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## ApoE promotes the proteolytic degradation of A $\beta$

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### Abstract

Apolipoprotein E is associated with age-related risk for Alzheimer's disease and plays critical roles in A $\beta$  homeostasis. We report that ApoE plays a previously unappreciated role in facilitating the proteolytic clearance of soluble A $\beta$  from the brain. The endolytic degradation of A $\beta$  peptides within microglia by neprilysin and related enzymes is dramatically enhanced by ApoE. Similarly, A $\beta$  degradation extracellularly by insulin degrading enzyme is facilitated by ApoE. The capacity of ApoE to promote A $\beta$  degradation is dependent upon the ApoE isoform and its lipidation status. The enhanced expression of lipidated ApoE, through the activation of liver X receptors, stimulates A $\beta$  degradation. Indeed, aged Tg2576 mice treated with the LXR agonist GW3965 exhibited a dramatic reduction in brain A $\beta$  load. GW3965 treatment also reversed contextual memory deficits. These data demonstrate a novel mechanism through which ApoE facilitates the clearance of A $\beta$  from the brain and suggest that LXR agonists may represent a novel therapy for AD.

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Alzheimer's disease (AD) is characterized by the accumulation and deposition of A $\beta$  peptides within the brain, leading to the perturbation of synaptic function and neuronal loss that typifies the disease (Tanzi and Bertram, 2005). Genetic analysis of familial forms of AD has established the centrality of APP processing and A $\beta$  production to disease pathogenesis. A $\beta$  peptides are normally produced by neurons in the brain and cleared through efflux into the peripheral

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circulation (Zlokovic et al., 2005) and through their degradation by proteinases within the brain (Hardy and Selkoe, 2002).

An isoform of apolipoprotein E, ApoE4, has been shown to confer dramatically increased risk for late onset AD (LOAD) (Roses et al., 1995); however, the basis for this remains one of the major unanswered questions of disease pathogenesis. ApoE plays critical roles in regulating brain A $\beta$  peptide levels, as well as their deposition and clearance (Holtzman, 2001; Zlokovic et al., 2005). Thus, processes that regulate ApoE expression and functional state could affect its ability to influence brain A $\beta$  homeostasis. ApoE is the predominant apolipoprotein in the brain and is synthesized and secreted mainly by astrocytes (but also by microglia (Xu et al., 2006; Xu et al., 2000)) within unilamellar HDL-like particles (Fagan et al., 1999). ApoE is lipidated principally through the action of the ATP-binding cassette transporter ABCA1 (and related transporters) which acts in a variety of cell types to transfer both phospholipids and cholesterol to ApoE (Hirsch-Reinshagen et al., 2004; Wahrle et al., 2004) and in this way ApoE acts to traffic lipids throughout the brain. The lipidation status of ApoE is an important functional parameter, governing its conformation (Fisher and Ryan, 1999), intrinsic stability (Hirsch-Reinshagen et al., 2004; Wahrle et al., 2004), and interactions with membrane receptors (Dergunov et al., 2000; Ladu et al., 2006). Importantly, ApoE binds to A $\beta$  and this, too, is influenced by its lipidation status (Tokuda et al., 2000). Studies with APP transgenic mice have demonstrated ApoE isoform-specific effects on the propensity of A $\beta$  to be deposited in the brain (E4>E3>E2), the nature of the deposits and a gene dosage-related influence on the magnitude of these effects (Holtzman, 2004). ApoE lipidation status is also a significant determinant of whether its interaction with A $\beta$  leads to efflux of the peptides from the brain, or alternatively to the formation of fibrils and their deposition into plaques (Bell et al., 2007; LaDu et al., 1995; Morikawa et al., 2005; Tokuda et al., 2000). Recently, three independent studies have reported that inactivation of the *AbcA1* gene in APP-expressing transgenic mice resulted in reduced levels of ApoE. Remarkably, these mice exhibited a seemingly paradoxical elevation of brain A $\beta$  peptide levels and a doubling of A $\beta$  plaque burden, without a significant effect on A $\beta$  generation (Hirsch-Reinshagen et al., 2005; Koldamova et al., 2005a; Wahrle et al., 2005). The outcomes of these studies strongly suggested lipidated forms of ApoE act to enhance the clearance of A $\beta$  peptides from the brain. The aim of the present study was to establish the mechanism by which ApoE and its lipidation affect A $\beta$  homeostasis.

Liver X Receptors (LXRs) are ligand-activated transcription factors which induce the expression of *ApoE*, *AbcA1* and other genes of lipid metabolism (Beaven and Tontonoz, 2006). LXRs act physiologically as cellular cholesterol sensors and are activated by oxysterols. There are two LXR isoforms, LXR $\alpha$  and LXR $\beta$ , both of which are expressed in the brain (Wang et al., 2002) and their activation results in the rapid and robust increase in the levels of lipidated forms of ApoE (Jiang et al., 2003). Thus, regulation of LXR transcriptional activity provides a mechanism to regulate brain ApoE levels and its lipidation status.

The brain possesses robust intrinsic A $\beta$  clearance mechanisms (Tanzi et al., 2004). A $\beta$  peptides are proteolytically degraded within the brain principally by neprilysin (NEP) (Iwata et al., 2000) and insulin degrading enzyme (IDE, insulysin) (Kurochkin and Goto, 1994). Genetic inactivation of these genes (Farris et al., 2003; Iwata et al., 2001) or administration of inhibitors of these proteinases into the brain results in substantial elevation of A $\beta$  levels in the brain and induction of plaque deposition (Dolev and Michaelson, 2004). Conversely, overexpression of IDE or neprilysin lowered brain A $\beta$  levels and reduced plaque formation (Hemming et al., 2007; Leissring et al., 2003). It has been argued that the predominant mode of A $\beta$ <sub>42</sub> clearance from the brain is through its proteolytic degradation since this peptide is not efficiently exported through the vasculature (Deane et al., 2004). Microglia, the brain's resident macrophages, play an essential role in A $\beta$  clearance through their ability to take up and degrade soluble and fibrillar forms of A $\beta$  (Rogers et al., 2002). Moreover, both microglia and astrocytes secrete proteinases,

including IDE that mediate the degradation of A $\beta$  peptides in the extracellular milieu (Qiu et al., 1998).

Despite considerable effort, the cellular mechanisms through which ApoE influences A $\beta$  clearance remain unresolved. We report that ApoE acts to facilitate the proteolytic degradation of A $\beta$ , a previously unappreciated action of this apolipoprotein. Moreover, the lipidation status of ApoE is a critical determinant of its ability to stimulate A $\beta$  degradation and this finding provides a mechanistic explanation of the increased A $\beta$  levels and deposition observed in APP expressing mice lacking the *ApoE* gene. Importantly, we demonstrate that elevation of lipidated forms of ApoE, through activation of LXRs, results in reduced A $\beta$  peptide and plaque levels in an animal model of AD and is associated with improved contextual memory. Therapeutic agents which increase the abundance of highly lipidated forms of ApoE, including LXR agonists, may attenuate disease pathogenesis and represent a promising strategy for the treatment of AD.

## RESULTS

### Microglia efficiently take up and degrade soluble A $\beta$

We first investigated the contribution of microglia to the clearance of soluble A $\beta$ . We found that these cells rapidly take up soluble forms of A $\beta$  from the extracellular milieu (Figure 1A) through a non-saturable (Figure 1B) macropinocytotic uptake mechanism (Mandrekar et al., submitted for publication). The internalized A $\beta$  is rapidly trafficked to a late endosomal/lysosomal compartment (Figure 1C). Indeed, incubation of BV-2 microglia with soluble A $\beta_{42}$  resulted in the loss of A $\beta$  from the medium, with complete clearance of the peptide within 24 hours (Figure 1D) that is reflective of both of its uptake into microglia as well as degradation by proteinases secreted by these cells into the extracellular milieu. Microglia efficiently degrade internalized soluble A $\beta$  and we observed little or no resecretion of A $\beta$  into the medium (Figure 1E). The intracellular degradation of A $\beta$  by microglia is carried out principally by neprilysin and related proteases whose activity can be inhibited by phosphoramidon and thiorphan (Iwata et al., 2000; Tanzi et al., 2004), because in the presence of these protease inhibitors the degradation of internalized soluble A $\beta$  was dramatically inhibited (Figure 1F).

### HDL apolipoproteins enhance the cellular degradation of soluble A $\beta$ by microglia

ApoE has been postulated to facilitate the clearance of A $\beta$  peptides from the brain and we tested whether ApoE and related apolipoproteins influenced the ability of microglia to degrade soluble A $\beta$ . ApoA-I is another major HDL-associated apolipoprotein both in the periphery and in the brain (Koch et al., 2001). Importantly, it functions and is trafficked in a manner similar to ApoE (Smith et al., 2004; Wang et al., 2001). ApoA-I is also lipidated by ABCA1 (Denis et al., 2004) and avidly binds A $\beta$  peptides (Harr et al., 1996; Koldamova et al., 2001); thus, providing an independent measure of HDL apolipoprotein function. BV-2 microglia (Figure 2A) or primary microglia (Figure 2B) were incubated with soluble A $\beta_{42}$  in the presence of exogenously supplied ApoE or ApoA-I. Coincubation with purified human ApoE or ApoA-I resulted in a dramatic stimulation of A $\beta$  clearance from the microglia. ApoE and ApoA-I elicited a similar reduction in intracellular A $\beta$  levels, demonstrating that the observed effects are reflective of the common actions of these apolipoproteins. The effect of ApoE on microglial A $\beta$  clearance is dose-dependent (Figure 2C). We measured the efficiency with which ApoE stimulated the clearance of A $\beta$  from microglia by monitoring in parallel both the cumulative uptake of A $\beta$  using fluorescently labeled A $\beta_{42}$  and the intracellular levels of the intact A $\beta_{42}$  peptide by ELISA (Figure 2D–F). Fluorescently labeled A $\beta$  is taken up by microglia and degraded, however, the fluorophore is retained within the cells and the total cellular fluorescence is an independent measurement of total A $\beta$  uptake (Mandrekar et al., submitted for publication). We found that the ApoE-stimulated reduction in intracellular A $\beta$  levels was

a consequence of enhanced degradation and was observed at both 3 and 9 hours of incubation. These data demonstrate that the degradation of soluble A $\beta$  within microglia is significantly enhanced by HDL apolipoproteins.

Activation of LXR increased the expression of ApoE and ABCA1 (Burns et al., 2006; Koldamova et al., 2003; Sun et al., 2003), resulting in elevated levels of lipidated ApoE in astrocytes (Supplemental Figure 1A and B). Microglia express both LXR isoforms and respond to LXR agonists (Zelcer et al., 2007). These cells also express and secrete lipidated forms of ApoE (Xu et al., 2000). We tested whether elevation of lipidated ApoE levels would result in enhanced A $\beta$  degradation by microglia by treating these cells with an agonist of LXRs. BV-2 microglia (Figure 3A) or primary microglia (Figure 3B) were pretreated with LXR-specific agonist GW3965 for 18 hours and then incubated with exogenous soluble A $\beta_{42}$  for an additional 3 or 24 hours, respectively. The GW3965-treated microglia exhibited significantly lower levels of intracellular A $\beta_{42}$  due to the enhanced degradation of A $\beta$  in the drug-treated cells compared to vehicle-treated controls. Similar effects were observed using another LXR agonist T0901317 (Figure 3C). LXR activation was verified by the observation of increased expression of its target gene products, ABCA1 and ApoE (Figure 3A and B). Significantly, the combined treatment with both the LXR agonist and provision of exogenous HDL apolipoproteins resulted in a further reduction of intracellular A $\beta$  levels, evaluated either by Western analysis (Figure 3A, B) or by ELISA (Figure 3C). Control experiments established that the uptake of soluble A $\beta$  was not affected by the presence of either exogenous ApoA-I or LXR activation (Supplemental Figure 2A). LXR activation did not affect the expression of neprilysin in microglia (Supplemental Figure 3). These data suggest that LXR activation facilitates the degradation of A $\beta$  peptides and this is associated with induction of LXR target genes including ABCA1 and ApoE and elevation of lipidated ApoE levels.

### **ApoE is essential for efficient intracellular degradation of soluble A $\beta$ by microglia**

To directly assess whether ApoE was essential for efficient intracellular A $\beta$  degradation, wild type or *ApoE*<sup>-/-</sup> microglia were isolated and incubated with soluble A $\beta_{42}$ . Loss of ApoE does not impair soluble A $\beta$  uptake by microglia (Supplemental Figure 2C). Wild type microglia efficiently degrade A $\beta$ , resulting in lower cellular A $\beta$  levels when examined 24 hours later. However, the degradation of soluble A $\beta$  by *ApoE*-deficient microglia was significantly impaired (Figure 4A, B). When the *ApoE* null microglia were provided with exogenous ApoE (Figure 4C), it restored their ability to efficiently degrade soluble A $\beta$ , verifying that intracellular A $\beta$  degradation is facilitated by ApoE. Treatment of *ApoE*-deficient microglia with GW3965 was without effect, demonstrating the effect of LXR agonists on A $\beta$  degradation is dependent on expression of ApoE (Figure 4D).

Human ApoE isoforms confer different risk for LOAD (E4>E3>E2) (Roses et al., 1995). We investigated whether the individual ApoE isoforms differ in their ability to facilitate the proteolytic degradation of soluble A $\beta$  by microglia. We isolated ApoE-containing HDL particles produced by immortalized astrocytes derived from mice in which human *ApoE2*, *e3* or *e4* alleles were knocked into endogenous mouse *ApoE* locus (Morikawa et al., 2005). Primary microglia from *ApoE*<sup>-/-</sup> animals were treated with A $\beta_{42}$  in the absence or the presence of the same amount of purified human ApoE-containing HDL particles. While all isoforms of human ApoE increased clearance of soluble A $\beta$ , ApoE2 exhibited the strongest effect whereas ApoE4 was significantly less efficient in promoting the degradation of soluble A $\beta$  (Figure 4E). Addition of ApoE particles did not significantly influence the uptake of soluble A $\beta$  by *ApoE*<sup>-/-</sup> microglia. These data demonstrate an ApoE isoform-specific promotion of the proteolytic degradation of soluble A $\beta$  that is consistent with the linkage of these isoforms to disease risk.

## The lipidation status of ApoE influences its ability to promote the intracellular degradation of soluble A $\beta$

The degree of ApoE lipidation and HDL particle size is governed primarily by the expression levels of ABCA1 (Supplemental Figure 1A, D, F and G) which, in turn, is directly regulated by LXRs (Supplemental Figure 1A) (Lee and Parks, 2005). We investigated whether ABCA1 affects the ability of ApoE to facilitate the proteolytic degradation of soluble A $\beta$  by microglia. We isolated primary microglia from *Abca1*<sup>-/-</sup>, +/- and wild type mice and compared their ability to degrade exogenously supplied soluble A $\beta$ . The *Abca1* genotype did not significantly influence the uptake of soluble A $\beta$  by microglia (Supplemental Figure 2C). Microglia expressing *Abca1* were able to efficiently degrade soluble A $\beta$ . However, *Abca1* null microglia, which cannot efficiently lipidate ApoE, exhibited significantly higher intracellular A $\beta$  levels, reflecting their impaired ability to degrade the soluble A $\beta$  peptides (Figure 5A). When we induced the expression of ABCA1 and ApoE through the activation of LXRs we observed a significant enhancement of A $\beta$  clearance in wild type cells, however, this effect was substantially diminished in microglia from *Abca1* null animals (Figure 5B, C). LXR agonist treatment resulted in induction of cellular ABCA1 and ApoE levels in a genotype related manner, providing an internal control for LXR action. These data demonstrate that ABCA1 acts to facilitate soluble A $\beta$  clearance through mechanisms that are reliant upon apolipoprotein function and are subject to regulation by LXR agonists, supporting the conclusion that lipidated forms of ApoE act to facilitate intracellular A $\beta$  degradation by microglia.

## ApoE lipidation status influences the extracellular degradation of A $\beta$ by IDE

The proteolytic degradation of A $\beta$  occurs both within cells and in the extracellular milieu of the brain. A $\beta$  peptides are degraded in the extracellular space and interstitial fluid of the brain principally by insulin degrading enzyme (IDE) with minor contributions by other secreted proteinases (Qiu and Folstein, 2006). IDE has been reported to be secreted by microglia and to degrade soluble A $\beta$  (Qiu et al., 1998), but it is unknown whether astrocytes act similarly. We first verified that active IDE is synthesized and secreted by astrocytes (Supplemental Figure 4) and upon addition of soluble A $\beta$  to the astrocyte-conditioned medium the A $\beta$  peptide is efficiently degraded (Figure 6A,B). The degradation of A $\beta$  was inhibited in the presence insulin, a competitive inhibitor of IDE, supporting the notion that IDE (and/or related proteinases whose activity is also inhibited by insulin) was responsible for A $\beta$  proteolysis in the conditioned media.

Next, we tested if ApoE promoted the degradation of soluble A $\beta$  extracellularly by IDE in the same fashion as we observed within microglia and to ascertain if the ABCA1-dependent lipidation was important for this process. Astrocytes are the predominant source of ApoE in the brain and secrete it into the extracellular space, thus we employed conditioned medium from these cells containing both ApoE and IDE and evaluated A $\beta$  degradation. Soluble A $\beta$ <sub>42</sub> was added to astrocyte-conditioned medium and A $\beta$  levels were measured 24 hours later. We found conditioned medium from wild type astrocytes efficiently degraded A $\beta$  (Figure 6C). However, conditioned medium obtained from *Abca1*<sup>-/-</sup> astrocytes that contain poorly lipidated forms of ApoE (Supplemental Figure 1) exhibited significantly higher (four fold) levels of remaining A $\beta$  measured either by Western analysis (Figure 6C) or by ELISA (Figure 6D). The reduced levels of ApoE in *Abca1*<sup>-/-</sup> conditioned media after 24 hours are consistent with previous reports that suggested the stability of ApoE is regulated by its lipidation status (Hirsch-Reinshagen et al., 2004; Wahrle et al., 2004). The reduced degradation of A $\beta$  is not due to differences in the levels of IDE or overall enzymatic activity in the conditioned medium derived from *Abca1*<sup>-/-</sup> compared to wild type astrocytes (Supplemental Figure 4). These data demonstrate that the lipidation status of ApoE regulates its stability and influences extracellular soluble A $\beta$  proteolysis by IDE (and/or related proteinases).

We then employed an *in vitro* assay to verify that the lipidation status of ApoE regulated the degradation of A $\beta$  by IDE. We immunoprecipitated ApoE-containing HDL particles from the conditioned medium of wild type and *ApoE*  $-/-$  astrocytes and then incubated the particles with A $\beta$ <sub>42</sub> in the presence of recombinant IDE for 1 hour. Parallel samples from *ApoE*  $-/-$  astrocytes provide a negative control for this experiment. We found that A $\beta$ , in the presence of nascent highly lipidated forms of ApoE derived from wild type animals, was more efficiently degraded by IDE. In contrast, poorly-lipidated forms of ApoE derived from *ApoE*  $-/-$  mice were much less effective in facilitating soluble A $\beta$  degradation (Figure 6 E). These data demonstrate that the capacity of IDE to degrade A $\beta$  is governed by the lipidation status of ApoE.

### The LXR agonist GW3965 reduces plaque pathology and improves memory in Tg2576 mice

We investigated the effects of LXR agonist treatment on AD-related pathology in aged Tg2576 mice that overexpress hAPP<sup>swE</sup> (Hsiao et al., 1995) and exhibit extensive deposits of A $\beta$  by one year of age (Kawarabayashi et al., 2001). One year old Tg2576 mice were treated orally with the LXR agonist GW3965 for 4 months. We found that GW3965-treated animals exhibited many fewer plaques (Figure 7 A), with an approximate 50% reduction in overall plaque number (Figure 7B) and 67% reduction in plaque load (Figure 7 C) in the hippocampus compared with Tg2576 mice fed a control diet. We found a corresponding approximate 50% reduction in the levels of total A $\beta$ <sub>40</sub> (Figure 7D) and A $\beta$ <sub>42</sub> (Figure 7E) peptides in the brains of the drug-treated animals. Importantly, LXR agonist treatment did not affect APP expression levels, or alter the levels of C-terminal APP fragments (Figure 7F). Thus, drug treatment did not significantly affect APP metabolism or A $\beta$  generation *in vivo*. GW3965 administration resulted in the elevation in brain levels of ApoE and ABCA1, demonstrating the ability of this drug to act within the brain upon LXR target genes (Figure 7F). LXRs have been demonstrated to exhibit robust anti-inflammatory actions in the brain (Zelcer et al., 2007), and we observed that the LXR-mediated reduction in A $\beta$  deposition was accompanied by a corresponding decrease in the number of activated, plaque-associated microglia and interleukin-6 mRNA levels (Supplemental Figure 5A–D). These data demonstrate that LXR activation results in reduced A $\beta$  deposition in an animal model of AD that is postulated to occur through facilitating A $\beta$  clearance from the brain.

Finally, as pre-plaque Tg2576 mice have been shown to display an A $\beta$ -dependent deficit in contextual memory (Jacobsen et al., 2006), we examined the effects of GW3965 on memory processes in these animals. Twenty week-old Tg2576 mice were treated orally for 6 days with GW3965 or vehicle and trained and tested for contextual memory following the contextual fear conditioning procedure. Treatment of Tg2576 mice with GW3965 resulted in significantly improved contextual memory scores that were not significantly different from those observed in the wild-type animals (Figure 8). No effect of GW3965 treatment was observed in wild-type animals. Together, these results demonstrated that LXR activation not only ameliorates amyloid plaque pathology and neuroinflammation, but also improves the memory in an animal model of AD.

## DISCUSSION

Sporadic, late onset AD is the most common form of the disease and the only established genetic risk factor is possession of one or more ApoE4 alleles (Roses, 1996; Tanzi and Bertram, 2005). It remains unclear how ApoE participates in disease pathogenesis. ApoE is implicated as a critical regulator of the propensity of A $\beta$  to be deposited within the brain. Examination of APP expressing mice lacking the murine *ApoE* gene failed to develop compact amyloid plaques, but exhibited elevated levels of A $\beta$  peptides within the brain (Bales et al., 1997). Strikingly, when human ApoE isoforms were expressed in these animals there was a delay in the onset of

plaque deposition and a significant reduction in plaque burden that was isoform-specific (E2>E3>E4) and gene dose-dependent (DeMattos et al., 2004). These data suggested that ApoE influenced A $\beta$  deposition and promoted A $\beta$  clearance, with the ApoE4 isoform being less effective than other isoforms in facilitating removal of these peptides from the brain.

Age-related impairment of A $\beta$  homeostatic mechanisms has been postulated to be a critical determinant of disease risk. A $\beta$  peptides are normally generated at high levels in the brain (approximately 8%/hour) and are cleared at an equivalent rate in both humans and mice (Bateman et al., 2006). Thus, even modest reductions in clearance of soluble A $\beta$  could result in elevated levels of A $\beta$  peptides and ultimately their deposition within the brain. The clearance of A $\beta$  peptides from the brain is accomplished by their proteolytic degradation within the brain and by their efflux through the blood-brain barrier (Tanzi et al., 2004). ApoE has been postulated to facilitate A $\beta$  clearance across the blood-brain barrier, owing to its ability to form complexes with A $\beta$  and to be exported to the peripheral circulation via LRP1 in vascular endothelial cells (Deane et al., 2004). However, a recent quantitative analysis has revealed that ApoE:A $\beta$  complexes are inefficiently cleared through this mechanism and lipidated forms of ApoE are not significantly trafficked by this process (Bell et al., 2007). These findings suggest that ApoE promotes A $\beta$  clearance from the brain principally through intrinsic proteolytic mechanisms. The physiological importance of intrinsic A $\beta$  proteolysis has been demonstrated *in vivo*. Mice in which *Nep* or *Ide* have been genetically inactivated exhibit higher A $\beta$  levels and enhanced plaque deposition in the brain (Farris et al., 2003; Farris et al., 2007; Iwata et al., 2001).

The present study documents a novel mechanism through which ApoE stimulates the degradation of soluble A $\beta$  peptides within the brain. We show that ApoE facilitates the proteolytic degradation of soluble A $\beta$ , both within microglia and in the extracellular milieu, through the action of two distinct classes of proteinases. Significantly, the proteolysis of A $\beta$  is dramatically enhanced in the presence of lipidated apolipoproteins (Figure 9). The transfer of lipids to ApoE is accomplished principally by ABCA1 and several recent studies have provided direct evidence for the functional importance of this process in A $\beta$  homeostasis. Three independent studies in different hAPP mice lacking the *Abca1* gene, have demonstrated that the absence of *Abca1* results in low levels of poorly-lipidated ApoE in the brain. Importantly, *Abca1* inactivation did not affect APP processing or A $\beta$  production. However, these animals exhibited much greater A $\beta$  plaque loads and higher levels of insoluble forms of ApoE, reflective of the codeposition of these molecules (Hirsch-Reinshagen et al., 2005; Koldamova et al., 2005a; Wahrle et al., 2005). Thus, the dominant effect of *Abca1* inactivation *in vivo* is to impair A $\beta$  clearance, without a significant impact on A $\beta$  synthesis.

We demonstrate that the proteolytic degradation of A $\beta$  is stimulated by LXR agonist treatment and is reliant upon the LXR target genes, *ApoE* and *Abca1*. ApoE secreted from wild type glia is fully lipidated (Hirsch-Reinshagen et al., 2004; Wahrle et al., 2004) and enables soluble A $\beta$  to be efficiently degraded by proteinases. Our findings are consistent with the previous report that A $\beta$  present within lipidated ApoE:A $\beta$  complexes isolated from human brains was more susceptible to proteolytic degradation than with purified ApoE, which is poorly lipidated (Russo et al., 1998). We favor a model in which ApoE interacts with A $\beta$  acting to chaperone its proteolysis. However, we cannot exclude the possibility that enhanced intracellular proteolytic degradation of A $\beta$  might arise from HDL-mediated changes in cellular membrane lipid composition, as *Abca1* null microglia retain a partial response to the LXR agonist GW3965. Indeed, LXR has been demonstrated to induce the expression of other cholesterol transporters and apolipoproteins involved in HDL metabolism such as *Abcg1* and *ApoD*, respectively (Zelcer et al., 2007). The ApoE-dependent soluble A $\beta$  clearance mechanisms reported here appear to be distinct from those that occur in adult astrocytes which involve the ApoE receptor LRP1 and deposited forms of A $\beta$  (Koistinaho et al., 2004).

Much of the initial interest in LXR action centered on the roles of these receptors and ABCA1 on neuronal APP processing and A $\beta$  generation. LXR activation in cell culture models yielded conflicting data and was shown to either increase (Fukumoto et al., 2002) or suppress A $\beta$  production (Burns et al., 2006; Koldamova et al., 2003; Sun et al., 2003). Subsequent studies in mice have uniformly found that LXR agonist treatment of wild type (Burns et al., 2006) or young APP-expressing mice (Koldamova et al., 2005b; Lefterov et al., 2007; Riddell et al., 2007) resulted in a significant decrease in brain A $\beta$  levels, and we report similar results in aged Tg2576 mice treated with GW3965. Moreover, we have recently reported that hAPP mice in which either *Lxra* or *Lxr $\beta$*  have been knocked out exhibit a significant increase in A $\beta$  plaque pathology (Zelcer et al., 2007), consistent with the view that LXR activation facilitates A $\beta$  clearance from the brain. These recent studies also found that LXRs inhibited the microglia-mediated inflammatory response and inflammatory gene expression, owing to their ability to functionally inactivate the promoters of proinflammatory genes (Zelcer and Tontonoz, 2006). We have reproduced this finding in aged Tg2576 mice with existing plaque pathology. Thus, LXRs can act to ameliorate AD pathogenesis through their action on both A $\beta$  clearance and suppression of the plaque-related inflammatory response.

The behavioral impairment associated with overexpression of hAPP has been postulated to be due to elevated levels of soluble forms of A $\beta$  within the brain (Comery et al., 2005; Riddell et al., 2007; Walsh and Selkoe, 2004). We found that treatment of Tg2576 mice with the LXR agonist GW3965 resulted in a dramatic improvement in contextual memory and this finding, too, is consistent with the LXR-stimulated A $\beta$  clearance.

In summary, we provide data documenting a previously unappreciated action of ApoE in facilitating the proteolytic clearance of A $\beta$  peptides from the brain. We postulate that ApoE acts both within microglia and in the extracellular space to affect the clearance of A $\beta$  through promoting its proteolysis by at least two distinct classes of proteinases. Importantly, the ApoE4 isoform, which is associated with increased risk for AD, exhibits an impaired ability to promote A $\beta$  proteolysis compared to the ApoE2 and ApoE3 isoforms. There is a remarkable correspondence in the effects of LXR activation and *Abca1* inactivation (or overexpression) in animal models of AD that support an essential role for the lipidation of ApoE in removing soluble A $\beta$  from the brain. The present study provides a mechanistic explanation of both the effects of *Abca1* inactivation (or overexpression (Wahrle et al., 2008)) and the *Lxra* and *Lxr $\beta$*  knockouts (Zelcer et al., 2007) on A $\beta$  plaque pathology in animal models of AD. Our data suggests that therapeutic agents which increase the levels of lipidated forms of ApoE, including LXR agonists, represent a new and potentially efficacious therapy for AD.

## Experimental Procedures

**Animals**—Tg2576 mice or wild type littermates (5 animals/group), 12 months of age, were fed the AIN-76A standard rodent diet alone or containing GW3965 (120 mg/kg; 33 mg/kg/day) *ad libitum* for 4 months. The animals were sacrificed, and the right hemispheres were fixed and processed for immunohistochemical analysis. The left hemispheres were snap-frozen on dry ice and subject to serial extraction of total RNA, DNA and protein. Wild type mice (C57BL/6), *ApoE* null mice (B6129P2-*ApoE*<sup>tm1Unc/J</sup>) and *Abca1* heterozygote mice (DBA/1-*Abca1*<sup>tm1Jdm/J</sup>) were obtained from Jackson Laboratory (Bar Harbor, ME). *Abca1*<sup>+/+</sup>, *+/+* and *-/-* mice were bred from *Abca1* heterozygote mice (DBA/1-*Abca1*<sup>tm1Jdm/J</sup>). Tg2576 mice were maintained by crossing to B6SJLF1/J animals.

**Cell culture**—BV2 microglia were maintained in DMEM containing 2% FBS. Primary microglia and astrocytes were prepared from P0–P3 mice and purified microglia and astrocyte cultures were obtained as previously described (Koenigsknecht and Landreth, 2004; Zander et al., 2002). Immortalized astrocytes derived from human ApoE knockin mice (Morikawa et al.,



2005) were maintained in DMEM containing 10% FBS, 500  $\mu\text{g/ml}$  G418 and 1 mM sodium pyruvate.

**Preparation of A $\beta$  peptides**—A $\beta_{42}$  or A $\beta_{40}$  lyophilized powder (American Peptide Company, Sunnyvale, CA) was dissolved to a final concentration of 1 mg/mL in DMSO and stored at  $-80^{\circ}\text{C}$  until use. Fluorescently labeled peptides were prepared by first dissolving the peptides to a concentration of 2 mmol/L in sterile H<sub>2</sub>O. The A $\beta_{42}$  was then labeled with Cy3 (Amersham Biosciences, Pittsburgh, PA) or Alexa488 fluorophores (Invitrogen, Carlsbad, CA) using manufacturer's protocol. The A $\beta_{42}$  reaction mixture was allowed to fibrillize at  $37^{\circ}\text{C}$  overnight after which unincorporated dyes were removed by ultracentrifugation at  $4^{\circ}\text{C}$ . The pellet was then resuspended in DMSO, sonicated and ultracentrifuged. This was subsequently repeated until the fA $\beta$  was solubilized in DMSO. The supernatant contains the operationally defined "soluble A $\beta$ " which exhibits an electrophoretic mobility corresponding to 4 kDa on SDS-PAGE and is predicted to consist of primarily monomeric and small oligomeric species (Shen and Murphy, 1995).

**Western Blot analysis**—Protein concentrations of cell lysates or brain extracts were measured using the BCA method (Pierce, Rockford, IL). For Western blot analysis of A $\beta$ , 10–20% Tricine gels or 4–15% Bis-Tris gels (Invitrogen, Carlsbad, CA) were used. The following primary antibodies were used: anti-human A $\beta$ , 6E10 (Covance, Dedham, MA); anti-ApoE (Calbiochem, San Diego, CA); anti- $\beta$ -actin; anti- $\beta$ -tubulin (Santa Cruz Biotechnology, Santa Cruz, CA); anti-ABCA1 (Novus Biologicals, Littleton, CO); anti-APP CTF (Millipore, Billerica, MA).

**Immunohistochemistry and image analysis**—Post-fixed hemispheres were sectioned sagittally into 10  $\mu\text{m}$  sections using a cryostat. Sections were mounted, air-dried, and then stored at  $4^{\circ}\text{C}$  until use. For A $\beta$  immunohistochemistry, sections (3 per mouse, about 1.2 mm–1.5mm from the midline, spaced 0.1 mm apart from each other) were incubated in 70% formic acid for 3 min and the endogenous tissue peroxidase activity was quenched by incubation with 1% peroxide in methanol for 10 min. Sections were then microwaved in distilled water for 3 min, and then incubated with blocking solution (5% normal goat serum and 0.1% Triton X-100 in PBS) for 1 hour. Sections were incubated with primary antibody in the blocking solution overnight at  $4^{\circ}\text{C}$ . The antigens were detected by secondary antibodies using standard ABC-DAB methods. Sections were counterstained with hematoxylin. The 6E10 antibody against human A $\beta$  was used to stain A $\beta$  plaques. Images were analyzed using Image Pro-Plus software (Media Cybernetics, Silver Spring, MD).

**A $\beta$  ELISAs**—A $\beta_{40}$  and A $\beta_{42}$  ELISAs were performed using commercial kits (Invitrogen, Carlsbad, California) following the manufacturer's instructions. Alternatively, A $\beta_{1-x}$ , A $\beta_{40}$  or A $\beta_{42}$  ELISAs were performed using 6E10 as the capture antibody and monoclonal 4G8, anti-A $\beta_{40}$  or anti-A $\beta_{42}$  HRP-conjugated antibodies (Covance, Dedham, MA) as the detection antibodies. Synthetic A $\beta_{40}$  or A $\beta_{42}$  were used to generate a standard curve for each experiment. The plates were developed using TMB substrate kit (Pierce, Rockford, IL) and the reaction was stopped by addition of equal volume of 1 M HCl. The results were read using a Spectramax colorimetric plate reader (Molecular Devices, Sunnyvale, CA).

**Conditioned media harvesting**—Confluent primary astrocytes were washed twice with PBS (pH7.4) and incubated with fresh serum-free F12-DMEM media for 24 hours. Immortalized astrocytes were washed twice with PBS (pH7.4) and incubated with fresh serum-free DMEM supplemented with 1 mM sodium pyruvate. Conditioned media were collected and sterilized by filtering through 0.22  $\mu\text{m}$  filters.

**Flow cytometry**—BV2 or primary microglia were incubated with 2  $\mu\text{g/ml}$  Alexa488 labeled  $\text{A}\beta_{42}$  in serum-free DMEM or F12 DMEM, respectively, in the presence of 10  $\mu\text{g/ml}$  BSA for 3 hours, washed extensively with PBS and fixed with 4% paraformaldehyde for 5 min. Cells were washed again with PBS and collected for flow cytometry using Beckman-Coulter XL flow cytometer. The total amount of  $\text{A}\beta$  internalized was determined by fluorescent intensity of Alexa488 as measured by flow cytometry.

**Live cell imaging**—For live cell imaging, microglial BV-2 cells were plated overnight in DMEM containing 2% FBS on Delta T tissue culture plates. Cells were incubated with 2  $\mu\text{g/ml}$  Cy3- $\text{A}\beta$  and 100 nmol/L LysoTracker Green DND-26 (Invitrogen, Carlsbad, CA) in serum-free DMEM containing 10  $\mu\text{g/ml}$  BSA and imaged using a Zeiss LSM 510 confocal microscope.

**Intracellular  $\text{A}\beta$  clearance**—BV2 or primary mouse microglia cells were incubated with DMSO or 1  $\mu\text{mol/L}$  GW3965 for 18 hours at 37°C. Cells were then treated with 2  $\mu\text{g/ml}$  soluble  $\text{A}\beta_{42}$  in serum-free medium containing 10  $\mu\text{g/ml}$  BSA for 3 hours or 24 hours respectively in the presence or absence of drug. For experiments using apolipoproteins, purified human HDL ApoA-I (Sigma, Saint Louis, MO), purified human plasma ApoE (rPeptide, Athens, GA), native ApoE2, ApoE3, or ApoE4 particles were applied at the same time as soluble  $\text{A}\beta_{42}$ .  $\text{A}\beta_{42}$  levels in the cell lysates were determined by immunoblotting with the anti- $\text{A}\beta$  antibody 6E10. Briefly, cells were extensively washed with PBS to ensure removing  $\text{A}\beta$  which is attached to the cell surface. Cells were then lysed in ice-cold RIPA buffer (Upstate Biotechnology, Lake Placid, NY), sonicated briefly, then collected by centrifugation at 13,000 rpm at 4 °C for 15 min. The samples were resolved by 4–15% Bis-Tris SDS-PAGE. ABCA1, ApoE (for positive controls) and  $\beta$ -actin or  $\beta$ -tubulin levels (for loading controls) in the cell lysates were measured by Western blot. For ELISA measurements, cells were treated similarly, washed with PBS and lysed in 1% SDS.  $\text{A}\beta_{42}$  levels were measured using ELISA and normalized to total protein.

**$\text{A}\beta$  degradation in astrocyte-conditioned medium**—Conditioned media from *Abca1* +/+, +/- and -/- astrocyte cultures were incubated with 1  $\mu\text{g/ml}$   $\text{A}\beta_{42}$  for 0 or 24 hours.  $\text{A}\beta_{42}$  in the media was assessed using Western analysis or ELISA.

**In vitro IDE assay**—ApoE-containing HDL particles were immunoprecipitated from the conditioned medium using the anti-ApoE antibody. Recombinant (500 ng/ml) IDE (R&D Systems, Minneapolis, MN) was incubated with 2  $\mu\text{g/ml}$   $\text{A}\beta_{42}$  for 1 hour in the presence of ApoE containing HDL particles. The samples were resolved by 4–15% Bis-Tris SDS-PAGE. The levels of IDE, ApoE and  $\text{A}\beta$  were monitored using Western analysis.

**Contextual Fear Conditioning Studies**—Following six days of treatment by oral gavage with 50 mg/kg/day GW3965 or vehicle, 20 week old Tg2576 mice (n=11/genotype/treatment) were trained and tested on two consecutive days as described previously (Comery et al., 2005).

**Statistical analysis**—To compare differences between the experimental groups, two-tailed *t*-test or one-way analysis of variance (ANOVA) followed by the Tukey-Kramer multiple comparison test was performed using GraphPad Prism software (GraphPad Software, San Diego, CA). Contextual memory was analyzed using a two-way ANOVA and post hoc pairwise comparison made using SAS Statistical Software (SAS Institute, Cary, NC).

## Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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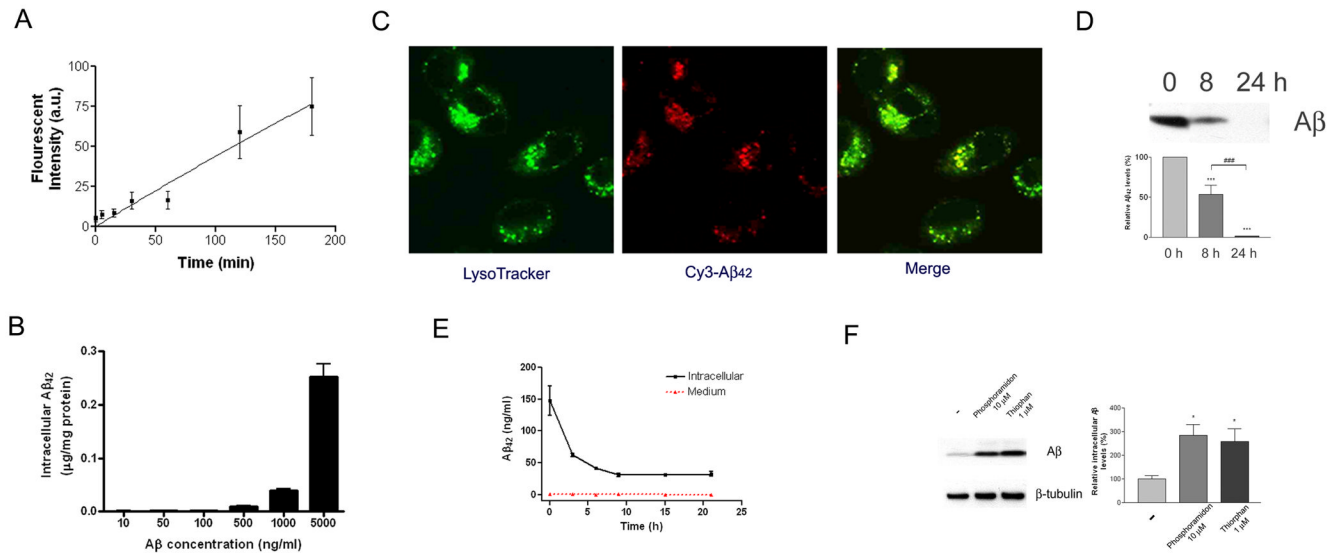
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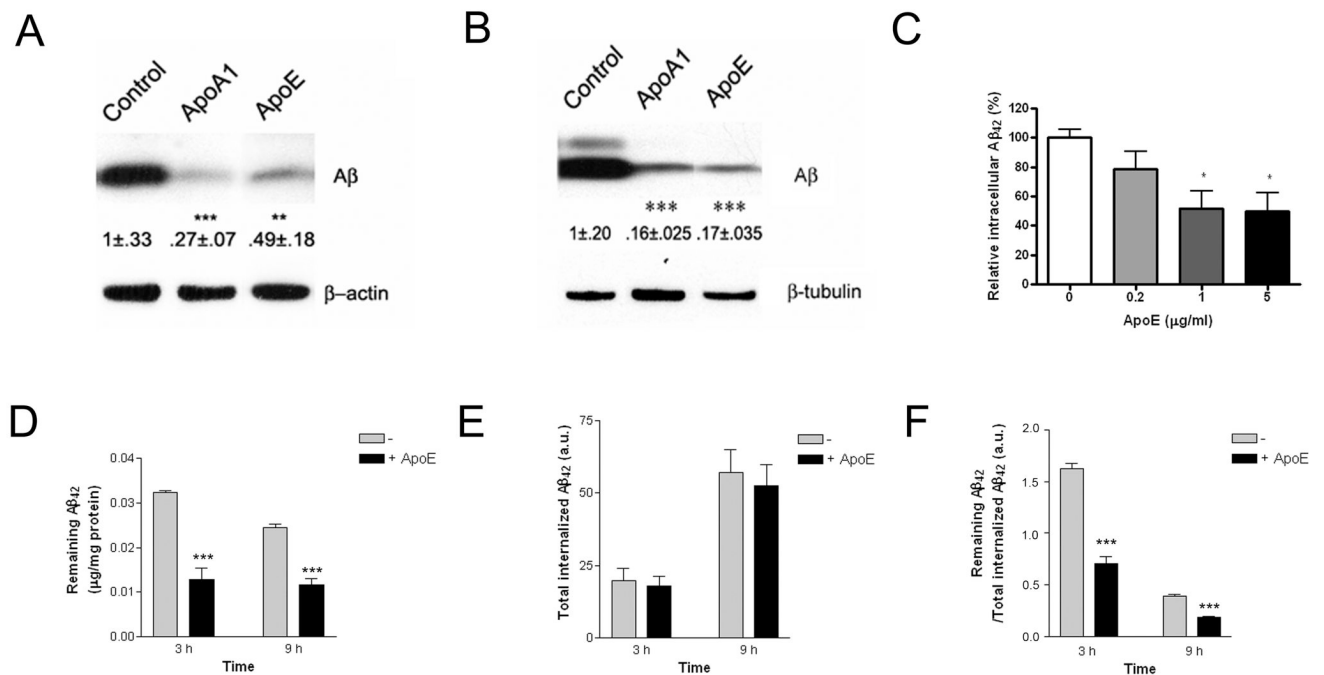
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### Figure 1. Microglia efficiently take up and degrade soluble A $\beta$

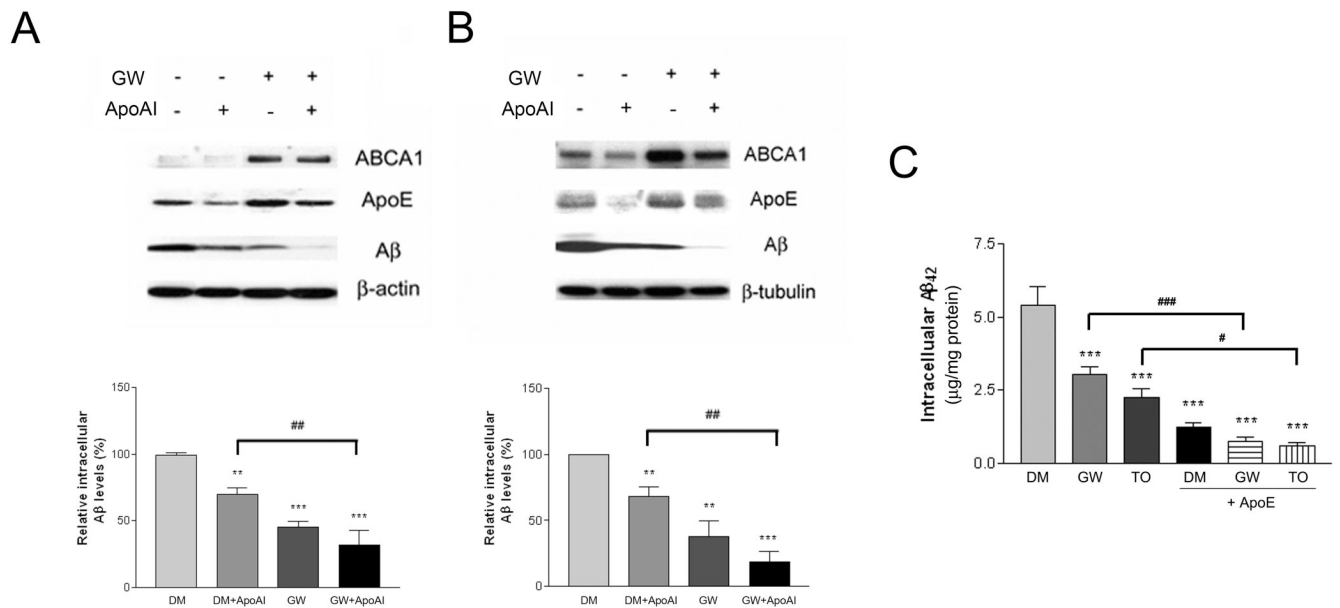
**(A)** Kinetics of soluble A $\beta$  uptake by microglia. BV2 microglia were incubated with exogenous Alexa488-labeled soluble A $\beta_{42}$  for the indicated periods. The uptake of fluorescently-labeled soluble A $\beta$  was measured using flow cytometry. **(B)** Soluble A $\beta$  is dose-dependently taken up by microglia. BV2 microglia were incubated for 3 hours with indicated concentrations of unmodified soluble A $\beta_{42}$ . The intracellular A $\beta_{42}$  levels were measured using ELISA. The results were normalized to total cellular protein. **(C)** Soluble A $\beta$  is rapidly trafficked to lysosomes for degradation. Confocal imaging of live BV-2 microglia 15 min after addition of 2  $\mu$ g/ml soluble Cy3-A $\beta_{42}$  demonstrated localization of A $\beta$  (red) within lysosomes. Lysosomes were stained using LysoTracker (green). **(D)** Exogenous soluble A $\beta$  was efficiently cleared from the medium by microglia. BV2 microglia were incubated with 2  $\mu$ g/ml A $\beta_{42}$  for the indicated time. The media were collected and immunoblotted for A $\beta$  (\*\* $P$ <0.001; ### $P$ <0.001). **(E)** Soluble A $\beta$  can be degraded by microglia. BV2 microglia were incubated for 3 hours with 2  $\mu$ g/ml unmodified soluble A $\beta_{42}$ . After washout of the remaining A $\beta_{42}$  in the media, the intracellular A $\beta_{42}$  levels were measured using ELISA. The media were also monitored for resecreted A $\beta$ . **(F)** Neprilysin and related proteinases mediate intracellular degradation of soluble A $\beta$  by microglia. Primary microglia from wild type mice were pretreated with vehicle, 10  $\mu$ M phosphoramidon or 1  $\mu$ M thiorphan for 18 hours. The cells were then incubated with 2  $\mu$ g/ml A $\beta_{42}$  in the presence of vehicle or drug for an additional 24 hours. The data represent the outcome of 3 independent experiments (\* $P$ <0.05).



**Figure 2. HDL Apolipoproteins increase soluble Aβ degradation in microglia**

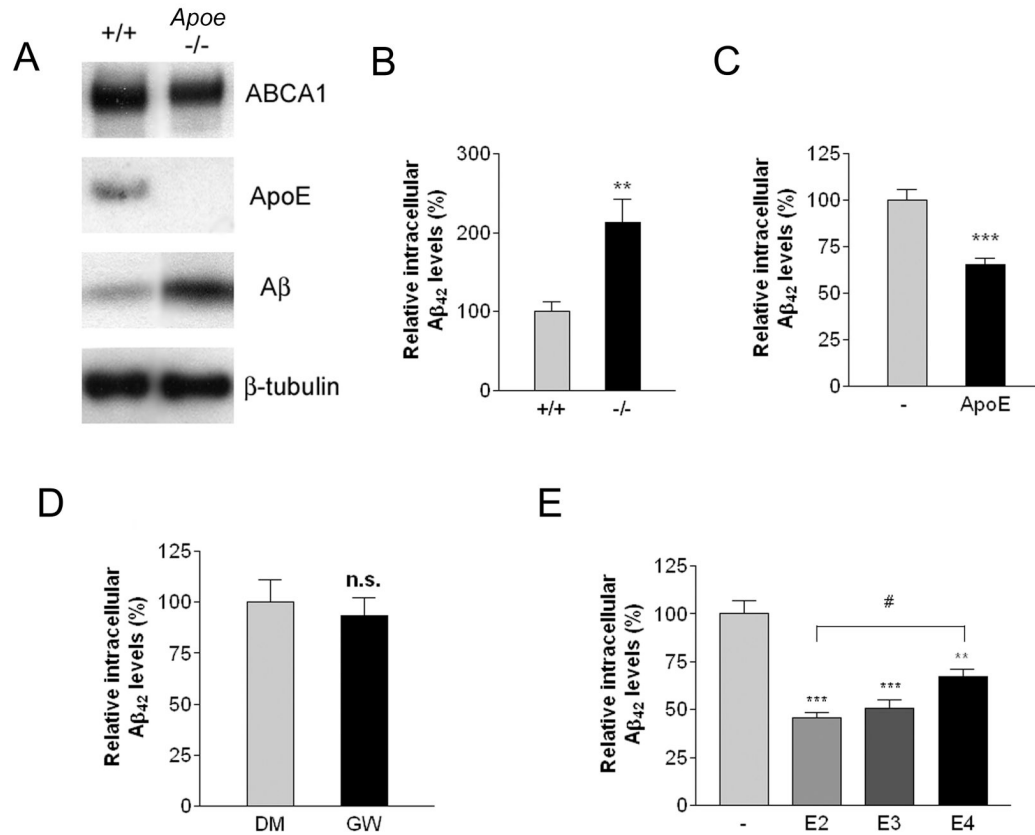
BV2 microglia (A) or primary microglia from wild type mice (B) were incubated with 2 μg/ml Aβ<sub>42</sub> in the presence or absence of 20 μg/ml ApoA-I or 5 μg/ml ApoE for 3 hours or 24 hours respectively. The intracellular Aβ<sub>42</sub> levels were quantified from three independent experiments and data normalized to loading controls. Values are expressed as percentage relative to the value of controls (\*\**P*<0.01; \*\*\**P*<0.001). (C) ApoE dose-dependently increased Aβ degradation in microglia. BV2 microglia were incubated with 2 μg/ml Aβ<sub>42</sub> in the presence of increasing concentrations of purified human ApoE (0, 0.2, 1 or 5 μg/ml) for 3 hours, intracellular Aβ<sub>42</sub> levels were quantified using ELISA and data normalized to total protein (\**P*<0.05). (D–F) ApoE enhanced soluble Aβ degradation. BV2 microglia were incubated with 2 μg/ml Aβ<sub>42</sub> or 2 μg/ml Alexa488-labeled Aβ<sub>42</sub> in the presence or absence of 1 μg/ml purified human ApoE for indicated time. (D) The remaining intracellular Aβ<sub>42</sub> levels were quantified using ELISA and data normalized to total protein (\*\*\**P*<0.001). (E) The total internalized Aβ<sub>42</sub> over the same intervals was monitored using flow cytometry. (F) The efficiency of Aβ degradation was expressed as a ratio of remaining intact Aβ<sub>42</sub> over total internalized Aβ<sub>42</sub> (\*\*\**P*<0.001).





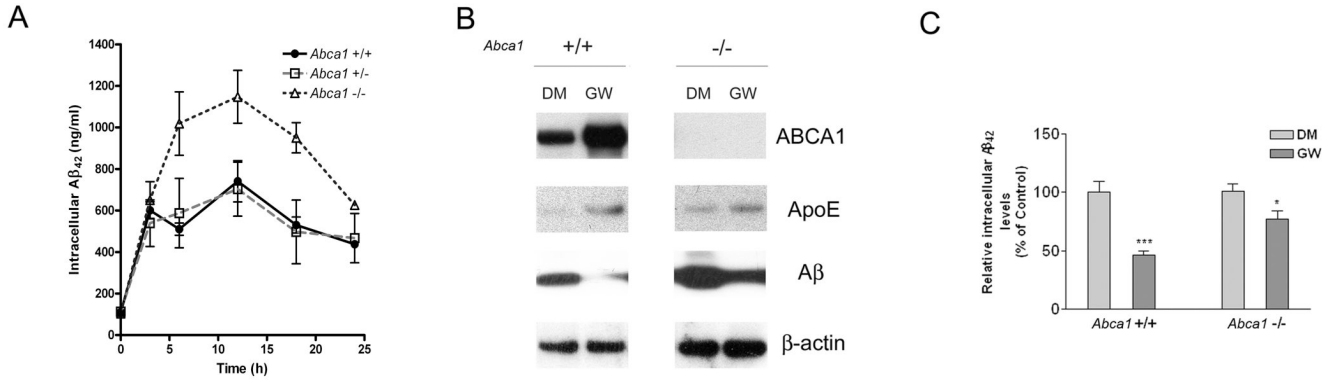
### Figure 3. LXR activation enhances intracellular A $\beta$ degradation in microglia

BV2 microglia (A) or primary microglia from wild type mice (B) were pretreated with DMSO or 1  $\mu$ M GW3965 for 18 hours. The cells were then incubated with 2  $\mu$ g/ml A $\beta_{42}$  in the presence or absence of 20  $\mu$ g/ml ApoA-I for 3 hours or 24 hours, respectively. Cellular lysates were subjected to SDS-PAGE and Western blotted for ABCA1, ApoE, A $\beta$  and  $\beta$ -actin or  $\beta$ -tubulin. The levels of A $\beta$  were normalized to  $\beta$ -actin or  $\beta$ -tubulin as loading controls. Relative intracellular A $\beta$  levels were quantified from three independent experiments (\*\* $P$ <0.01; \*\*\* $P$ <0.001; ## $P$ <0.01). (C) Primary microglia from wild type mice were pretreated with DMSO, 1  $\mu$ M GW3965 or 1  $\mu$ M T0901317 for 18 hours. The cells were then incubated with 2  $\mu$ g/ml A $\beta_{42}$  in the presence or absence of 5  $\mu$ g/ml purified human plasma ApoE for 24 hours. Intracellular A $\beta$  levels were quantified using ELISA for A $\beta_{42}$  and normalized to total protein. The data represent the outcome of 4 independent experiments (\*\*\* $P$ <0.001; # $P$ <0.05; ### $P$ <0.001).



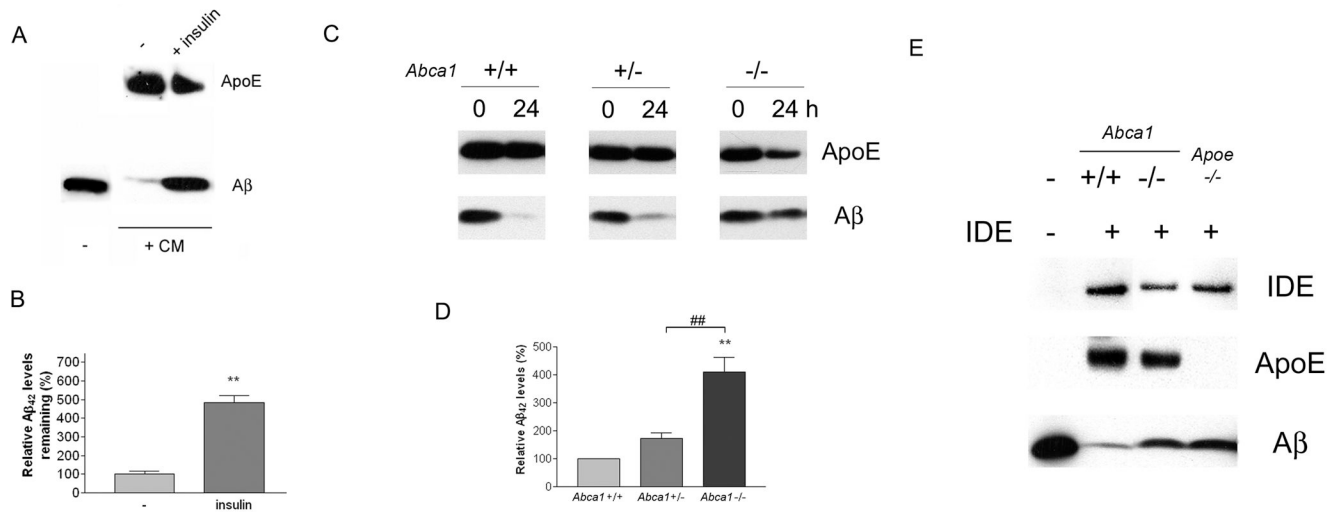
#### Figure 4. ApoE is essential for efficient degradation of Aβ

(A) Loss of ApoE resulted in intracellular accumulation of Aβ. Primary microglia from wild type or *ApoE*<sup>-/-</sup> mice were treated with 2 μg/ml Aβ<sub>42</sub> for 24 hours. The lysates were subjected to SDS-PAGE and Western blotted for ABCA1, ApoE, Aβ and β-tubulin as a loading control. Blots shown are representative of three independent experiments. (B) Intracellular Aβ levels were quantified using ELISA for Aβ<sub>42</sub> and normalized to total protein. The data represent the outcome of 5 independent experiments (\*\**P*<0.01). (C) Exogenous ApoE rescued the Aβ degrading deficiency in *ApoE*<sup>-/-</sup> microglia. Primary microglia from *ApoE*<sup>-/-</sup> mice were incubated with 2 μg/ml Aβ<sub>42</sub> in the absence or presence of 1 μg/ml ApoE for an additional 24 hours. Intracellular Aβ<sub>42</sub> levels were quantified using ELISA and the data normalized to total protein (\*\*\**P*<0.001). (D) ApoE is required for LXR-mediated effect on Aβ degradation. Primary microglia from *ApoE*<sup>-/-</sup> mice were pretreated with DMSO or 1 μM GW3965 for 18 hours. The cells were then incubated with 2 μg/ml Aβ<sub>42</sub> for an additional 24 hours. Intracellular Aβ<sub>42</sub> levels were quantified using ELISA and data normalized to total protein. (E) Native ApoE particles enhanced Aβ degradation in an isoform-dependent manner. Primary microglia from *ApoE*<sup>-/-</sup> mice were treated with 2 μg/ml Aβ<sub>42</sub> in the absence or the presence of 200 ng/ml purified ApoE-containing native HDL particles isolated from immortalized astrocytes expressing the human ApoE isoforms for 24 hours. The levels of remaining intracellular Aβ were quantified using ELISA for Aβ<sub>42</sub>. The data represent the outcome of 4 independent experiments (\*\**P*<0.01; \*\*\**P*<0.001; #*P*<0.05).



**Figure 5. ABCA1 influences the intracellular degradation of Aβ by microglia**

**(A)** Loss of *Abca1* impairs Aβ degradation in primary microglia. Primary microglia from *Abca1* +/+, +/- and -/- mice were treated with 2 μg/ml Aβ<sub>42</sub> for 24 hours. The cellular levels of Aβ<sub>42</sub> were measured in lysates by ELISA. The data were normalized to total protein and represent the outcome of 5 independent experiments. **(B)** Primary microglia from *Abca1* +/+ and -/- mice were pretreated with DMSO or 1 μM GW3965 for 18 hours. The cells were then incubated with 2 μg/ml Aβ<sub>42</sub> in the presence or absence of 1 μM GW3965 for 24 hours. Cellular lysates were subjected to SDS-PAGE and Western blotted for ABCA1, ApoE, Aβ and β-actin. Representative blots from 3 independent experiments are shown. **(C)** Primary microglia from *Abca1* +/+ and -/- mice were treated as above. Intracellular Aβ levels were quantified using ELISA for Aβ<sub>42</sub> and normalized to total protein. The data represent the outcome of 6 independent experiments (\**P*<0.05; \*\*\**P*<0.001).

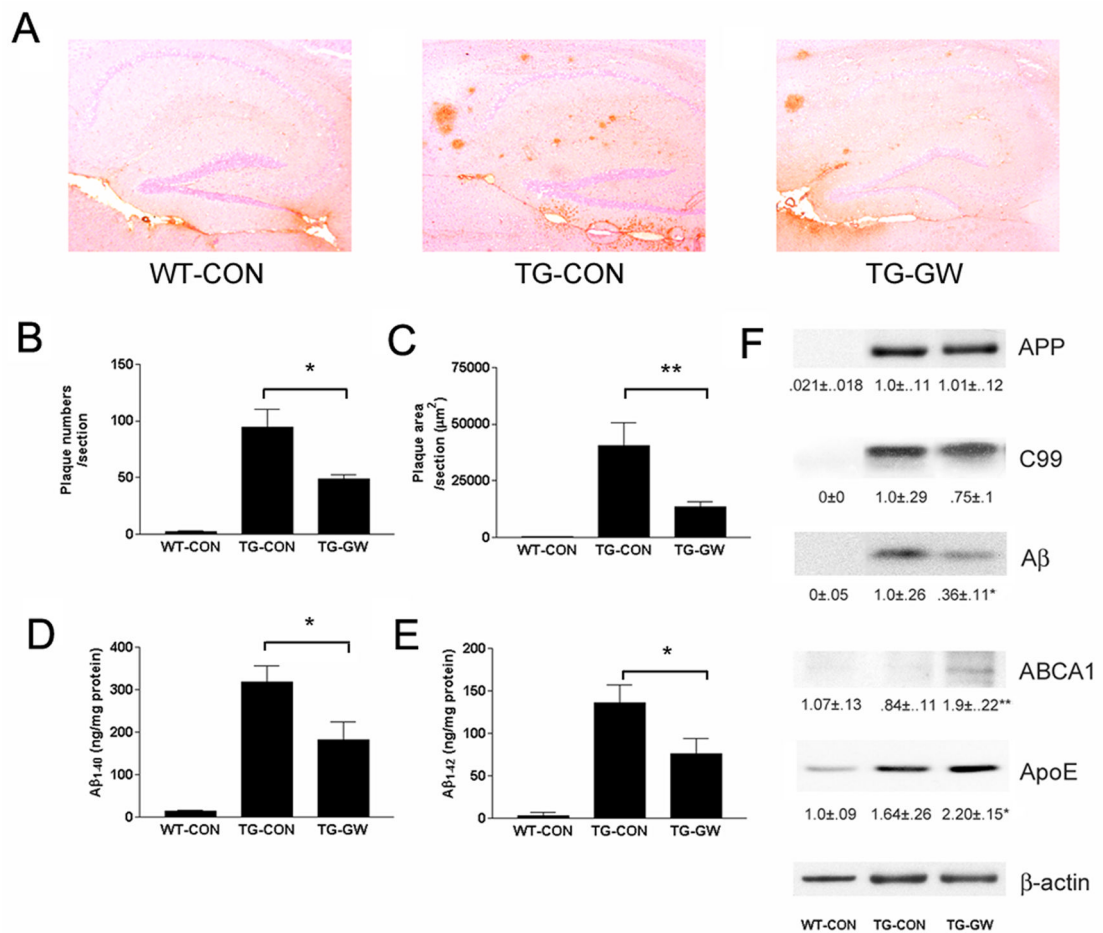


**Figure 6. Extracellular degradation of soluble A $\beta$  is dependent on IDE and related proteinases, and is influenced by the lipidation status of ApoE**

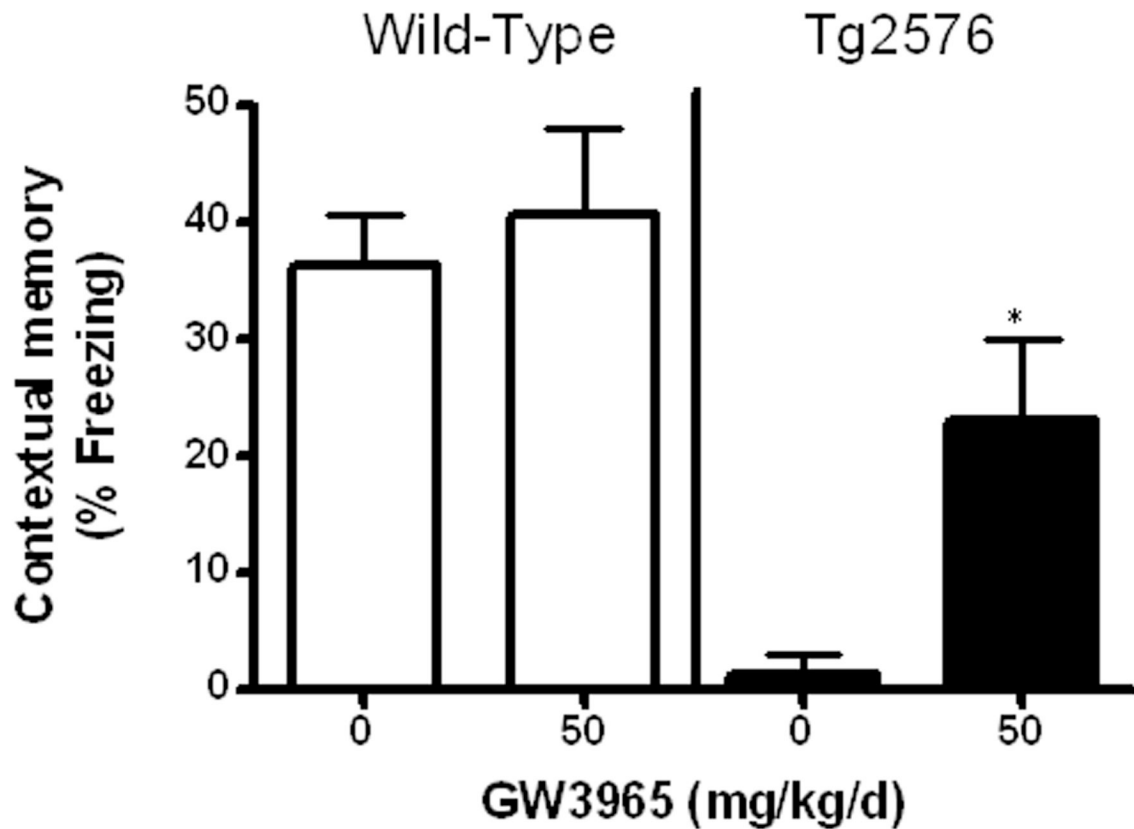
**(A) Astrocyte-conditioned medium degrades soluble A $\beta$  by IDE and related proteinases.**

Astrocyte-conditioned medium from wild type mice was incubated with A $\beta$ <sub>42</sub> in the presence or absence of 10  $\mu$ M insulin for 24 hours. The samples were then subjected to SDS-PAGE and Western blotted for ApoE and A $\beta$ . **(B)** After 24 hours, the samples were analyzed for A $\beta$ <sub>42</sub> using ELISA. The data represent the outcome of 3 independent experiments (\*\* $P < 0.01$ ). **(C) Lipidation status of ApoE regulates A $\beta$  degradation in astrocyte-conditioned medium.**

Conditioned media from astrocytes derived from *Abca1* +/+, +/-, and -/- mice were incubated with 1  $\mu$ g/ml A $\beta$ <sub>42</sub> for 0 or 24 hours and the reaction mixtures were subjected to SDS-PAGE and Western blotted for ApoE and A $\beta$ . Representative blots from three independent experiments are shown. **(D)** The amount of A $\beta$ <sub>42</sub> remaining in the medium after 24 hours was quantified using ELISA. The data represent the outcome of 3 independent experiments (\*\* $P < 0.01$ ; ## $P < 0.01$ ). **(E) ApoE-containing HDL particles promote A $\beta$  degradation by recombinant IDE.** ApoE-containing HDL particles were collected from astrocyte-conditioned medium derived from wild type, *Abca1* -/- and *ApoE* -/- mice by immunoprecipitation and the complexes were incubated with 500 ng/ml recombinant IDE and 2  $\mu$ g/ml A $\beta$ <sub>42</sub> for 1 hour. The reaction mixtures were then resolved by SDS-PAGE and Western blotted for IDE, ApoE and A $\beta$ .

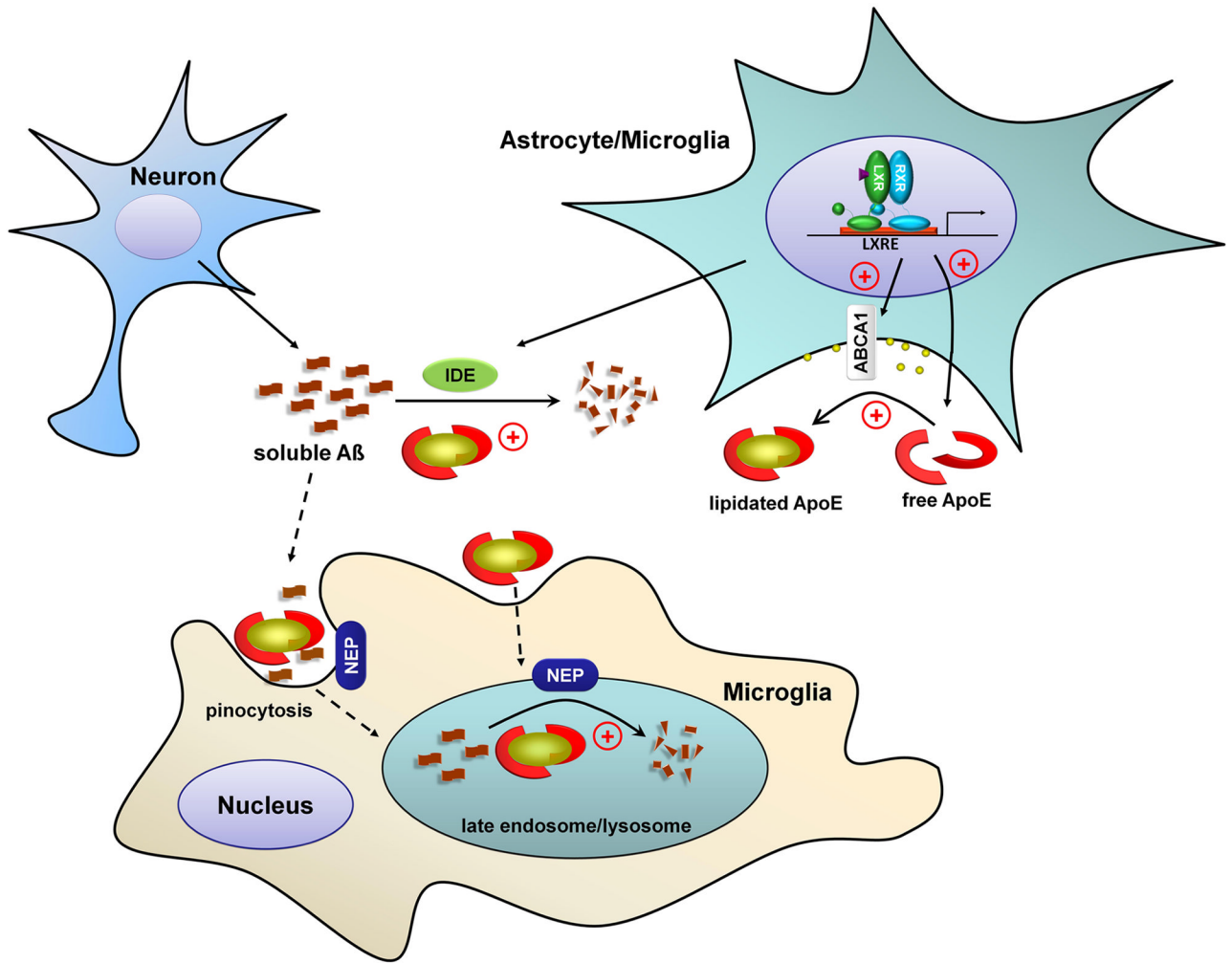


**Figure 7. LXR agonist treatment reduces Aβ levels and ameliorates plaque burden in Tg2576 mice** (A) Aged Tg2576 mice (12 month-old) or genetically similar controls were treated for 4 months with normal chow or chow containing GW3965 (120 mg/kg, 33 mg/kg/day). Aβ plaque burden was monitored by 6E10 staining in the hippocampus. Plaque number and plaque area were quantified in (B) and (C) respectively (n=5, \*P<0.05; \*\*P<0.01). The levels of Aβ<sub>40</sub> (D) and Aβ<sub>42</sub> (E) were quantified using ELISA (\*P<0.05). (F) The levels of full-length APP, C99 C-terminal fragment, total Aβ, ABCA1 and ApoE were monitored by Western analysis. The results were normalized to β-actin (n=5, \*P<0.05; \*\*P<0.01). Quantification of the data is shown under the Western blots.



**Figure 8. GW3965 significantly improves contextual memory in Tg2576 mice**

Tg2576 or wild type mice (20 week-old) were orally treated with vehicle or 50 mg/kg/day of GW3965 for 6 days and subjected to contextual memory assessment as outlined in the methods. Treatment of the LXR agonist improved hippocampal-dependent contextual memory in the heterozygous Tg2576 mice ( $n = 11$ ,  $*P < 0.05$ ). No significant effect of treatment was observed in contextual memory in the wild-type littermate control mice. No significant effect of genotype was observed in hippocampal-independent cue conditioning (data not shown).



**Figure 9. ApoE and its lipidation status regulates the proteolytic degradation of soluble A $\beta$**   
 The level of soluble A $\beta$  is homeostatically controlled by its production by neurons and its subsequent clearance. Soluble A $\beta$  can be cleared by proteolytic enzymes including NEP and IDE acting both intracellularly and extracellularly. ApoE is principally synthesized and secreted by glia. The lipid transporter ABCA1 mediates the lipidation of ApoE. Liver X receptors regulate the expression of both ABCA1 and ApoE and their activation results in increased levels of lipidated ApoE. The degradation of soluble A $\beta$  both intracellularly and extracellularly is enhanced by increasing ApoE and its lipidation.