ApoE influences amyloid- β (A β) clearance despite minimal apoE/A β association in physiological conditions

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Apolipoprotein E gene (APOE) alleles may shift the onset of Alzheimer's disease (AD) through apoE protein isoforms changing the probability of amyloid- β (A β) accumulation. It has been proposed that differential physical interactions of apoE isoforms with soluble A β (sA β) in brain fluids influence the metabolism of A β , providing a mechanism to account for how APOE influences AD risk. In contrast, we provide clear evidence that apoE and sAß interactions occur minimally in solution and in the cerebrospinal fluid of human subjects, producing apoE3 and apoE4 isoforms as assessed by multiple biochemical and analytical techniques. Despite minimal extracellular interactions with sA β in fluid, we find that apoE isoforms regulate the metabolism of sAß by astrocytes and in the interstitial fluid of mice that received apoE infusions during brain Aß microdialysis. We find that a significant portion of apoE and sAβ compete for the low-density lipoprotein receptor-related protein 1 (LRP1)-dependent cellular uptake pathway in astrocytes, providing a mechanism to account for apoE's regulation of $sA\beta$ metabolism despite minimal evidence of direct interactions in extracellular fluids. We propose that apoE influences sAß metabolism not through direct binding to sAß in solution but through its actions with other interacting receptors/transporters and cell surfaces. These results provide an alternative frame work for the mechanistic explanations on how apoE isoforms influence the risk of AD pathogenesis.

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lzheimer's disease (AD), the most common cause of de-Amentia, is a neurodegenerative disease pathologically characterized by extracellular accumulation of amyloid- β (A β), intracellular accumulation of tau, neuronal and synaptic loss, brain atrophy, and inflammation (1-3). There is significant evidence that the accumulation of A β peptides in the brain generates a cascade of events that initiates neurodegeneration and clinical dementia of the Alzheimer's type (4-7). Thus it is important to understand factors that govern A β metabolism and how they lead to A β accumulation. The human apolipoprotein E gene (APOE) encodes the apoE protein, a key protein involved in lipid metabolism (8). It has been established that APOE strongly influences the risk for sporadic, late-onset AD (9-11). The presence of one copy of the APOE E4 allele increases the risk of late-onset AD by about 3.7 times, and the presence of two copies increases the risk by about 12 times (9) (www.alzgene.org/meta.asp?geneID=83), relative to the ɛ3 isoform. ApoE is an exchangeable apolipoprotein with three major polymorphic forms: apoE2 (Cys112, Cys158), apoE3 (Cys112, Arg158), and apoE4 (Arg112, Arg158) (8, 12). The differences in amino acids at these positions are important because they change the charge and structural properties of the protein that influence the functional properties of apoE isoforms. Several mechanisms have been proposed to explain the mechanism by which apoE isoforms affect AD risk. An abundance of evidence from in vitro, animal, and human studies

suggests that a major reason that apoE isoforms affect risk for AD is that they differentially modulate A β clearance and accumulation in the brain. ApoE4 increases A β aggregation and impairs clearance relative to other apoE isoforms (12–21). A recent in vivo study shows that the clearance of soluble A β in the brain interstitial fluid (ISF) depends on the isoform of human apoE expressed (apoE4 < apoE3 ≤ apoE2) (21).

ApoE is a component of many cerebral amyloid deposits, including A_β deposits in AD and prion protein deposits in Creutzfeldt-Jakob disease (22, 23). The identified role of apoE isoforms in influencing AD onset, the association of apoE with extracellular amyloid plaques, and the ability of apoE to affect Aß aggregation and clearance in vivo led us to hypothesize, as have others, that apoE isoforms interact directly with $A\beta$ to influence $A\beta$ metabolism. In the past two decades, numerous in vitro studies have tested this hypothesis, reporting that that apoE interacts with monomeric, fibrillar, and/or oligomeric AB with isoform specificity depending upon the type of apoE (lipidated or nonlipidated) and Aβ used in these studies (10, 24-34; reviewed in ref. 13). It has been suggested that the differential direct interaction of apoE isoforms with Aß influences A_β clearance and/or aggregation in the CNS, thus providing a mechanistic explanation for how apoE isoforms influence the risk of AD pathogenesis.

Significance

It has been proposed that differential physical interactions of apolipoprotein E (apoE) isoforms with soluble amyloid- β (A β) in brain fluids influence the metabolism of A β , providing a major mechanism to account for how *APOE* influences Alzheimer's disease risk. The current study challenges this proposal and clearly shows that lipoproteins containing apoE isoforms are unlikely to play a significant role in A β metabolism by binding directly to A β in physiological fluids such as cerebrospinal fluid or interstitial fluid. Our in vitro and in vivo results suggest that apoE isoforms influence A β metabolism by competing for the same clearance pathways within the brain.

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ApoE, along with apolipoprotein A1, plays an important role as a cholesterol and phospholipid acceptor in reverse cholesterol transport and subsequently in the distribution of cholesterol among different cells (35, 36). Lipoproteins and their transporters/receptors form an efficient transport system to distribute lipids, peptides, and other water-soluble and insoluble molecules. Given the role of apoE in modulating the metabolism of $A\beta$, apoE-containing lipoproteins in the brain also could act as efficient acceptors by binding directly to A β secreted from neurons, thus playing a major role in the clearance or aggregation of AB in an isoform-dependent manner in the CNS. To date, most apoE/Aβ-binding studies in solution have been performed with synthetic preparations of Aß using supraphysiological (high micromolar to millimolar) concentrations and have not assessed the formation of apoE/AB complexes quantitatively. Importantly, the precise amount of direct binding between physiological preparations of apoE and AB occurring in solution and biological fluids, the molecular nature of these interactions, and how the direct interaction of $apoE/A\beta$ influences Aß metabolism have not been detailed. Moreover, most apoE-A β interaction studies were not done in the presence of primary brain-derived cells to test whether apoE could act as an extracellular acceptor for $A\beta$ to alter its metabolism.

In this study, we hypothesized that apoE lipoproteins function as extracellular acceptors/carriers for $\hat{A}\beta$ with isoform specificity to regulate subsequent A^β clearance or aggregation. Our current findings reject this hypothesis, because we show that the association of human apoE particles (reconstituted, cell-secreted, and human CSF) with A β (synthetic, cell-secreted, and CSF) in the presence and absence of cells is minimal in extracellular solutions. Despite minimal interaction in extracellular solution, apoE isoforms influence the clearance of cell-secreted and synthetic sA β in astrocytes and the clearance of extracellular A β in a mouse model of β-amyloidosis during in vivo microdialysis. ApoE isoforms block the uptake and subsequent degradation of A β in astrocytes by competing for the same cellular clearance pathways. Our results suggest that the ability of apoE to influence A β metabolism probably is mediated not through direct binding to $A\beta$ in extracellular solutions but rather through its actions with other interacting cellular receptors, transporters, and/or cellular/membrane surfaces.

Results

Association of apoE with Cell-Secreted Soluble A_β and Synthetic Soluble $A\beta$ in the Presence and Absence of Cells Is Minimal. For apoE- and A\beta-binding studies, we primarily used cell-derived soluble A β (sA β) and a range of concentrations of sA β and apoE that are of physiological relevance in biological fluid. H4 neuroglioma cells expressing human amyloid precursor protein (APP) with the Swedish mutation (APP695 Δ NL) were used to collect cell-derived $A\beta$ secreted into the culture medium. We characterized the H4 APP695 Δ NL cell-derived A β by sizeexclusion chromatography (SEC); the elution profiles show that A β exists as a monomeric species, as was further confirmed by Western blot analysis showing that the A β collected was less than 6 kDa (Fig. 1A). The H4 APP695ΔNL cell-secreted Aβ consists of ~90-95% Aβ40 and 5-10% Aβ42 (Fig. S1), a ratio similar to the A640:42 ratio found in human CSF in normal subjects and in patients with late-onset AD (37, 38). In this study, we used apoE from multiple sources: (i) reconstituted lipidated apoE isoform particles (rapoE2, rapoE3, and rapoE4); (ii) astrocyte-secreted human apoE particles and astrocyte-secreted mouse apoE containing HDL, purified as previously described (39, 40); and (iii) human apoE present in human cerebrospinal fluid (CSF) and isolated by SEC (41). The rapoE isoform particles were prepared using specified ratios of cholesterol and phospholipids and were characterized by nondenaturing gradient gel electrophoresis (Fig. 1B and SI Materials and Methods). The rapoE isoforms were heterogeneous in size with particle diameters ranging from



Fig. 1. Characterization of rapoE isoforms and cell-secreted sA β . (A) H4 APP695 Δ NL cells were incubated with serum-free Opti-MEM medium for 24 h. Then the medium was collected and concentrated fivefold with a 3-kDa cutoff concentrator and subjected to SEC. The SEC-purified A β was separated in a 16.5% tricine gel with nondenaturing PAGE and immunoblotted with 82E1 (anti-A β 1–5) antibody (*Right*). (*B*) rapoE2, rapoE3, and rapoE4 particles (8 µg) were loaded on a 4–20% Tris-glycine gel for native PAGE to assess particle size using a standard containing proteins of specified hydrodynamic radii; particles that were concentrated for in vivo microdialysis experiments in Fig. 6 also were analyzed by native PAGE. (C) [³H]-cholesterol-labeled H4 APP695 Δ NL cells were incubated with 20 µg rapoE3 (1:50:10) and rapoE4 (1:50:10) particles at different time points to assess cholesterol efflux relative to liposomes, expressed as the percentage of radiolabeled cholesterol released into the medium. Differences were assessed using Student's t tests (n = 4; P < 0.05).

10–17 nm, similar to the range of HDL-like particles isolated from human CSF. Although the particle sizes differed slightly by rapoE isoform in the preparation shown in Fig. 1*B*, there were no obvious differences across preparations, and most preparations contained rapoE isoforms of the same size (21). To test the lipid mobilization efficiency and thus the functionality of rapoE isoform particles, the isoforms were tested for their cholesterol efflux efficiency relative to liposomes not containing rapoE. The apoE isoform particles were efficient acceptors, promoting cholesterol efflux in a dose-dependent manner (Fig. 1*C*), whereas liposomes effluxed relatively little cholesterol over the time course. There were no significant isoform-specific effects on the efflux properties as observed in previous studies (Fig. 1*C*) (42).

First, we wanted to test whether apoE could act as an efficient external A β acceptor by binding directly to A β . The molar ratio of Aβ:apoE present in human CSF is 1:50-75 (38, 43), so we performed Aß and apoE-binding studies in this physiological range of molar ratios of A_β:apoE. To study the direct binding of apoE and A β in the presence of cells that secrete A β , H4 APP695 Δ NL cells were labeled with ³H-cholesterol and were incubated with rapoE3 for 3, 6, or 12 h with molar ratios of A β :apoE (as observed at the end of the incubation time) of 1:50, 1:35, and 1:25. Samples collected at different time points were subjected to KBr gradient ultracentrifugation. We examined the density distribution of rapoE3 and $A\beta$ in the KBr gradient. More than 95% of lipidated rapoE3, as determined by ³H cholesterol, was located in the region of the KBr gradient with density less than 1.15 g/cm³ (Fig. 2*A*). Using A β ELISA, the KBr gradient-separated samples were analyzed over the entire range of lipidated and poorly lipidated particles to determine the extent to which AB was associated with rapoE3 particles and whether the association changes with



Fig. 2. KBr density gradient studies showing that the association of apoE with cell-secreted sA β in the presence and absence of cells is minimal. (A) H4 APP695∆NL cells were incubated with 20 µg of lipidated rapoE3 particles and liposome particles. Medium was collected at indicated time points and subjected to ultracentrifugation in a KBr density gradient. Aliquots of the density gradient were used to determine radioactivity and density, rapoE3 (1:50:10) particles were distributed primarily below the density of 1.15 g/cm³. (B) H4 APP695ΔNL cells were incubated with 20 µg of lipidated rapoE3 particles (1:50:10) and liposomes (phospholipid:cholesterol ratio, 50:10) for 3, 6, and 12 h [Aβ:apoE molar ratios of 1:50 (3 h), 1:35 (6 h), and 1:25 (12 h)]. Medium was collected at indicated time points and subjected to ultracentrifugation in a KBr density gradient. Aliquots of the density gradient were used for A β_{1-x} ELISA. (C) (Left) The bar graph represents the percentage of sA β present in the region with a KBr density less than 1.15 g/cm³ (percent A β bound) and in the region with a KBr density greater than 1.15 g/cm³ (percent A β unbound). (*Right*) Time-dependent binding of A β and rapoE3. Differences were assessed using one-way ANOVA followed by a Dunnet post test (n = 5). (D) The bar graph represents the percentage of sA_β present in the region with a KBr density less than 1.15 g/cm³ (percent A_β bound) and in the region with a KBr density greater than 1.15 g/cm³ (percent A β unbound) when H4 APP695ΔNL cells were incubated with 20 µg of lipidated rapoE2, rapoE3, rapoE4 (1:50:10) particles and mouse astrocyte-derived apoE particles for 12 h (A_β:apoE molar ratio, 1:25). Differences were assessed using Student's t tests (n = 5). (E) The bar graph represents the percentage of sA β present in the region with a KBr density less than 1.15 g/cm³ (percent A β bound) and in the region with a KBr density greater than 1.15 g/cm³ (percent A β unbound) when H4 APP695 Δ NL CM-derived A β was incubated with 20 μ g of lipidated rapoE2, rapoE3, rapoE4 particles (in a ratio of 1:50:10) for 12 h (Aβ:apoE molar ratio, 1:25 at 12 h) without cells. Differences were assessed using one-way ANOVA followed by a Dunnet posttest (n = 4).

incubation time. We found that ~95% of A β is unbound at 12 h and is not associated with lipidated rapoE3 (Fig. 2 *B* and *C*). There were small and time-dependent increases in the amount of A β present in the lipoprotein fractions containing rapoE at 3 and 12 h (4.5 vs. 6.1%), but the increase observed was not significant (Fig. 2*C*). Surprisingly, quantification of both A β and apoE in the lipidated apoE fractions (density less than 1.15) indicates a ratio of one molecule of A β (1–40) for every 430 molecules of apoE3 at

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12 h (Fig. S2). Further, the association between cell-secreted A β and human apoE isoform particles did not differ among isoforms, nor did these particles differ from mouse apoE particles in their association with cell-secreted A β (Fig. 2D). These results suggest that lipidated apoE is a poor external acceptor of A β from CNS-derived cells and has an extremely low ability to associate with cell-secreted sA β .

To investigate further the binding ability of apoE with cellsecreted sA β in the absence of cells, the size-exclusion-purified A β from H4 APP695 Δ NL cells was incubated with rapoE3 and rapoE4 particles for 12 h (molar ratio of A β :apoE, 1:25). The collected samples were subjected to KBr gradient ultracentrifugation. We then examined the density distribution of A β in the KBr gradient. Similar to the findings shown in Fig. 2D, ~95–97% of A β was not associated with rapoE3 or rapoE4 particles (Fig. 2E). Further, we used synthetic sA β (molar ratio of A β :apoE, 1:25) in similar studies and found that more than 95% of A β is not associated with rapoE3 or rapoE4 (Fig. S3). These results further suggest that in the presence or absence of cells, in solution, and regardless of the source, A β is an extremely poor binding partner of lipidated apoE isoforms.

When a KBr-gradient ultracentrifugation technique is used, the presence of high salt and the high centrifugal force possibly could dissociate Aß from apoE. To circumvent this potential limitation, we used SEC in physiological salt concentrations to isolate and quantify the A β associated with rapoE isoforms and astrocytesecreted apoE isoforms. The APP695∆NL conditioned media (CM) (Aβ,100 ng/mL) was incubated with rapoE3 (apoE3:phospholipids:cholesterol ratio, 1:50:10), rapoE4 (apoE4:phospholipids: cholesterol ratio, 1:50:10) or astrocyte-secreted, immunopurified apoE3 or apoE4 for 3 or 6 h (molar ratio of Aβ:apoE, 1:25). The collected samples were separated on a size-exclusion column, and the distribution of apoE and $A\beta$ in the fractions was analyzed further. Using sensitive sandwich ELISAs for apoE and A β , we found that apoE and A β were found in separate peaks, and more than 95% of AB was not associated with rapoE3, rapoE4, astrocyte-secreted apoE3, or astrocyte-secreted apoE4 (Fig. 3 A-C). The H4 APP neuroglioma cells secrete mostly monomeric sA β (Fig. 1). Recent studies have suggested that apoE associates with higher-order A β more efficiently than with monomeric A β . To address this question, we used 7PA2 cells that previously have been shown to secrete low- and high-order A β species (44). The 7PA2 CM (Aβ,100 ng/mL) was incubated with astrocyte-secreted apoE3 and apoE4 for 6 h (molar ratio of A β :apoE, 1:20). The collected samples were separated on a size-exclusion column, and the distribution of apoE and A β in the fractions was analyzed. A β was eluted in multiple peaks corresponding to high and lower molecular weights, suggesting that both high- and low-order species of A β are present in the medium (Fig. S44). However, in the presence of astrocyte-secreted apoE3 or apoE4, the SEC elution profile of A β remained the same, and there was no change in peak height or pattern, thus suggesting minimal association of apoE3 and apoE4 with higher- or lower-order A β species secreted by 7PA2 cells (Fig. S4B).

It has been suggested that the ability of apoE to efflux cholesterol is compromised by its putative interaction with A β (45). To investigate this possibility, we performed cholesterol efflux of rapoE isoform particles in the presence of cell-secreted sA β with A β concentrations up to eightfold higher than the A β levels found in CSF (2–15 nM) (38). The rapoE isoforms increased cholesterol efflux in a time-dependent fashion (Fig. 3D). However, there was no compromise in the cholesterol efflux properties of apoE isoforms in the presence of cell-secreted sA β at any concentration tested (Fig. 3D). These results further suggest that apoE isoform functions are not compromised in the presence of physiological or supraphysiological concentrations of A β .

Physical separation processes such as chromatographic or highspeed ultracentrifugation techniques possibly could dissociate



Fig. 3. SEC studies showing that the association of apoE with cell-secreted sA β is minimal. (A and B) H4 APP695 Δ NL CM-derived sA β (100 ng/mL) was incubated with 10 µg of astrocyte-derived apoE3 and apoE4 particles for 6 h (A) or 20 µg of rapoE3 and rapoE4 particles (1:50:10) for 3 or 6 h (B). Medium was subjected to SEC with a Superose 6/10 column. Fractions were analyzed for apoE and A β by ELISAs. (C) The bar graph represents the percentage of sA β present in the region of apoE elution, detected by ELISA (fraction number, 25–32) (percent A β bound) and in the region in which apoE is not detectable with ELISA (fraction number, 34–40) (percent A β unbound). Differences were assessed using *t* tests (*n* = 5). (*D*) [³H]-cholesterol–labeled H4 cells were incubated with 20 µg rapoE3 particles (1:50:10) and the indicated concentrations of A β purified from H4 APP695 Δ NL CM at different time points to assess cholesterol efflux, expressed as the percentage of radiolabeled cholesterol released into the medium. The efflux properties of apoE3 did not change significantly in the presence or absence of A β . Significance was assessed using one-way ANOVA followed by a Dunnet posttest (*n* = 4).

protein complexes if the interactions between proteins are very weak. To circumvent this issue, we used fluorescence correlation spectroscopy (FCS), a highly sensitive single-molecule technique that measures the diffusion time of molecules and does not involve a physical separation process (46). If $A\beta$ binds directly to apoE in solution, the diffusion time of $A\beta$ should increase



Fig. 4. FCS study showing that the association of apoE with synthetic A β is negligible. (*A* and *B*) Diffusion time was obtained from incubation of 100 nM TMR, 100 nM TMR-A β 40 and TMR-A β 42 and 100 nM rapoE2, rapoE3, and rapoE4 particles (molar ratio 1:1) for 2 h or 24 h (n = 4). (*C*) Diffusion time was obtained from incubation of 100 nM TMR-A β 42 and 100 nM astrocyte-derived apoE2, apoE3, and apoE4 for 24 h (n = 4).

significantly because of the change in the size of the $A\beta$ /apoE complex. Synthetic A640 and A642 were labeled with tetramethylrhodamine (TMR) and incubated with rapoE2, rapoE3, and rapoE4 particles for 2 or 24 h. The diffusion time for TMR alone was 24 µs and increased to 62 µs when fluorescein was tagged with A β 40 and A β 42 (Fig. 4*A*). However, no increase in the diffusion time was observed when TMR-labeled A640 and A642 were incubated with rapoE2, rapoE3, and rapoE4 particles even at A_β: apoE molar ratios of 1:1 (Fig. 4A and B). Further, we performed binding studies with astrocyte-secreted apoE2, apoE3, and apoE4 and TMR-labeled Aβ42 (at Aβ42:apoE molar ratios of 1:1 and 5:1) for 24 h. The diffusion time of A β was not increased in the presence of astrocyte-secreted apoE even with high Aß:apoE molar ratios (Fig. 4C). These results further confirm that lipidassociated apoE interacts negligibly or extremely poorly with sAß derived from synthetic or cellular sources in solution.

Association of apoE with sA β in Human apoE3 and apoE4 CSF Is Minimal. Previous studies have reported that $A\beta$ is associated with apoE in human CSF (26, 41). However, these studies were conducted with concentrated CSF (with a 10-kDa cutoff membrane) to 10-30 times their original volumes, and the amount of apoE/Aβ complex formation or association was not determined. To investigate further the putative associations in human CSF, we separated apoE and A β from nonconcentrated CSF (0.5 mL) collected from age-matched APOE E3 homozygous and APOE E4 homozygous human subjects by SEC and analyzed the distribution of apoE and A β in the fractions. ApoE and A β eluted in separate peaks, and more than 95% of $A\beta$ was not coeluted with apoE3 and apoE4 peaks (Fig. 5A), similar to our observations with cell-secreted A β incubated with rapoE particles. Further, there were no significant differences between apoE isoforms in the coelution profile (Fig. 5A). Quantification of A β and apoE in the peak fractions containing both apoE and A β indicates an A β : apoE ratio of 1:3,000-5,000 (Fig. 5A). The recovery of apoE and A β using 0.5 mL of CSF by SEC columns was ~40%. To improve the recovery, we mildly concentrated pooled CSF samples from a 4.0-mL sample to four times the original volume with a 3-kDa cutoff concentrator and performed SEC. The recovery of apoE and A β increased significantly (50–60%), as did the presence of Aβ in the apoE peak (Fig. S5). However, the stoichiometry of apoE:Aβ remained the same in concentrated and nonconcentrated samples (ratio of Aβ:apoE, 1:3,000–5,000) (Fig. S5). To understand whether apoE in CSF binds to the cell-secreted exogenous A β , the nonconcentrated pooled human CSF (1.0 mL) was incubated with cell-secreted sA_β for 6 h. ApoE and A_β were separated by size exclusion, and the distribution of apoE and sA β in the fractions was analyzed. ApoE and sA β eluted in distinct peaks, and more than 95% of A β (exogenous as well as endogenous sA β) was not present in the same fractions as apoE3 and apoE4 (Fig. 5B). The exogenously added cell-secreted sA β eluted along with the endogenous CSF sA β peak. These results strongly suggest that the majority of apoE in physiological solutions (i.e., CSF) is not associated with $A\beta$.

Mechanism of A\beta Clearance Through apoE. Our current results suggest that sA β binds very poorly to apoE in lipoproteins, and therefore the ability of apoE-containing lipoproteins to influence A β clearance may not be mediated through its direct binding to A β in the extracellular fluid of the CNS where A β plaques accumulate. Astrocytes produce the majority of apoE in the CNS, and previous evidence suggests that they are one of the main cell types in the brain that play a central role in the cellular clearance of A β (47–49). To test whether particles containing the apoE isoform influence A β clearance by astrocytes, we cultured immortalized astrocytes derived from apoE2, apoE3, and apoE4 knock-in or apoE-KO mice (apoE2, apoE3, and apoE4 astrocytes secreted 4–6 ng/mL of apoE into culture medium/h; Fig. S6) and incubated them with sA β (cell-secreted H4APP CM, 100 ng/mL sA β) to assess cellular A β



Fig. 5. Association of apoE with sA β in human APOE $\epsilon 3/\epsilon 3$ and APOE $\epsilon 4/\epsilon 4$ CSF is minimal. (A) CSF from APOE $\epsilon 3/\epsilon 3$ and APOE $\epsilon 4/\epsilon 4$ human subjects (0.5 mL) was subjected to SEC. Fractions were analyzed for apoE and A β_{1-x} by ELISAs as described in SI Materials and Methods. Differences were assessed using Student's t tests (n = 4). (B) Pooled CSF from human subjects (0.8 mL) was incubated with A β purified from CM of H4 APF695 Δ NL cells for 6 h and subjected to SEC. Fractions were analyzed for apoE and A β_{1-x} by ELISAs as described in SI Materials and Methods. Differences were assessed using Student's t tests (n = 4).

clearance. After 12-h incubation, the medium was collected, and cells were homogenized in lysis buffer after trypsin treatment to remove extracellular membrane-bound Aß. Aß levels in the medium and cell homogenates were determined by ELISA to examine cellular uptake and clearance. The apoE-KO cells had significantly higher cellular clearance of A β than cells producing apoE2, apoE3, or apoE4 (Fig. 6A). There was a significant increase in the clearance of A β in the presence of apoE2 compared with apoE4; however, there was no significant difference in cellular clearance between apoE2 vs. apoE3, nor was there a difference between apoE3 and apoE4 (Fig. 6A). The A β present in the medium was significantly lower in the presence of apoE-KO astrocytes than in the presence of cells producing apoE3 or apoE4. This result is consistent with the observation of increased cellular levels of $A\beta$ in apoE-KO cells and decreased cellular Aβ in the cells producing apoE3 and apoE4 (Fig. 6B). To confirm our results and assess another cellular source of A β , we used A β CM from 7PA2 cells and incubated the AB with apoE2, apoE3, apoE4, and apoE-KO astrocytes to evaluate AB uptake. We observed similar results, in that the apoE-KO cells had significantly higher cellular clearance of A β than did cells secreting apoE2, apoE3, or apoE4 (EKO > E2 > E3 > E4) (Fig. 6C). In other words, we observed that cells expressing apoE4 had lower clearance of 7PA2 CM-derived A_β than cells expressing apoE2 or apoE3 (Fig. 6C), similar to our observations shown in Fig. 64. Overall, these results suggest that apoE is not required for the cellular clearance of $sA\beta$ in astrocytes, and apoE



Fig. 6. Aβ and apoE share clearance pathways without direct binding. (*A* and *B*) To assess the effect of apoE isoforms on Aβ uptake, apoE2, apoE3, apoE4, and apoE-KO astrocytes were incubated with CM from H4 APP695 Δ NL cells secreting sAβ (sAβ, 100 ng/mL) for 12 h. The amount of Aβ in the cell lysate (A) and medium (*B*) then was assessed by ELISA as described in *Materials and Methods*. (*C*) apoE2, apoE3, apoE4, and apoE-KO–expressing astrocytes were incubated with CM from TPA2 cells (sAβ, 100 ng/mL) for 12 h. The amount of Aβ in the cell lysate then was assessed by ELISA as described in *SI Materials and Methods*. (*D*–*F*) To determine whether apoE and Aβ compete for the same cellular uptake pathways, apoE-KO astrocytes were incubated with CM from H4 APP695 Δ NL cells (sAβ, 100 ng/mL) and indicated concentrations of rapoE2, rapoE3, and rapoE4 (apoE:phospholipid:cholesterol ratio, 1:50:10) for 12 h. The amount of Aβ in the cell lysate (*D*) and in the medium from *D* (*E*) then was assessed by ELISA as described in *SI Materials and Methods*. To determine the role of LRP1 in apoE-dependent Aβ competition, MEF1 (LRP1-expressing cells), and MEF2 (LRP1-KO cells) were incubated with CM from H4 APP695 Δ NL cells (sAβ, 100 ng/mL) and 10⁻⁷ M rapoE3, rapoE3, and rapoE4 (apoE:phospholipid:cholesterol ratio, 1:50:10) for 12 h. The amount of Aβ in the cell lysate (*S*) ApoE3, and E4 (apoE:phospholipid: cholesterol ratio, 1:50:10) for 12 h. The amount of Aβ in the cell lysate (*D*) and in the medium from *D* (*E*) then was assessed by ELISA as described in *SI Materials and Methods*. (*G*) ApoE-KO astrocytes were incubated with CM from H4 APP695 Δ NL cells (sAβ, 100 ng/mL) and 10⁻⁷ M rapoE3, rapoE3, and rapoE4 (apoE:phospholipid:cholesterol ratio, 1:50:10) for 12 h. (*F*) The amount of Aβ in the cell lysate then was assessed by ELISA as described in *SI Materials and Methods*. (*G*) ApoE-KO astrocytes were incubated with CM from H4 APP695 Δ NL cells (sAβ, 100 ng/mL) and 10⁻⁷ M rapoE3, rapoE4, rapoE4 (apoE:ph

isoforms significantly inhibit the uptake of $sA\beta$, either by competing for the same pathways or through other effects on $A\beta$ clearance pathways in astrocytes.

To determine whether apoE and A β compete for the same cellular clearance pathways, we performed a competition assay with lipidated apoE isoforms and sA β in apoE-KO astrocytes. Cellsecreted sA β (H4APP CM; A β , 100 ng/mL) was incubated with apoE-KO astrocytes in the presence of increasing concentrations of rapoE2, rapoE3, and rapoE4 particles (apoE:phospholipid:cholesterol ratio, 1:50:10), and the cellular clearance of A β was determined. ApoE isoform particles significantly inhibited the cellular clearance of A β by astrocytes (Fig. 6D). The inhibition of A β cellular clearance by apoE exhibited saturation at ~100 nM of apoE (Fig. 6D), suggesting that apoE blocked receptor-mediated uptake of A β . The significant increase in A β levels in media with increasing concentrations of apoE isoforms (Fig. 6*E*) is consistent with the decreased A β clearance by the cells in the presence of apoE. ApoE isoform particles were able to block ~50% of the total A β cellular clearance in this setting, suggesting the presence of other pathways of A β uptake independent of externally applied apoE. Further, we tested the role of lipidation of apoE isoforms in inhibiting A β clearance in astrocytes. Cell-secreted sA β (H4APP CM; A β , 100 ng/mL) was incubated with apoE-KO astrocytes in the presence of increasing concentrations of poorly lipidated rapoE2, rapoE3, and rapoE4 particles (apoE:phospholipid:cholesterol ratio, 1:5:1) and the cellular clearance of A β was determined. rapoE isoform particles significantly inhibited the uptake of A β by astrocytes as observed with highly lipidated apoE isoforms (Fig. S7). We

did not observe a difference among apoE isoforms in the competition for blocking A β uptake by astrocytes.

Recent studies have suggested that low-density lipoprotein receptor-related protein 1 (LRP1) and LDL receptor (LDLR) play a major role in the metabolism of A β in brain (47, 50-53). The equal efficiency of apoE2, apoE3, and apoE4 in blocking Aβ uptake/clearance in astrocytes ruled out the possibility that LDLR plays a role in the competitive inhibition of $A\beta$ uptake. It also has been suggested that LDLR has a role in A β metabolism that is independent of apoE (47, 54). To test whether LRP1 has any role in the apoE-sensitive clearance of $sA\beta$ in cells, we used mouse embryonic fibroblasts, MEF1 (LRP1-expressing cells) and MEF2 (LRP1-deficient cells) cells to test the clearance of cellsecreted sA β in the presence of extracellular apoE2, apoE3, and apoE4. The levels of sAβ clearance in MEF1 cells were similar to the levels in astrocytes, and 50% of the sA β clearance was inhibited in the presence of apoE3 and apoE4 (apoE, 100 nM) (Fig. 6F). The LRP1-deficient cells had significantly lower uptake of sA β , as observed in previous studies (50). However, the effect of apoE on the inhibition of $A\beta$ uptake was completely absent in LRP1-deficient cells, suggesting that LRP1 plays a role in apoEdependent competition of sA β uptake in cells (Fig. 6F). To determine further the role of LRP1 in apoE-dependent competition in sA β clearance by astrocytes, we assessed the effects of an LRP1-blocking antibody in apoE-KO astrocytes in the presence and absence of apoE3 and apoE4. The LRP1 antibody significantly decreased the uptake of sA β (~30%) in the absence of apoE (Fig. 6F). However, the effect of apoE (apoE, 100 nM) on Aβ uptake in the presence of the LRP1-blocking antibody did not exceed the maximum inhibition (i.e., ~50% inhibition as observed in Fig. 6D), further suggesting that LRP1 may play a significant role in the apoE-dependent inhibition of cellular uptake of sAß by astrocytes (Fig. 6F).

Activation of endoplasmic reticulum (ER) stress response has been reported in Arginine 61 apoE mice and in macrophages isolated from apoE4 mice (55, 56). We wanted to test whether ER stress is activated in astrocytes expressing the apoE isoform and whether this activation could account for the observed difference in sA β clearance in the apoE isoform (Fig. 6A and C). To test whether apoE isoforms are associated with differential activation of ER stress in our experimental conditions, we cultured immortalized astrocytes derived from apoE2, apoE3, and apoE4 knock-in or apoE-KO mice and incubated them with sAß (cell-secreted H4APP CM,100 ng/mL sA β) for 12 h. After 12-h incubation, cells were homogenized, and Western blot analysis was performed for ER stress markers. ApoE4 cells had significantly higher levels of the ER stress-marker proteins binding immunoglobulin protein (BiP), inositol-requiring enzyme 1 α (IRE1 α), and C/EBP homology protein (CHOP), whereas programmed death-1 (PD-1) levels were similar in all genotypes (Fig. S8). ApoE2, ApoE3, and ApoE-KO cells had similar levels of all ER stress markers tested (Fig. S8). These results suggest that cells expressing apoE4 have higher ER stress, which may affect the general metabolism and specifically the clearance of A β . However, the similar levels of ER stress markers in apoE2, apoE3, and apoE-KO cells does not fit the pattern of $sA\beta$ clearance seen in the presence vs. absence of apoE. This result suggests that ER stress and the inhibition of sA β clearance by apoE in these cells probably are not linked phenomena.

To determine further the effects of increasing apoE concentration on A β in vivo, we assessed A β present in the ISF of human-APP transgenic mice (PDAPP-V717F) by in vivo microdialysis (21, 57) while simultaneously infusing rapoE particles containing apoE2 or apoE4. We characterized the particles before in vivo microdialysis by nondenaturing PAGE (Fig. 1B) before and after the concentration procedure needed for infusion in vivo and found that these particles were essentially the same as those used in our in vitro studies. An equimolar amount of rapoE2 or rapoE4 particles was infused directly around an

ISF AB by ~18.2% compared with a baseline period of steadystate ISF A^β obtained in PDAPP/E2 mice, although this effect did not reach statistical significance (Fig. 7). In contrast, infusion of the same amount of rapoE4 particles in PDAPP/E4 mice significantly increased ISF AB levels by 36.6% above baseline levels. These results further suggest that apoE and A β may compete for similar clearance pathways in vivo, as suggested by our studies in astrocytes (Fig. 6), and that the extent of this competition may depend on the isoform. Discussion The experiments in this study provide multiple lines of evidence that sA β is a very poor binding partner of apoE-containing lipo-

proteins and that the influence of apoE on sA β metabolism may not require direct binding of apoE with sA β in solution. We used native, cell-derived sA β and lipidated apoE from multiple sources to study the direct association of $sA\beta$ and apoE isoforms. Using density-gradient centrifugation and SEC as separation techniques and a nonseparation technique such as FCS, we show both in vitro and in vivo that more than 95% of sAß is not associated with apoE-containing lipoproteins and binds to apoE isoforms in extracellular solution only to a very small extent. Our surprising findings strongly suggest that lipoproteins containing apoE isoforms likely do not play a significant role in sAß metabolism by binding directly to $A\beta$ in physiological fluids such as CSF or ISF. Our in vitro and in vivo results also suggest that apoE isoforms influence sAß metabolism through a different mechanism: by competing for the same clearance pathways within the brain.

implanted 38-kDa microdialysis probe during brain ISF sampling

of freely behaving PDAPP/E2 or PDAPP/E4 mice to examine

the acute effects of rapoE treatment on extracellular A β levels.

Treatment with rapoE2 particles increased steady-state levels of

Several intriguing questions regarding apoE and sA β binding remain and should motivate further study. What is the physiological relevance of the observed small percentage (3-6%) of sA β associated with apoE lipoproteins? The half-life of sA β in the brain ISF of mouse models overexpressing h-APP is $\sim 0.5-1.5$ h (E4 > E2 = E3) and in the CSF of humans is 6 h (21, 58). The turnover rate of A β 40/42 in human CSF is around 8%/h (58, 59). Considering the slow binding kinetics of apoE and sA β (~5% of binding in 12 h) (Fig. 2C) and the faster clearance rate of A β in extracellular fluid, a significant role of apoE/Aß complexes in

В

increase in [eA_{β1-x}]

%

infusior

30

20

PDAPP/E2

PDAPP/E4

following rApoE

А

2.5

-2.0 1.5 1.5 1.0 1.0

0.0

PDAPP/E2

rApoE2

Relative change in [eA_{β1-x}]



PDAPP/E4

rApoE4

mediating clearance differences of sA β in solution is exceedingly improbable. The low percentage of binding could contribute to A β aggregation pathways in the long term. However, no significant apoE isoform-dependent association of A β was observed in apoE-containing lipoproteins in plasma or CSF in our studies or in others (41, 60).

Human CSF contains 2-3 nM of A\beta and 150-300 nM of apoEcontaining lipoproteins (molar ratio: 1-2 molecules of Aß for every 75-150 molecules of apoE). This extracellular location is an excellent space to study apoE and Aß interactions and the subsequent metabolic fates of both molecules. Previous studies from our laboratory and by others have shown that some fraction of A β associates with apoE containing lipoproteins (41, 60). However, several studies suggest that the majority of the endogenous A β is present in the CSF lipoproteins (26, 61). In one study, the authors used a 10-kDa cutoff concentrator to concentrate 30 mL of CSF to 1 mL for SEC separation, but the amount of AB filtered through the 10-kDa cutoff concentrator and the molar ratios of apoE and A β present in the coeluted apoE-A β fractions were unclear from this study (26). A separate study used gradient ultracentrifugation to examine apoE-Aß interactions, finding that 70% of endogenous A β is present in the lipoproteins of CSF and plasma (61). However, by using a similar gradient ultracentrifugation technique, it was reported that 90% of endogenous AB was present along with free protein (26). Our study convincingly shows that a very small percent of $A\beta$ actually is associated with or coelutes in fractions that contain lipoproteins.

What is the relevance of apoE and $sA\beta$ surface-binding studies? The binding affinities of lipid-free and lipidated apoE isoforms and A β by surface plasmon resonance (SPR) spectroscopy or platebased assays have been reported to be in the low nanomolar range; however, the maximal molar apoE:Aß ratios on the chip surface have been estimated as 130:1 (62). Apolipoproteins such as apoAI and apoAII also are reported to bind to $A\beta$ with high affinity and in a range similar to that found for apoE (34, 62). We have repeated the surface affinity binding studies using SPR, finding that the binding of lipidated apoE and A β is in the nanomolar range (Table S1). However, the observed K_d values were higher than previously reported, and lowering the lipidation of apoE isoforms significantly increased the binding affinity to $A\beta 40$ and $A\beta 42$ (Table S1). These results suggests that a small percentage of apoE or AB protein conformations may change during the process of binding to a surface, facilitating the high-affinity binding of apoE and A β on surfaces such as cells or the extracellular matrix. Whether apoE or $sA\beta$ binding to the surface changes the structure of $apoE/sA\beta$ to assist their binding and whether these surface-binding properties of apoE and A β complex formation are physiologically relevant requires further investigation. It will be important to understand whether parenchymal A β interacts with apoE on surfaces to influence A β clearance or aggregation in an isoform-dependent manner.

Our data raise the question of whether apoE must bind to $sA\beta$ to influence its metabolism. Multiple pathways exist in brain for AB clearance, including blood-brain barrier-mediated clearance, cellular uptake, and passive elimination (47, 52, 63-65). Astrocytes represent a major cell type mediating clearance of $A\beta$ from the extracellular space in the brain. Recently, it was shown that LDLR plays an important role in the clearance of sAß in astrocytes and modifies plaque pathology in an APP mouse model independent of apoE, suggesting that sA β can be metabolized by cells by binding directly to lipoprotein receptors such as LDLR without requiring apoE to facilitate this clearance (47, 54). Human and mouse apoE dose-dependently increase Aß levels and plaque burden in APP transgenic mouse models (16, 66, 67). The levels of apoE and $A\beta$ pathology are inversely correlated in mice, suggesting that decreasing the apoE level augments A β clearance and/or reduce A β aggregation. Indeed, we found that acutely increasing apoE2 or apoE4 levels in the ISF of PDAPP/TRE mice resulted in greater retention of $A\beta$ in the ISF. Furthermore, particles containing apoE isoforms decrease the uptake of A β by astrocytes. ApoE and sA β are transported through similar pathways, probably through lipoprotein receptors/transporters, resulting in competition with each other for cellular uptake that is not dependent on a direct interaction.

The isoform specificity for A β uptake is not completely consistent among different sources of A β , including CM of H4APP and 7PA2 cells A β (Fig. 6). A β present in the CM of H4 APP and 7PA2 cells resulted in differences among apoE isoforms in A β clearance, differences that were more robust using CM obtained from 7PA2 cells. This result probably reflects differences in the A β species present in the CM; i.e., H4APP cells produced mostly monomeric A β , whereas the CM of 7PA2 had both monomeric and higher-order species. The role of apoE isoforms in mediating the uptake and degradation of higher-order species of A β (in physiologically relevant concentrations) remains to be determined.

In skin fibroblasts, it was shown that $A\beta$ competes with apoE containing very low-density lipoprotein (VLDL) for uptake into cells in a manner that does not depend on the apoE isoform (68). In the present study, we show that apoE-containing lipoproteins efficiently blocked 50% of sAß uptake into astrocytes with no isoform-specific differences. The saturation of A β uptake in apoE competition studies suggests that the uptake mechanism functions through a receptor(s). The absence of apoE isoform-dependent differences observed in our apoE competition assay (apoE-KO cells incubated exogenously with increasing concentrations of apoE2/E3/E4 and a constant amount of sAB) compared with results obtained with human apoE isoform-expressing astrocytes (only endogenously produced apoE and exogenous A β incubation) likely can be explained by differences in exogenous vs. endogenous apoE uptake/recycling pathways. In peripheral cells, it was observed that apoE intracellular transport pathways are different for apoE that is endogenously produced (and recycled) and for exogenously incubated (and recycled) apoE (69). In brief, our results suggest that apoE isoforms block the clearance of $A\beta$ in astrocytes by competing for the same clearance pathways and/or through defective apoE4 and/or apoE-receptor recycling, as observed in human hepatoma cells and neuronal cells, respectively (70, 71). Consistent with our in vitro studies, we sampled the hippocampal ISF of a mouse model of β -amyloidosis expressing human apoE isoforms (PDAPP/TRE2 and PDAPP/TRE4) after acute infusion of lipidated apoE2 and apoE4. Infusion of either apoE2 or apoE4 particles elevated the concentration of sA β in the brain ISF, suggesting that apoE and A β may compete for the same clearance pathways in vivo, as was suggested by our studies in astrocytes (Fig. 6). However, the extent of this competition is isoform dependent in vivo, whereas we do not observe this dependence under in vitro conditions. The precise reasons for this difference between in vitro and in vivo results are unknown. The difference may result from the contribution of multiple cell types, the blood-brain barrier, and the extracellular matrix present in the brain that are involved in the clearance of $A\beta$ in an apoE-dependent fashion.

Elucidation of receptors that are responsible for the uptake of endogenous and exogenous apoE and sA β will be important to understand further how apoE influences sA β metabolism in the CNS. We have identified LRP1 as a potential candidate in apoEdependent sA β uptake in astrocytes and in MEF cells (Fig. 6 E and F). LRP1 is an endocytic receptor involved in the metabolism of various extracellular ligands (72). Studies have shown that LRP1 is involved in the metabolism of A β in the brain by directly binding to $A\beta$ for cellular uptake/clearance or indirectly through other pathways (73-75). ApoE isoforms interact directly with LRP1 and hence have the potential to compete with $sA\beta$ in the cellular clearance process. The association and dissociation rates of the ligands of LRP1 determine the competition efficiency and subsequent uptake and metabolism of ligands. Hence, apoE isoforms with differential binding and recycling kinetics with LRP1 could differentially influence $A\beta$ clearance in brain. It is important to understand further how apoE isoforms and levels

modulate LRP1 function on the cell surface to affect sAß clearance in vivo.

In conclusion, we present several lines of evidence that do not support the existence of significant direct interactions of apoE isoforms with sA β in CNS fluids. Instead, we propose that the ability of apoE to influence AB clearance or aggregation probably is mediated not through direct binding to $sA\beta$ in solution but rather through its actions with LRP1 and other interacting receptors/transporters. Evidence indicates that decreasing apoE levels in the CNS and modulating the level and activity of apoE receptors and transporters may provide opportunities for developing AD therapeutics.

Materials and Methods

Aβ Clearance Assays. Human apoE isoform-expressing (apoE2, apoE3, and apoE4) and apoE-KO immortalized astrocytes, MEF1 (LRP1-expressing cells, and MEF2 (LRP1-KO) cells were plated in six- or 12-well plates and were grown to confluence. To measure A β uptake, cells first were washed three times with serum-free medium (SFM) followed by the addition of fresh SFM. CM containing H4 APP695 Δ NL cell A β or CM containing 7PA2 cell-secreted A β then was added to the medium at an approximate concentration of 100 ng/mL, and the cells were incubated at 37 °C for 12 h. Then the medium was collected, and the cells were washed twice with PBS. To remove cell surfacebound A β , the cells were incubated with 0.05% trypsin/EDTA for 2 min and then were washed twice with PBS. Cells were removed and homogenized by adding 50 mM Tris Hcl, 150 mM NaCl, 1% Triton X-100 (pH 7.6), and complete protease inhibitor mixture (Roche), and the cell lysates were cleared by centrifugation at 14,000 \times g (Beckman Coulter, Rotor number 301.5) for 12 min. Protein content was measured in all cell lysates using a bicinchoninic acid (BCA) protein assay (Thermo Scientific). The AB and apoE in the media and cell lysates were measured by sandwich ELISAs for A β using m266 (anti-A β 13-28) and biotinylated 3D6 (anti-Ap1-5) antibodies and HJ6.2 and biotinylated HJ6.1 to detect apoE (21, 47, 76). In LRP1-blocking experiments, nonimmune IgG or anti-LRP1 IgG (75 µg/mL) was added 1-1.5 h before apoE and Aβ treatment.

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For apoE/A_β competition assays, apoE-KO cells were plated into 12-well plates, and CM containing either H4 APP695 Δ NL cell-derived A β or 7PA2 cellderived A $\!\beta$ was added to the medium at a concentration of 100 ng/mL. Where indicated, cells were incubated with 10⁻¹⁵ M, 10⁻¹² M, 10⁻¹⁰ M, 10⁻⁹ M, 10^{-8} M, 10^{-7} M, and 10^{-6} M rapoE2, rApoE3, rApoE4 (1:50:10 or 1:5:1) at 37 °C for 12 h. Then the medium was collected, and the cells were washed twice with PBS. To remove cell surface-bound Aß, the cells were incubated with 0.05% trypsin/EDTA for 2 min and then were washed twice with PBS. The cells were removed and were homogenized by 50 mM Tris-Hcl, 150 mM NaCl, 1% Triton X-100 (pH 7.6), and complete protease inhibitor mixture (Roche), and the cell lysates were cleared by centrifugation at 14,000 $\times\,g$ (Beckman Coulter, Rotor number 301.5). Protein content was measured in all cell lysates using a BCA protein assay (Thermo Scientific). The A_β and apoE in the media and cell lysates were measured by sandwich ELISAs m266 (anti-Aβ13-28) and 3D6b (anti-Aβ1-5) and HJ 6.2 and HJ6.1b, respectively (21, 47, 76).

Statistics. All data are presented as mean ±SEM, and different conditions were compared using one-way ANOVA followed by Dunnett's and Bonferroni post hoc test to compare control with treatment groups. The Student's t test was used for comparing conditions with only two groups. Statistical significance (*P < 0.05, **P < 0.01, ***P < 0.001) was determined using GraphPad Prism Software.

Additional materials and methods are available in SI Methods and Materials.

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