani A, Pike CJ, Whitemore ER, Walencewicz AJ, Cotman CW (1993): Apoptosis is induced by β -amyloid in cultured central nervous system neurons. *Proc Natl Acad Sci U S A* 90:7951–7955. [PubMed: 8367446]
9. Bradt BM, Kolb WP, Cooper NR (1998): Complement-dependent proinflammatory properties of the Alzheimer's disease β -peptide. *J Exp Med* 188:431–438. [PubMed: 9687521]
10. Suo Z, Tan J, Placzek A, Crawford F, Fang C, Mullan M (1998): Alzheimer's β -amyloid peptides induce inflammatory cascade in human vascular cells: the roles of cytokines and CD40. *Brain Res* 807:110–117. [PubMed: 9757011]
11. Hensley K, Carney JM, Mattson MP, Aksenova M, Harris M, Wu JF, et al. (1994): A model for β -amyloid aggregation and neurotoxicity based on free radical generation by the peptide: relevance to Alzheimer disease. *Proc Natl Acad Sci U S A* 91:3270–3274. [PubMed: 8159737]
12. Murakami K, Irie K, Ohigashi H, Hara H, Nagao M, Shimizu T, et al. (2005): Formation and stabilization model of the 42-mer A β radical: implications for the long-lasting oxidative stress in Alzheimer's disease. *J Am Chem Soc* 127:15168–15174. [PubMed: 16248658]
13. Lee VM, Trojanowski JQ (2006): Progress from Alzheimer's tangles to pathological tau points towards more effective therapies now. *J Alzheimers Dis* 9:257–262. [PubMed: 16914864]
14. Rocca WA, Amaducci LA, Schoenberg BS (1986): Epidemiology of clinically diagnosed Alzheimer's disease. *Ann Neurol* 19:415–424. [PubMed: 3717905]
15. Elias MF, Elias PK, Sullivan LM, Wolf PA, D'Agostino RB (2003): Lower cognitive function in the presence of obesity and hypertension: the Framingham heart study. *Int J Obes Relat Metab Disord* 27:260–268. [PubMed: 12587008]
16. Kapogiannis D, Mattson MP (2011): Disrupted energy metabolism and neuronal circuit dysfunction in cognitive impairment and Alzheimer's disease. *Lancet Neurol* 10:187–198. [PubMed: 21147038]
17. Eisenstein M (2011): Genetics: finding risk factors. *Nature* 475:S20–22. [PubMed: 21760580]
18. Mahley RW (1988): Apolipoprotein E: cholesterol transport protein with expanding role in cell biology. *Science* 240:622–630. [PubMed: 3283935]

19. Mahley RW, Weisgraber KH, Huang Y (2009): Apolipoprotein E: structure determines function, from atherosclerosis to Alzheimer's disease to AIDS. *J Lipid Res* 50 Suppl:S183–188. [PubMed: 19106071]
20. Castellano JM, Kim J, Stewart FR, Jiang H, DeMattos RB, Patterson BW, et al. (2011): Human apoE isoforms differentially regulate brain amyloid- β peptide clearance. *Sci Transl Med* 3:89ra57.
21. Ye S, Huang Y, Mullendorff K, Dong L, Giedt G, Meng EC, et al. (2005): Apolipoprotein (apo) E4 enhances amyloid β peptide production in cultured neuronal cells: apoE structure as a potential therapeutic target. *Proc Natl Acad Sci U S A* 102:18700–18705. [PubMed: 16344478]
22. Harris FM, Brecht WJ, Xu Q, Tesseur I, Kekoni L, Wyss-Coray T, et al. (2003): Carboxyl-terminal-truncated apolipoprotein E4 causes Alzheimer's disease-like neurodegeneration and behavioral deficits in transgenic mice. *Proc Natl Acad Sci U S A* 100:10966–10971. [PubMed: 12939405]
23. Brecht WJ, Harris FM, Chang S, Tesseur I, Yu GQ, Xu Q, et al. (2004): Neuron-specific apolipoprotein E4 proteolysis is associated with increased tau phosphorylation in brains of transgenic mice. *J Neurosci* 24:2527–2534. [PubMed: 15014128]
24. Wang C, Najm R, Xu Q, Jeong DE, Walker D, Balestra ME, et al. (2018): Gain of toxic apolipoprotein E4 effects in human iPSC-derived neurons is ameliorated by a small-molecule structure corrector. *Nat Med* 24:647–657. [PubMed: 29632371]
25. Shi Y, Yamada K, Liddel SA, Smith ST, Zhao L, Luo W, et al. (2017): ApoE4 markedly exacerbates tau-mediated neurodegeneration in a mouse model of tauopathy. *Nature* 549:523–527. [PubMed: 28959956]
26. Uddin MS, Kabir MT, Al Mamun A, Abdel-Daim MM, Barreto GE, Ashraf GM (2018): APOE and Alzheimer's Disease: Evidence Mounts that Targeting APOE4 may Combat Alzheimer's Pathogenesis. *Mol Neurobiol* (Epub ahead of print).
27. Zhao N, Liu CC, Qiao W, Bu G (2018): Apolipoprotein E, Receptors, and Modulation of Alzheimer's Disease. *Biol Psychiatry* 83:347–357. [PubMed: 28434655]
28. Hass S, Fresser F, Kochl S, Beyreuther K, Utermann G, Baier G (1998): Physical interaction of ApoE with amyloid precursor protein independent of the amyloid A β region in vitro. *J Biol Chem* 273:13892–13897. [PubMed: 9593736]
29. Huang YA, Zhou B, Wernig M, Sudhof TC (2017): ApoE2, ApoE3, and ApoE4 Differentially Stimulate APP Transcription and A β Secretion. *Cell* 168:427–441 e421. [PubMed: 28111074]
30. Rezaei-Zadeh K, Shytle D, Sun N, Mori T, Hou H, Jeannot D, et al. (2005): Green tea epigallocatechin-3-gallate (EGCG) modulates amyloid precursor protein cleavage and reduces cerebral amyloidosis in Alzheimer transgenic mice. *J Neurosci* 25:8807–8814. [PubMed: 16177050]
31. Obregon D, Hou HY, Deng J, Giunta B, Tian J, Darlington D, et al. (2012): Soluble amyloid precursor protein-a modulates β -secretase activity and amyloid- β generation. *Nat Commun* 3:777. [PubMed: 22491325]
32. Zhu Y, Hou H, Rezaei-Zadeh K, Giunta B, Ruscin A, Gemma C, et al. (2011): CD45 deficiency drives amyloid- β peptide oligomers and neuronal loss in Alzheimer's disease mice. *J Neurosci* 31:1355–1365. [PubMed: 21273420]
33. Li S, Deng J, Hou H, Tian J, Giunta B, Wang Y, et al. (2014): Specific antibody binding to the APP672–699 region shifts APP processing from α - to β -cleavage. *Cell Death Dis* 5:e1374. [PubMed: 25118934]
34. Sawmiller D, Li S, Mori T, Habib A, Rongo D, Delic V, et al. (2017): Beneficial effects of a pyrroloquinolinequinone-containing dietary formulation on motor deficiency, cognitive decline and mitochondrial dysfunction in a mouse model of Alzheimer's disease. *Heliyon* 3:e00279. [PubMed: 28413833]
35. Sawmiller D, Habib A, Li S, Darlington D, Hou H, Tian J, et al. (2016): Diosmin reduces cerebral A β levels, tau hyperphosphorylation, neuroinflammation, and cognitive impairment in the 3xTg-AD mice. *J Neuroimmunol* 299:98–106. [PubMed: 27725131]
36. Haass C, Selkoe DJ (1993): Cellular processing of beta-amyloid precursor protein and the genesis of amyloid β -peptide. *Cell* 75:1039–1042. [PubMed: 8261505]

37. Habib A, Sawmiller D, Tan J (2017): Restoring Soluble Amyloid Precursor Protein α Functions as a Potential Treatment for Alzheimer's Disease. *J Neurosci Res* 95:973–991. [PubMed: 27531392]
38. Eggert S, Thomas C, Kins S, Hermey G (2018): Trafficking in Alzheimer's Disease: Modulation of APP Transport and Processing by the Transmembrane Proteins LRP1, SorLA, SorCS1c, Sortilin, and Calsyntenin. *Mol Neurobiol* 55:5809–5829. [PubMed: 29079999]
39. Zhang X, Song W (2013): The role of APP and BACE1 trafficking in APP processing and amyloid- β generation. *Alzheimers Res Ther* 5:46. [PubMed: 24103387]
40. Bukhari H, Glotzbach A, Kolbe K, Leonhardt G, Loosse C, Müller T (2017): Small things matter: Implications of APP intracellular domain AICD nuclear signaling in the progression and pathogenesis of Alzheimer's disease. *Prog Neurobiol* 156:189–213. [PubMed: 28587768]
41. Copenhagen PF, Kögel D (2017): Role of APP Interactions with Heterotrimeric G Proteins: Physiological Functions and Pathological Consequences. *Front Mol Neurosci* 10:3. [PubMed: 28197070]
42. Dawkins E, Small DH (2014): Insights into the physiological function of the β -amyloid precursor protein: beyond Alzheimer's disease. *J Neurochem* 129:756–769. [PubMed: 24517464]
43. Deyts C, Thinakaran G, Parent AT (2016): APP Receptor? To Be or Not To Be. *Trends Pharmacol Sci* 37:390–411. [PubMed: 26837733]
44. Sosa LJ, Caceres A, Dupraz S, Oksdath M, Quiroga S, Lorenzo A (2017): The physiological role of the amyloid precursor protein as an adhesion molecule in the developing nervous system. *J Neurochem* 143:11–29. [PubMed: 28677143]
45. Bu G (2009): Apolipoprotein E and its receptors in Alzheimer's disease: pathways, pathogenesis and therapy. *Nat Rev Neurosci* 10:333–344. [PubMed: 19339974]
46. Benjamin R, Leake A, McArthur FK, Ince PG, Candy JM, Edwardson JA, et al. (1994): Protective effect of apoE ϵ 2 in Alzheimer's disease. *Lancet* 344:473.
47. Suri S, Heise V, Trachtenberg AJ, Mackay CE (2013): The forgotten APOE allele: a review of the evidence and suggested mechanisms for the protective effect of APOE ϵ 2. *Neurosci Biobehav Rev* 37:2878–2886. [PubMed: 24183852]
48. Tokuda T, Calero M, Matsubara E, Vidal R, Kumar A, Permanne B, et al. (2000): Lipidation of apolipoprotein E influences its isoform-specific interaction with Alzheimer's amyloid β peptides. *Biochem J* 348 Pt 2:359–365. [PubMed: 10816430]
49. Krul ES, Tang J (1992): Secretion of apolipoprotein E by an astrocytoma cell line. *J Neurosci Res* 32:227–238. [PubMed: 1404494]

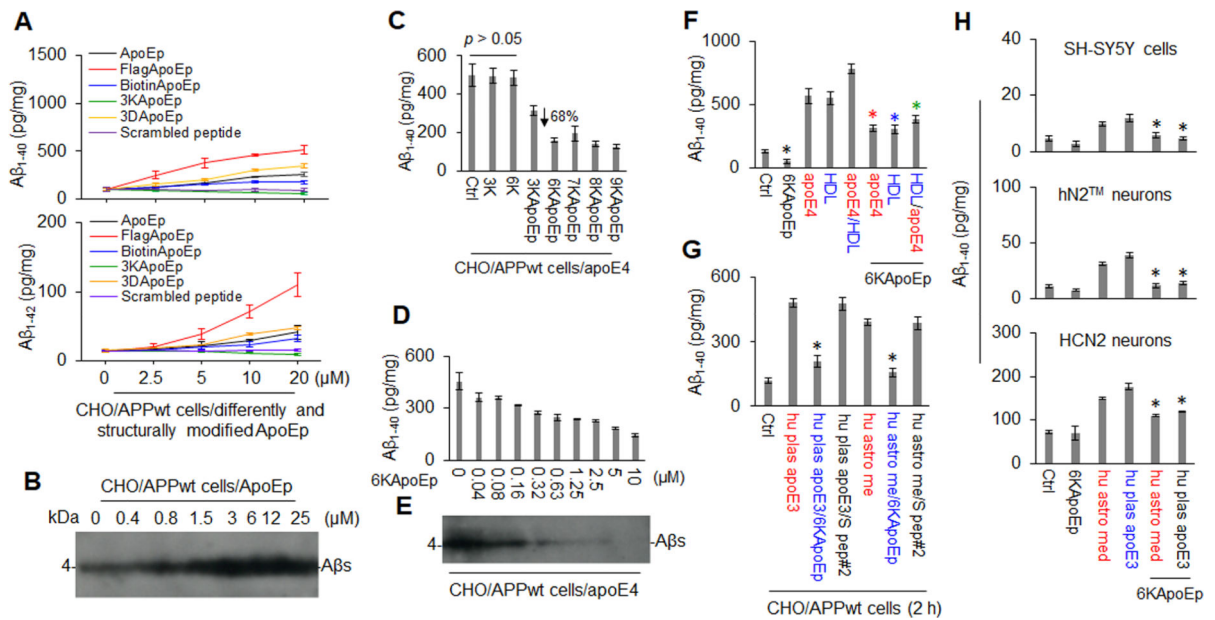


Figure 1. 6KApoEp markedly inhibits human recombinant and lipidated apoE-mediated A β production.

CHO/APPwt cells were treated with apoE LDLR binding domain peptide (aa 133–152, ApoEp), FlagApoEp, BiotinApoEp, 3K (lysine) ApoEp, 3D (aspartate) ApoEp, or scrambled peptide at 0 to 25 μ M, followed after 2 h by analysis of A β levels in conditioned media by A β ELISA (A) and after 16 h by analysis of A β in cell lysates by Western blot (WB) using 82E1 antibody (B). In addition, CHO/APPwt cells were treated with PBS (Ctrl) or 3K (3 lysines), 6K, 3KApoEp, 6KApoEp, 7KApoEp, 8KApoEp, or 9KApoEp at 10 μ M, or 6KApoEp at 0 to 10 μ M, for 15 min, followed by the addition of human recombinant apoE4 (apoE4) at 10 μ g/mL and then A β ELISA (2 h, C, D) and WB (16 h, E). CHO/APPwt cells, SH-SY5Y neuroblastoma cells, hN2TM human neurons and HCN2 cortical neurons were also treated with apoE4, human plasma derived-HDL (HDL) or human plasma-derived apoE3 (hu plas apoE3) at 10 μ g/mL, apoE4 pre-incubated with HDL for 1 h at 37°C (apoE4/HDL), or human astrocyte-derived media (hu astro med) at 1:4 dilution in the absence or presence of 6KApoEp or scrambled peptide (S pep#2) at 10 μ M, followed by A β ELISA (F-H). Pre-incubating with HDL has been previously reported to lipidate apoE (48). Hu astro med was obtained from human astrocytes (CCF-STTG1/ATCC® CRL-1718TM) as described previously (49). ELISA results representative of three independent experiments with each condition duplicated and presented as the mean (\pm s.d.) of A β (pg/mg of total intracellular protein). Asterisk indicates $p < 0.05$ for ApoEp-, FlagApoEp-, BiotinApoEp- or 3DApoEp-mediated A β production, compared with scramble peptide (#1)- or 3KApoEp, as determined by one-way ANOVA (A) and for apoE4-, hu plas apoE3, or hu astro media-mediated A β production in the presence compared with the absence of 6KApoEp as determined by t test (F-H).

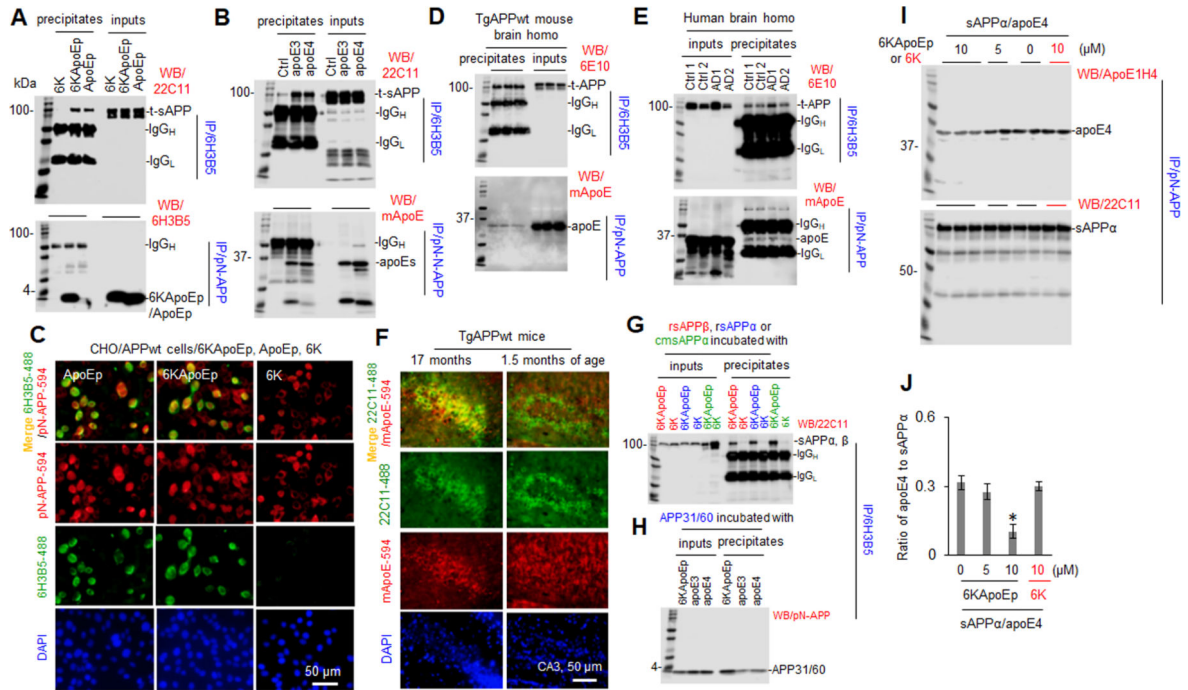


Figure 2. 6KApoEp physically interacts with N-terminal APP.

CHO/APPwt cells were treated with 6KApoEp, ApoEp, or 6K at 10 μ M (A), apoE3 or 4 at 10 μ g/mL, or PBS control (Ctrl, B) for 2 h followed by immunoprecipitation (IP) of the conditioned media with mouse monoclonal anti-apoE LDL binding domain antibody (6H3B5) or rabbit polyclonal anti-N-terminal APP41/66 antibody (pN-APP). Total secreted APP (t-sAPP), 6KApoEp, ApoEp, apoE3, and apoE4 4 in total conditioned media (inputs) and immunoprecipitates (precipitates) was then determined by WB using mouse anti-N-terminal APP antibody (22C11), 6H3B5, and rabbit anti-human apoE antibody (mApoE) (A, B), respectively. CHO/APPwt cells were also plated at 8 well-chambers at 1×10^5 /well for 24 h, treated with 6K, ApoEp, or 6KApoEp at 10 μ M for 2 h, fixed in 4% paraformaldehyde solution and stained with 6H3B5 and pN-APP (C). Alexa Fluor 488 goat anti-mouse IgG was used to detect ApoEp and 6KApoEp (green), while Alexa Fluor 594 donkey anti-rabbit IgG was used to detect N-terminal APP (red). DAPI costaining showed nuclear DNA. Brain tissue homogenates prepared from three TgAPPwt mice (two female/one male), two AD patients (AD1/male, AD2/female) and two normal age-matched control cortices (Ctrl1/male, Ctrl2/female) were IP with 6H3B5 or pN-APP and t-APP and apoE were determined by WB analysis using 6E10 and mApoE, respectively (D, E). Brain tissue sections from aged and young TgAPPwt mice ($n =$ two for each group, one female/one male) were also stained with 22C11 and mApoE at 4°C overnight, followed by staining with Alexa Fluor 488 goat anti-mouse IgG to detect cell surface APP (green) and Alexa Fluor 594 donkey anti-rabbit IgG to detect apoE (red, F). As in vitro confirmation, human recombinant sAPP β (rsAPP β , without A β domain), rsAPP α , CHO/APPwt cell conditioned media-derived sAPP α (cmsAPP α), or APP N-terminal peptide (aa 31–60, APP31/60) at 100 nM was incubated with 6KApoEp, 6K, apoE3, or apoE4 at 100 nM, at 37°C for 1 h followed by IP with 6H3B5. sAPP α / β and APP31/60 in total (inputs) and immunoprecipitates were determined by WB analysis using 22C11 and pN-APP, respectively (G, H). In addition, human sAPP α protein at 100 nM was

also incubated with apoE4 protein at 100 nM in the absence or presence of 6KApoEp or 6K at 5 or 10 μ M for 1 h at 37°C, followed by IP with pN-APP and determination of apoE4 and sAPP α in precipitates by WB analysis using ApoE1H4 and 22C11, respectively (**I**). Band density ratios of apoE4 to total sAPP α was determined by densitometry analysis (**J**). The results shown in **J** panel are representative of two independent experiments with each condition duplicated. Asterisk indicates $p < 0.05$ compared with the controls as determined by t -test. Overall, 6KApoEp, ApoEp, apoE3, and apoE4, but not 6K, were immunoprecipitated and localized with N-terminal APP in vitro and in vivo. ApoE was co-localized with N-terminal APP more in aged compared with young TgAPPwt mouse brains. 6KApoEp, but not 6K, reduced co-immunoprecipitation of sAPP α with apoE4. AD patient and age-matched control cortices provided by Banner Sun Health Research Institute (Sun City, AZ). The results shown in **A-D** panels are representative of two to three independent experiments.

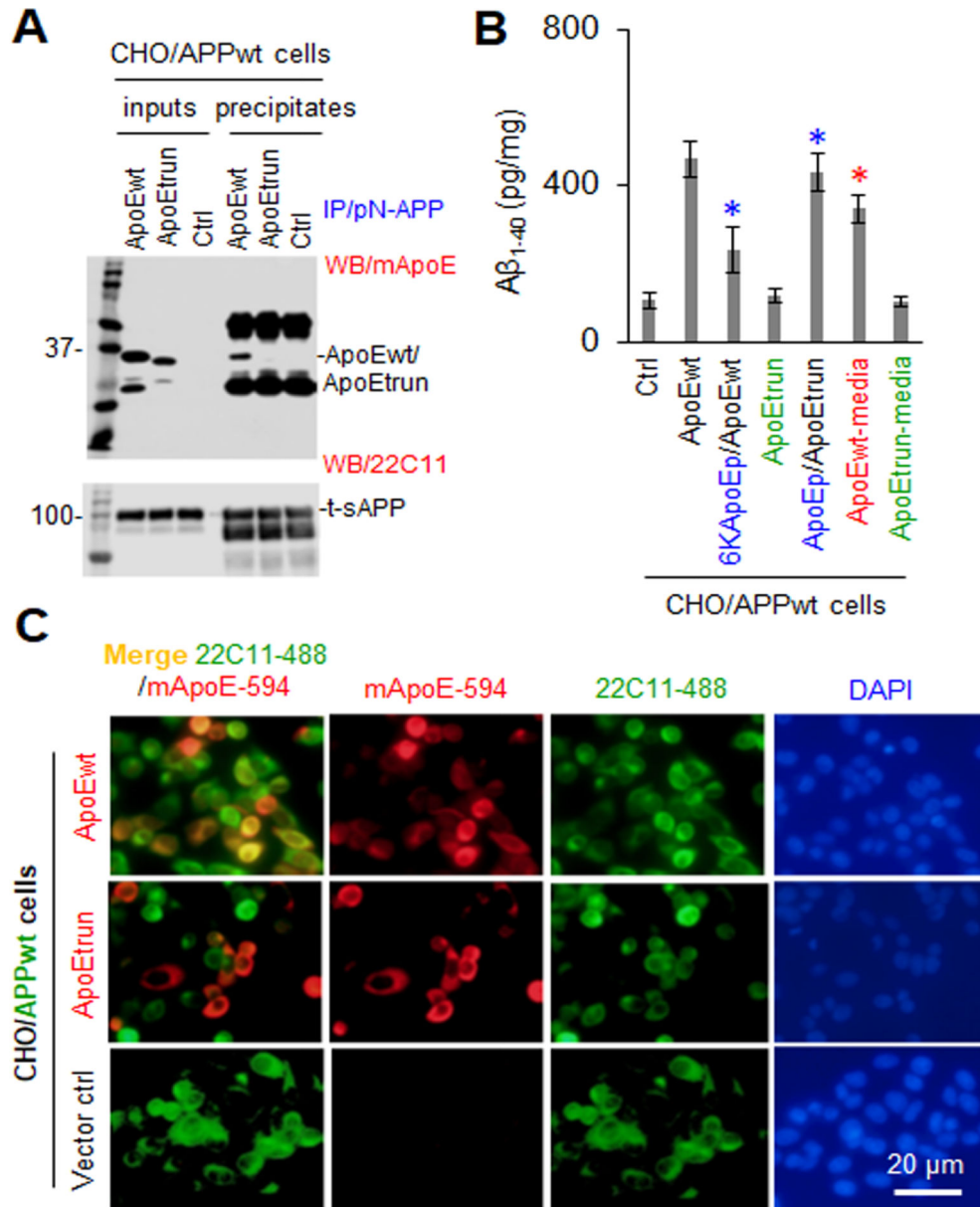


Figure 3. Truncated apoE lacking the LDLR binding domain failed to promote A β production in CHO/APPwt cells.

Conditioned media collected from CHO/APPwt cells co-transiently expressing human wild-type apoE4 (CHO/APPwt/ApoEwt), truncated apoE4 lacking the LDLR binding domain (CHO/APPwt/ApoEtrun), or control vector (CHO/APPwt/Ctrl) were immunoprecipitated with pN-APP and then apoE and sAPP α in total conditioned media (inputs) and immunoprecipitates were determined by WB analysis using mApoE and 22C11, respectively (A). In addition, CHO/APPwt/ApoEwt, CHO/APPwt/ApoEtrun, and CHO/APPwt/Ctrl cells were cultured on 24 well plates at 1×10^5 /well overnight and then treated with 6KApoEp or ApoEp at 10 μ M, and CHO/APPwt/Ctrl cells were treated with conditioned media collected from CHO/APPwt/ApoEwt or CHO/APPwt/ApoEtrun cells, for 2 h followed by A β ELISA (B). Furthermore, CHO/APPwt/ApoEwt, CHO/APPwt/ApoEtrun, or CHO/APPwt/Ctrl cells

were also plated at 8 well-chambers at 1×10^5 /well for 24 h, fixed in 4% paraformaldehyde solution and stained with mouse monoclonal anti-N-terminal APP (22C11) and rabbit monoclonal anti-apoE antibodies (mApoE, **C**). Alexa Fluor 488 goat anti-mouse IgG was used to detect cell surface APP (green), while Alexa Fluor 594 donkey anti-rabbit IgG was used to detect apoE (red). ApoEwt but not ApoEtrun was co-immunoprecipitated (**A**) and co-localized with cell surface APP (**C**, *left panels merged*) and enhanced A β production (**B**), indicating that the LDLR binding domain is necessary for apoE interaction with the N-terminal APP. Notably, 6KApoEp markedly reduced A β production in CHO/APPwt/ApoEwt cells, while ApoEp significantly restored A β production in CHO/APPwt/ApoEtrun cells (**B**). ELISA results are representative of three independent experiments with each condition duplicated and presented as the mean (\pm s.d.) of A β (pg/mg total intracellular protein). Asterisks indicates $p < 0.05$ for A β production in the presence compared with corresponding absence of 6KApoEp or ApoEp or in the presence of CHO/APPwt/ApoEwt compared with CHO/APPwt/ApoEtrun media as determined by *t* test.

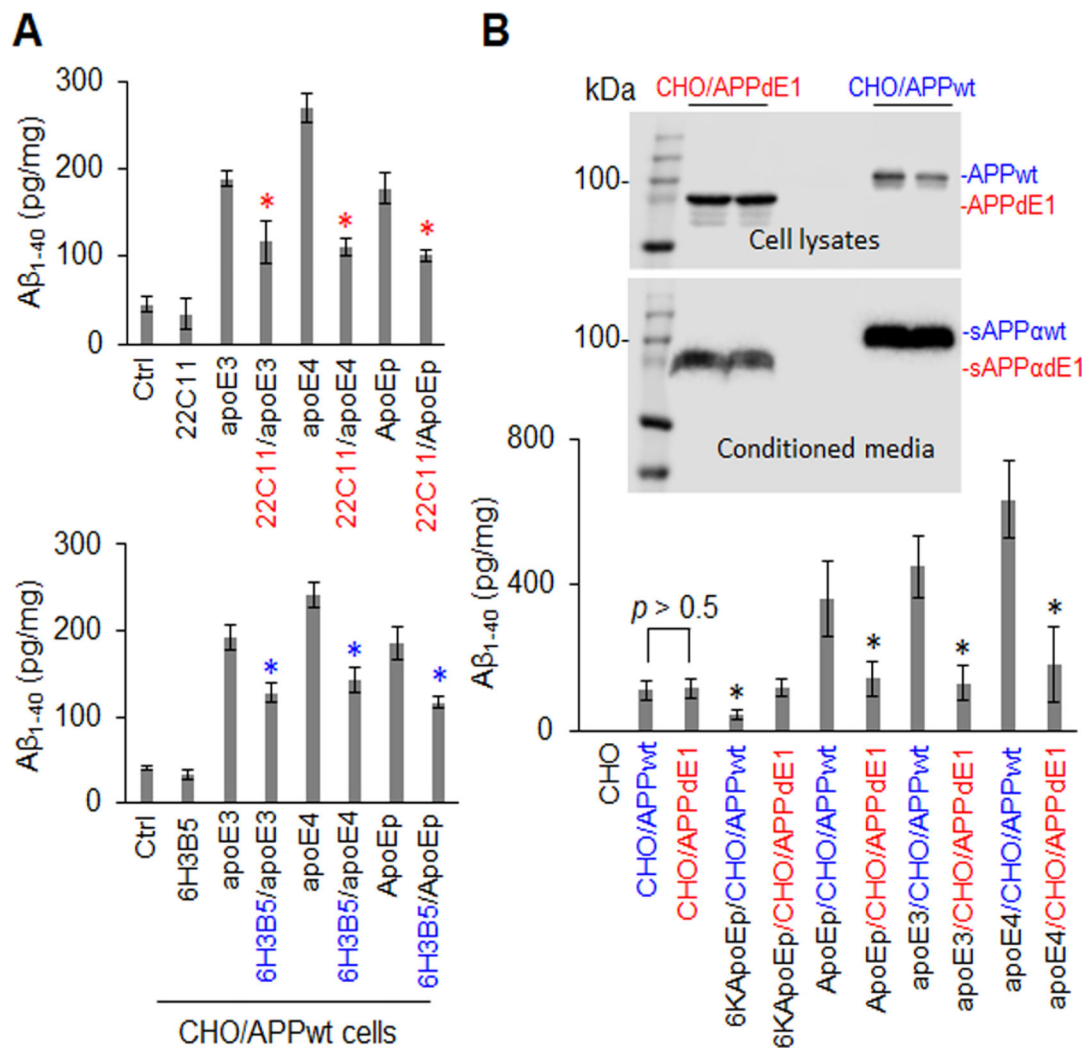


Figure 4. ApoE-promoted A β production was significantly attenuated by specific antibodies against the apoE LDLR binding domain or the N-terminal APP or by expression of N-terminally truncated APP.

CHO/APPwt cells were treated with apoE3 or apoE4 at 10 mg/mL or ApoEp at 5 μ M in the absence or presence of anti-N-terminal APP (22C11) or anti-apoE LDLR binding domain antibody (6H3B5) at 10 μ g/mL for 2 h followed by A β ELISA (a). CHO cells were also transfected with pCMV6 APP695 or pCMV6 E1 depleted APP695, yielding cells expressing wild-type APP695 (CHO/APPwt) or truncated APP695 lacking the N-terminal E1 region (CHO/APPdE1). After 24 h, the cells were treated with 6KApoEp or ApoEp at 10 μ M, or apoE3 or apoE4 at 10 μ g/mL, for 3 h followed by A β ELISA (B, lower panel). APPwt/dE1 in cell lysates and sAPP α wt/dE1 in conditioned media derived from the transfected CHO cells were determined by WB analysis using 6E10 (B, top panels). ApoE3-, apoE4-, and ApoEp-mediated A β production were markedly reduced by 22C11, 6H3B5, and truncation of the dE1 region of APP, indicating that apoE-mediated A β production is mediated by interaction of apoE LDLR binding domain with N-terminal APP. Isotype-matched control IgG failed to reduce apoE3, apoE4, or ApoEp-mediated A β production (data not shown). ELISA results are representative of three independent experiments with each condition

triplicated and presented as the mean (\pm s.d.) of A β (pg/mg of total intracellular protein). Asterisk indicates $p < 0.05$ compared with the corresponding absence of 22C11 or 6H3B5, or corresponding CHO/APPwt cells as determined by one-way ANOVA.

Author Manuscript

Author Manuscript

Author Manuscript

Author Manuscript

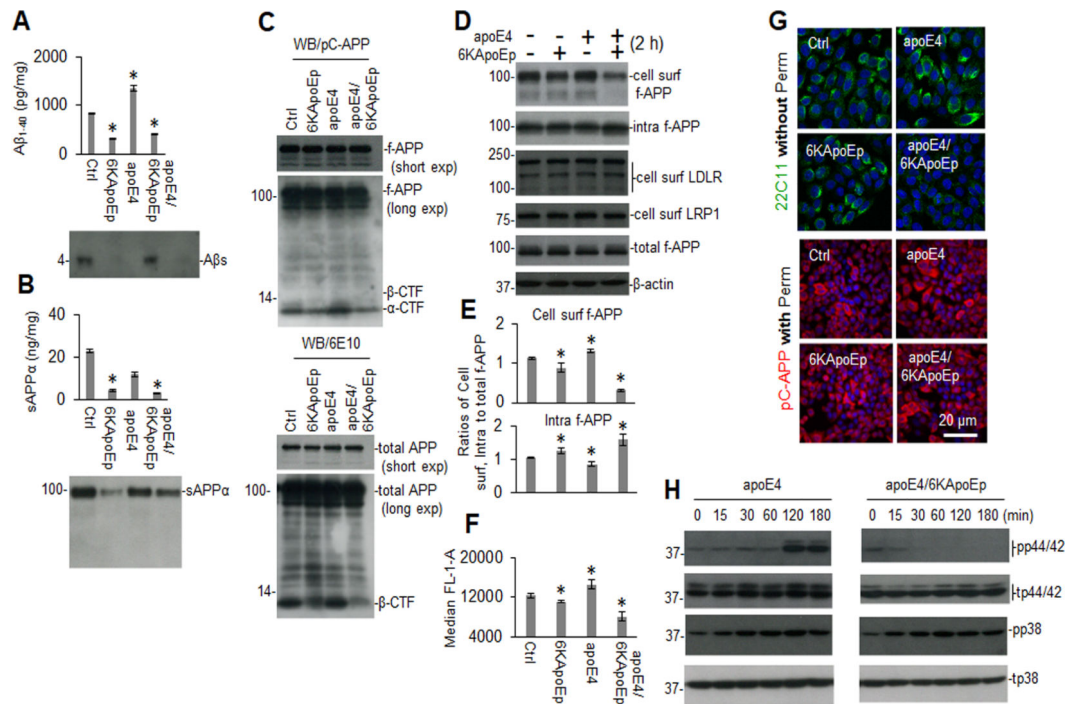
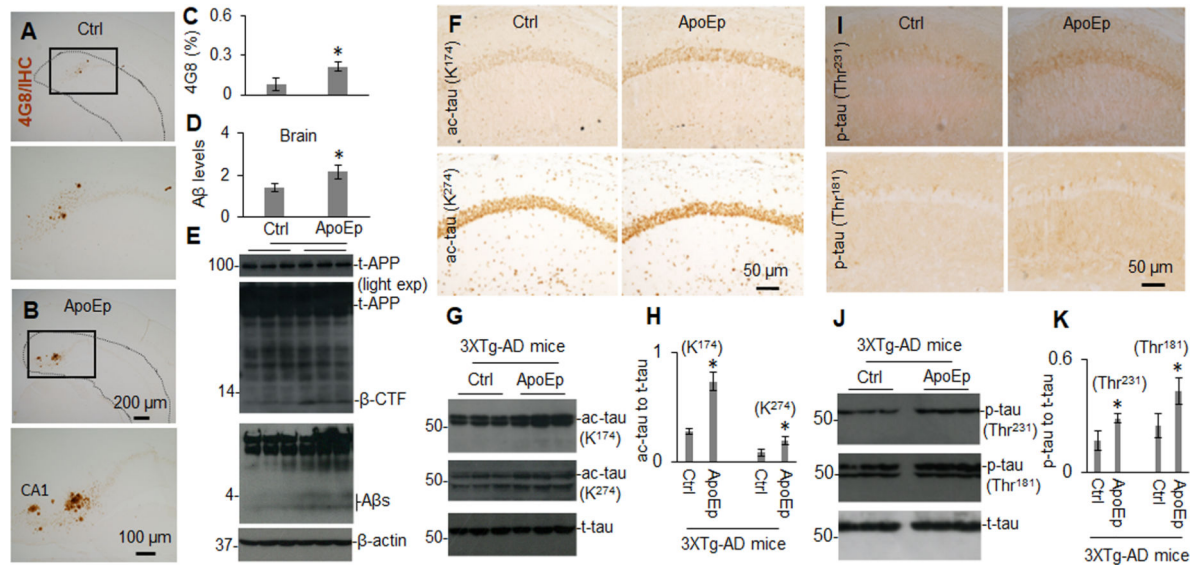
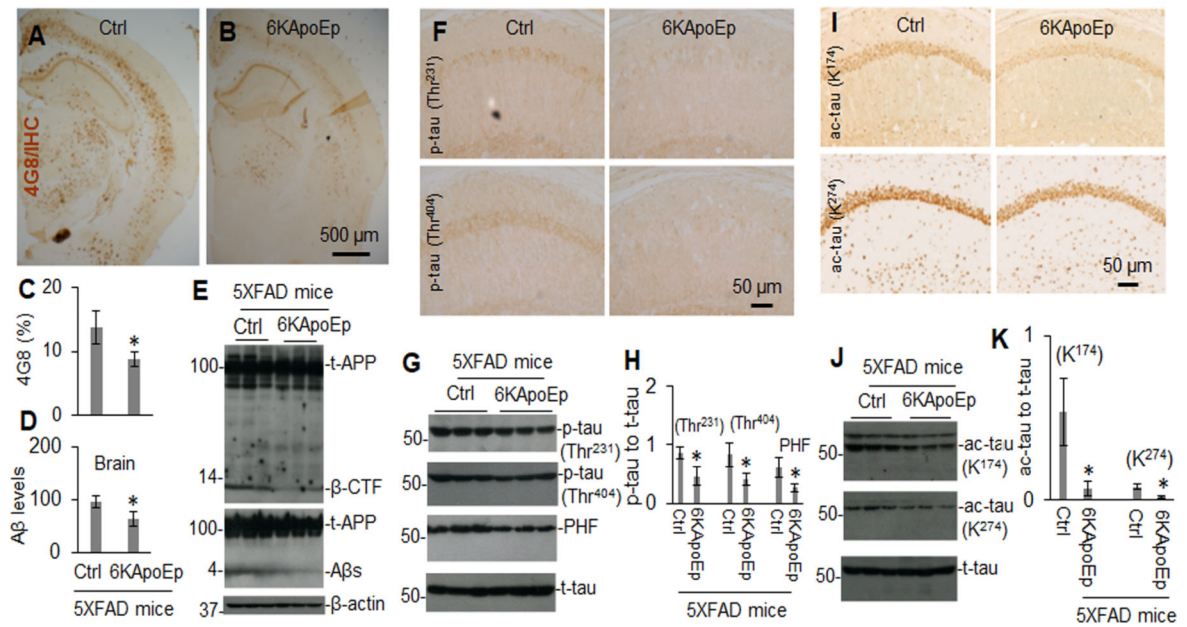


Figure 5. 6KApoEp inhibits APP trafficking to the cell surface and markedly mitigates p44/42 MAPK phosphorylation induced by apoE.

CHO/APPwt cells were treated with PBS control (Ctrl), 6KApoEp at 10 μ M, apoE4 at 10 μ g/mL, or 6KApoEp and apoE4, for 2 h followed by A β and sAPP α ELISA (A, B, top panels) and WB analysis using 82E1 or 2B3 antibody, respectively (A, B, lower panels). In addition, full-length APP (f-APP, short and long exp, exposure), α / β -CTF and total APP in cell lysates were also analyzed by WB using anti-C-terminal APP750/70 (pC-APP; C, top panel) and anti-N-terminal APP antibody (6E10; c, lower panel). Cell lysates were biotinylated and immunoprecipitated using Neutravidin beads and intracellular proteins obtained by Neutravidin depletion (intra) and cell surface proteins obtained by Neutravidin precipitation (cell surf) were analyzed for f-APP, LDLR, and LRP1 by WB using 6E10, anti-LDLR and anti-LRP1 antibodies (D). Band density ratios of cell surf or intra to total f-APP was determined by densitometry analysis (E). Cellular membrane associated full-length APP was analyzed by flow cytometry and presented as the mean (\pm s.d.) median FL-1-A (F). Cultured cells were stained with mouse anti-N-terminal APP (22C11) and rabbit anti-C-terminal APP (pC-APP) primary antibodies after permeabilization (perm, 0.05% Triton X-100, 5 min) or directly (without perm) and observed using an Olympus FV1000 laser scanning confocal microscope (G). Alexa Fluor 594 donkey anti-rabbit and Alexa Fluor 488 goat anti-mouse IgG was used to detect APP. In addition, CHO/APPwt cells were treated with apoE4 at 10 μ g/mL in the absence or presence of 6KApoEp at 10 μ M, for 0 to 180 min followed by determination of total p44/42 (t-p44/42), t-p38 and phosphorylated-p44/p42 (pp44/42/), and pp38 MAPK levels in cell lysates by WB analysis (H). ELISA results are representative of three independent experiments with each condition triplicated and presented as the mean (\pm s.d.). Asterisk indicates $p < 0.05$ compared with Ctrl as determined by t test.





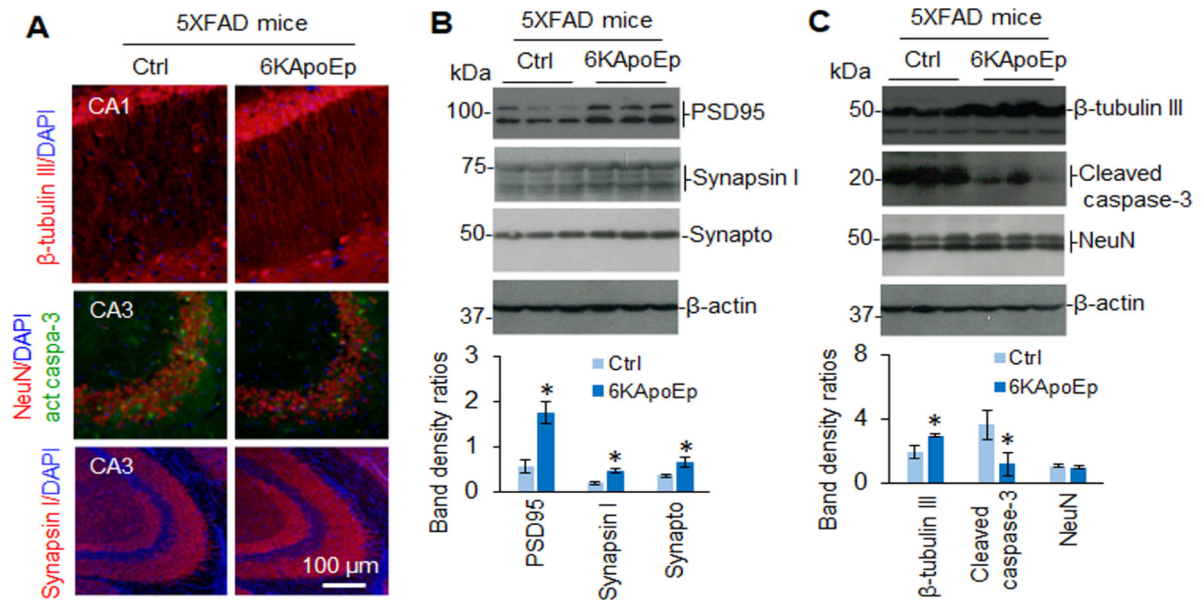


Figure 8. Peripheral administration of 6KApoEp significantly increases pre-synaptic synaptophysin and post-synaptic PSD95 protein expression.

5XFAD mice ($n =$ ten mice, five female/five male) at 6 weeks of age were treated with 6KApoEp at 250 μg/kg or vehicle control (Ctrl, PBS alone) i.p. daily for 12 weeks followed by analysis of neurogenesis in brain tissue coronal sections by IHC and IF. Neuronal (β-tubulin III, NeuN, and synapsin I), presynaptic synaptophysin (synapto), post-synaptic PSD95, and apoptotic cleaved caspase-3 were determined in brain tissue coronal sections by IF staining (A) and in brain homogenates by WB analysis (B, C, upper panels). Band density ratios of each protein to β-actin was determined by densitometry analysis (B, C, lower panels). WB band density ratios represented as the mean (\pm s.d.). Asterisk indicates $p < 0.05$ versus Ctrl.

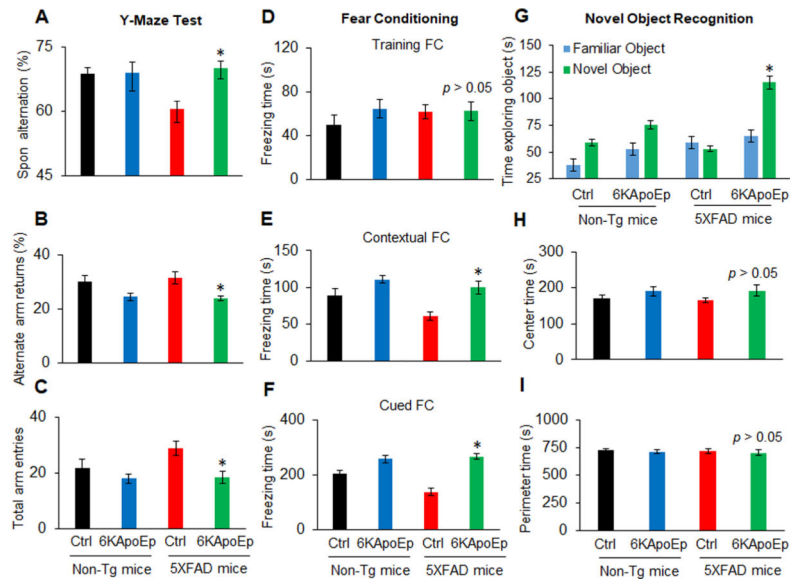


Figure 9. 6KApoEp treatment improves hippocampal-dependent learning and memory in 5XFAD mice.

Cognitive function and anxiety were determined after treatment of 5XFAD and non-transgenic control mice ($n =$ ten mice, five female/five male) with 6KApoEp or vehicle PBS (Ctrl). Spontaneous (Spon) alternation, alternate arm returns and total arm entries were determined for the Y-maze testing (A-C). Total freezing times during the training and contextual and cued testing after training were determined for the fear condition testing (D-F). In addition, total times exploring familiar and novel objects were determined for the novel object recognition testing (G) and total times exploring central and peripheral zones were determined for the open field testing (H, I). 5XFAD mice exhibited impaired cognitive functioning as shown by reduced spontaneous alternation, freezing in the context and cued testing and ability to discriminate between novel and familiar objects compared with non-transgenic mice, which was reversed by 6KApoEp treatment. 5XFAD mice or treatment with 6KApoEp did not exhibit altered anxiety, as determined by time spent in center or perimeter of the open field test. Data are presented as the mean (\pm s.d.). Asterisk indicates $p < 0.05$ as determined by ANOVA.

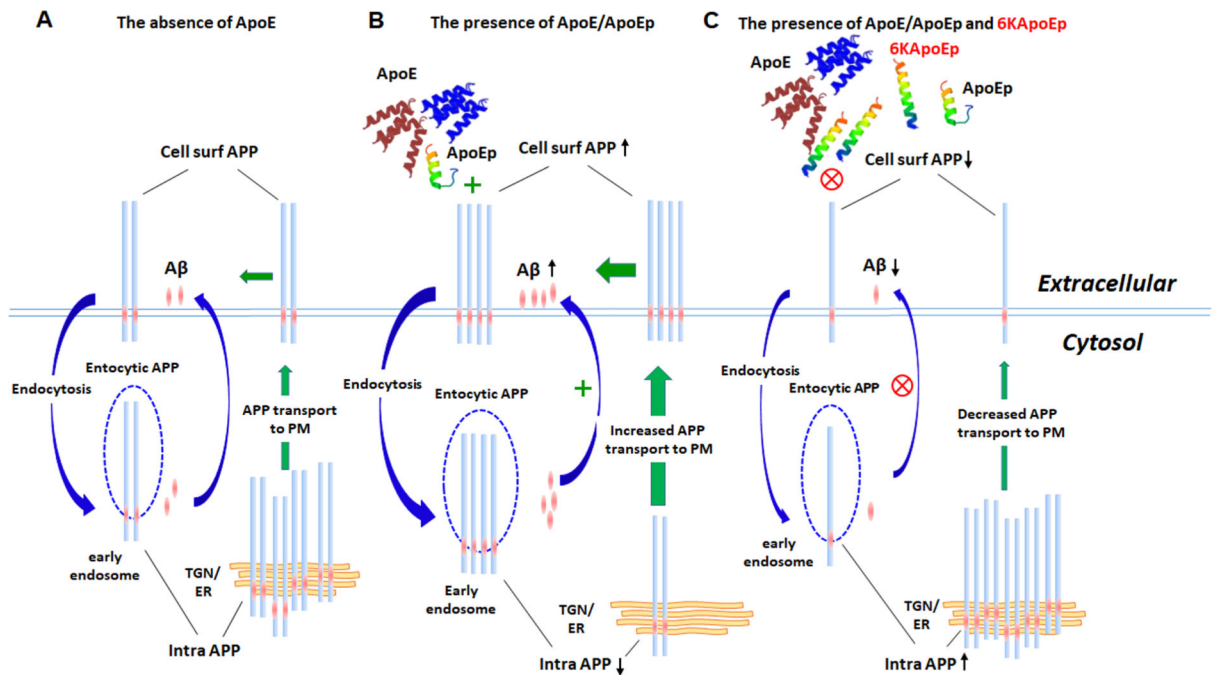


Figure 10. Schematic illustration of a novel apoE antagonist (6KApoEp) functionally blocking apoE or apoE LDLR binding domain peptide (ApoEp) interaction with N-terminal APP and inhibiting A β generation by reducing APP trafficking from trans-Golgi network and endoplasmic reticulum to the plasma membrane.

(A) Under physiological conditions, cell surface APP (Cell surf APP) auto internalizes into the cells *via* clathrin-mediated vesicles but β -site amyloid precursor protein cleaving enzyme 1 (BACE1) internalizes *via* ADP-ribosylation factor 6 (ARF-6) mediated endocytic pathway. APP β -cleavage occurs when lateral compartmentalization of APP and BACE1 meet in the early endosome at low pH and after additional γ -secretase cleavage generate A β , which can be secreted from the cell. In addition, newly synthesized APP travels through the secretory pathway from endoplasmic reticulum (ER) to the plasma membrane (PM) *via* the trans-Golgi network (TGN). (B) ApoE and/or apoE LDLR binding domain peptide (ApoEp) interaction with N-terminal APP increase A β generation by promoting internalization of APP *via* the endocytic pathway. In addition, binding of apoE and/or ApoEp with N-terminal APP markedly increases the trafficking of APP from ER and TGN to the plasma membrane, as evidenced by increased cell surface APP and decreased intracellular APP (Intra APP). (C) 6KApoEp inhibits apoE and/or ApoEp physical interaction with N-terminal APP, which decreases APP endocytosis and trafficking from ER and TGN to the plasma membrane, as evidenced by decreased cell surface and increased intracellular APP, reducing both sAPP α and A β generation.