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# Aβ42 Is Essential for Parenchymal and Vascular Amyloid

# **Deposition in Mice**

Eileen McGowan<sup>1,\*</sup>, Fiona Pickford<sup>1,4</sup>, Jungsu Kim<sup>1,4</sup>, Luisa Onstead<sup>1</sup>, Jason Eriksen<sup>1</sup>, Cindy Yu<sup>1</sup>, Lisa Skipper<sup>1</sup>, M. Paul Murphy<sup>1</sup>, Jenny Beard<sup>1</sup>, Pritam Das<sup>1</sup>, Karen Jansen<sup>1</sup>, Michael DeLucia<sup>1</sup>, Wen-Lang Lin<sup>1</sup>, Georgia Dolios<sup>2</sup>, Rong Wang<sup>2</sup>, Christopher B. Eckman<sup>1</sup>, Dennis W. Dickson<sup>1</sup>, Mike Hutton<sup>1</sup>, John Hardy<sup>3</sup>, and Todd Golde<sup>1,\*</sup>

1 Department Neuroscience Mayo Clinic College of Medicine Jacksonville, Florida 32224

2Department of Human Genetics Mount Sinai School of Medicine New York, New York 10029

3Laboratory of Neurogenetics National Institute on Aging National Institutes of Health Bethesda, Maryland 20892

# Summary

Considerable circumstantial evidence suggests that  $A\beta42$  is the initiating molecule in Alzheimer's disease (AD) pathogenesis. However, the absolute requirement for  $A\beta42$  for amyloid deposition has never been demonstrated in vivo. We have addressed this by developing transgenic models that express  $A\beta1-40$  or  $A\beta1-42$  in the absence of human amyloid  $\beta$  protein precursor (APP) overexpression. Mice expressing high levels of  $A\beta1-40$  do not develop overt amyloid pathology. In contrast, mice expressing lower levels of  $A\beta1-42$  accumulate insoluble  $A\beta1-42$  and develop compact amyloid plaques, congophilic amyloid angiopathy (CAA), and diffuse  $A\beta$  deposits. When mice expressing  $A\beta1-42$  are crossed with mutant APP (Tg2576) mice, there is also a massive increase in amyloid deposition. These data establish that  $A\beta1-42$  is essential for amyloid deposition in the parenchyma and also in vessels.

# Introduction

Overexpression of Alzheimer's disease (AD)-linked mutant human APP transgenes has been the most reliable means of promoting deposition of A $\beta$  in the brains of transgenic mice. As they age, these mutant APP mice develop robust amyloid pathology and other AD-like features, including decreased synaptic density, reactive gliosis, and some cognitive deficits. However, these mutant APP mouse models show little evidence of overt neuronal loss and neurofibrillary tangle (NFT) pathology (Hardy and Selkoe, 2002; Price et al., 1998). One potential problem with most of the widely studied mutant APP mice is that the high level of overexpression of mutant human APP may confound the phenotype. Overexpression of APP results in overproduction of APP fragments, which may have neuroprotective or neurotoxic functions. For example, secreted APP generated by  $\alpha$ -secretase (sAPP $\alpha$ ) can be neuroprotective, whereas the carboxyl terminal fragment generated by  $\beta$ -secretase cleavage (CTF $\beta$ ) and a caspasecleaved fragment of APP (C31) can be neurotoxic (Lu et al., 2000; Mattson, 2004; Yankner et al., 1989). Moreover APP and fragments such as the APP intracellular domain have signaling functions that may also contribute to a phenotype (LaFerla, 2002). Consequently, mice that overexpress individual AB peptides in the absence of overexpression of APP allow testing of hypotheses regarding (1) the role of select A $\beta$  species in the initiation and propagation of

<sup>4</sup>These authors contributed equally to this work.

<sup>\*</sup>Correspondence: mcgowan.eileen@mayo.edu (E.M.); golde.todd@mayo.edu (T.G.)

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amyloid deposition in vivo and (2) the specific contribution of each A $\beta$  peptide to the phenotype seen in AD mouse models.

Much of the data that support a pivotal role for  $A\beta42$  in AD have come from the study of mutations in the APP and presenilin genes that cause early-onset familial forms of AD (Selkoe, 1998). The vast majority of these mutations selectively increase the relative levels of  $A\beta42$ . However, even in typical late-onset AD there is evidence that  $A\beta42$ , a minor  $A\beta$  species, usually representing less then 20% of the total  $A\beta$  secreted, is both the earliest form and the predominant species deposited in the brain parenchyma (Golde et al., 2000). In contrast,  $A\beta40$ , the major  $A\beta$  peptide secreted by cells, appears to be the predominant species deposited in the cerebral vasculature (congophillic angiopathy, CAA) (Gravina et al., 1995; Iwatsubo et al., 1994). Transgenic mouse studies using mutant APP and PS transgenes have provided some insights into the effects that altering the ratio of  $A\beta40$  and  $A\beta42$  have on time to onset of deposition, type of deposit (e.g., diffuse versus compact), and extent of CAA (Borchelt et al., 1997; Herzig et al., 2004; Holcomb et al., 1998). However, such studies have not definitively identified which  $A\beta$  species are responsible for seeding amyloid deposition in either the parenchyma or vasculature.

To address this question, we have generated transgenic mice that express  $A\beta 1$ -40 or  $A\beta 1$ -42 without APP overexpression. For these studies we used cDNAs that express fusion proteins between the BRI protein, involved in amyloid deposition in Familial British (FBD) and Danish Dementia (FDD) and  $A\beta 1$ -40 (BRI- $A\beta 40$ ) or  $A\beta 1$ -42 (BRI- $A\beta 42$ ) (Lewis et al., 2001; Vidal et al., 1999, 2000)(Figure 1A). We have previously shown that transfection of BRI- $A\beta$  cDNAs results in high-level expression and secretion of the encoded  $A\beta$  peptide through proteolytic cleavage of the fusion protein at a furin cleavage site immediately preceding  $A\beta$  (Lewis et al., 2001). Efficient secretion of  $A\beta$  from the BRI fusion protein distinguishes this approach from studies using  $A\beta$  minigene constructs that generate high levels of intra-cellular  $A\beta$  and minimal secreted  $A\beta$  (LaFerla et al., 1995). The BRI- $A\beta$  transgenic mice we have generated provide substantial evidence that  $A\beta 1$ -42 but not  $A\beta 1$ -40 is sufficient to promote  $A\beta$  deposition in mice.

#### Results

#### Generation of BRI-Aß Transgenic Mice

The mouse prion promoter (MoPrp) (Borchelt et al., 1996) was used to drive BRI-A $\beta$ 40 and BRI-A $\beta$ 42 trans-gene expression. Linearized constructs were injected into fertilized oocytes from B6C3F1 mice. Multiple BRI-A $\beta$ 42 and BRI-A $\beta$ 40 founders were generated and maintained on a B6C3 background. Transgenic founders were screened by plasma-sandwich ELISAs. Six of the highest A $\beta$ 42-expressing founders, with plasma A $\beta$ 42 levels equivalent to or higher than A $\beta$ 42 levels in aged-matched Tg2576 mice (Hsiao et al., 1996), were selected for further breeding. Subsequently, the two lines that had the highest level of expression were chosen for further study. One BRI-A $\beta$ 1-40 founder had A $\beta$ 40 plasma levels higher than levels seen in age-matched Tg2576 mice (Kawarabayashi et al., 2001), and this line was also expanded.

Transgene expression and protein levels in the mice were determined by Northern blotting, in situ hybridization, and Western analysis (see Figure S1 in the Supplemental Data available online). Included in this analysis were BRI wild-type mice expressing the wild-type BRI protein under control of the MoPrP promoter. The BRI-A $\beta$ 40 line expressed the transgene at ~2- to 3-fold higher levels than the BRI-A $\beta$ 42A mice. In all lines, BRI-A $\beta$  mRNA was expressed in a pattern characteristic of the MoPrP promoter (Borchelt et al., 1996); with highest expression in the cerebellar granule cells and hippo-campus, followed by the cortex, pons, thalamus, and midbrain (Figures S1A and S1B). In the two highest expressing BRI-A $\beta$ 42 lines, BRI-A $\beta$ 42A

and BRI-A $\beta$ 42B, the fusion transgene was expressed at levels approximating endogenous mouse APP expression (BRI-A $\beta$ 42A) or at approximately half this level (BRI-A $\beta$ 42B). Fulllength BRI-A $\beta$  proteins levels (~37 kDa) were highest in BRI-A $\beta$ 40, followed by BRI-A $\beta$ 42A, with BRI-A $\beta$ 42B having the lowest protein levels (Figures S1D and S1E). Processed A $\beta$ species (4 kDa) were detected in both the BRI-A $\beta$ 40 and BRI-A $\beta$ 42 transgenic mice by immunoprecipitation Western blotting (Figure S1F).

Full-length BRI-A $\beta$ 40 or BRI-A $\beta$ 42 did not show significant alterations with age (Figure S2). BRI-A $\beta$ 40 and BRI-A $\beta$ 42 mice have a normal lifespan with no obvious behavioral abnormalities.

#### Specific Expression of A<sub>β</sub> Peptide Isoforms in Transgenic Mice

Although 4 kDa A $\beta$  species were detected by Western immunoprecipitation, MALDI-TOF mass spectrometry (IP/MS) (Wang et al., 1996) was used to definitively identify the specific A $\beta$  isoforms in PBS extracts from the brains of BRI-A $\beta$  mice in comparison to Tg2576 (Figure 1B). The BRI-A $\beta$ 42 lines show single peaks corresponding to human A $\beta$ 1-42. The relative peak heights between the two lines were consistent with other data indicating that the BRI-A $\beta$ 42A mice had higher expression than line BRI-A $\beta$ 42B. In BRI-40 mice, a single peak was detected that corresponded to human A $\beta$ 1-40. Several peaks were detected in extracts from Tg2576 mice that corresponded to human A $\beta$ 1-38, A $\beta$ 1-40, and A $\beta$ 1-42. No peaks were detected in the wild-type BRI transgenic or nontransgenic mice.

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A $\beta$  peptides in the brains of BRI-A $\beta$ 42 and BRI-A $\beta$ 40 mice from 3 to 22 months of age were characterized by A $\beta$  sandwich ELISAs (Figures 1C and 1D). Brains from Tg2576 mice were used as controls. RIPA-soluble A $\beta$ 1-40 levels in 3-month-old BRI-A $\beta$ 1-40 mice (80-120 pm/g) were ~2- to 3-fold higher than levels of A $\beta$ 1-40 in 3-month-old Tg2576 mice (~40 pm/gm). No appreciable change in the level of RIPA-soluble A $\beta$ 1-40 was observed up to 22 months of age, and there was no evidence for A $\beta$  accumulation in the RIPA-insoluble formic acid extractable (FA) fraction. Levels of RIPA-soluble A $\beta$ 1-42 in 3-month-old BRI-A $\beta$ 42A and BRI-A $\beta$ 42B brains were ~2- to 3- and ~1- to 1.5-fold higher than the A $\beta$ 1-42 levels in 3- to 6-month-old Tg2576 brains. Both BRI-A $\beta$ 42 lines showed large increases in RIPA-insoluble A $\beta$ 1-42 levels with age. In the oldest BRI-A $\beta$ 42 mice, the levels of RIPA-insoluble A $\beta$ 1-42 exceed the A $\beta$ 1-42 levels seen in aged Tg2576 mice. A $\beta$ 42 and A $\beta$ 40 could be readily detected in plasma from the BRI-A $\beta$ 42 mice, and conversely, no A $\beta$ 42 was detected in plasma from BRI-A $\beta$ 42 mice.

#### Extensive Vascular and Parenchymal Amyloid Pathology in BRI-Aβ42 but Not BRI-Aβ40 Mice

The biochemical analysis indicated that BRI-A $\beta$ 42 mice selectively accumulated detergentinsoluble A $\beta$  as they aged and that the BRI-A $\beta$ 40 mice did not. Immuno-staining with anti-A $\beta$  antibodies and thioflavin S staining showed that both lines of BRI-A $\beta$ 42 mice, but not BRI-A $\beta$ 40 mice, developed compact amyloid deposits and diffuse A $\beta$  deposits with age (Figure 2). In contrast to other mutant APP transgenic mice (Hsiao et al., 1996;Price et al., 1998), cerebellar A $\beta$  accumulation was the earliest and most consistent feature in both lines of BRI-A $\beta$ 42 mice. In the BRI-A $\beta$ 42A mice, cored plaques were observed as early as 3 months of age in the molecular layer, with diffuse A $\beta$  deposition becoming more prominent with age. Forebrain pathology occurred later and was more variable; occasional extracellular plaques were observed in the entorhinal/piriform cortices and the hippocampus at 6 months of age, but were not consistently present until at least 12 months in BRI-A $\beta$ 42A mice. The oldest mice had more widespread pathology, developing thioflavin S-positive-cored and diffuse plaques in the cerebellum, cortex, hippocampus, and olfactory bulb, in a similar manner to pathology

described for Tg2576 (Figure S4). Onset of amyloid pathology in the BRI-A $\beta$ 42B mice was more variable than in the BRI-A $\beta$ 42A mice; by 20 months of age, these mice had widespread cerebellar and forebrain pathology with a similar plaque distribution to that seen in the BRI-A $\beta$ 42A mice. The greatest amlyoid plaque burden was detected with N-terminal antibodies such as Bam-10 and 82E1 (Figures 2 and 4). Immunostaining with A $\beta$ 40 or A $\beta$ 42 end-specific antibodies was consistent with the notion that the deposited A $\beta$  was almost entirely composed of A $\beta$ 42 (Figure 2E). We could not rule out that trace amounts of A $\beta$ 40, derived from endogenous mouse APP, or limited proteolysis, were present in the core of some large compact plaques. Despite similar levels of BRI expression at the mRNA level, there was no pathology apparent in the brains of BRI wild-type mice at any age (up to 24 months; Figure 2L).

Congophilic amyloid angiopathy (CAA) was absent in the BRI-A $\beta$ 40 mice, but was a prominent feature in the BRI-A $\beta$ 42 mice (Figure 2K). EM analysis demonstrated that the CAA was fibrillar and closely associated with the basal lamina (Figures 3A and 3B), similar to CAA observed in AD. EM analysis of extracellular amyloid plaques showed dense amyloid cores with radiating fibrils (Figure 3C). Many bundles of dystrophic neurites were also present at the periphery of the dense cored extracellular plaques. Compact plaques were small relative to those seen in Tg2576 mice (mean plaque diameter  $38 \pm 5 \mu m$  at 3 months to  $53 \pm 3 \mu m$  at 16-18 months of age). Reactive astrogliosis was associated with the compact plaques (Figure 2J). There was little evidence of tau pathology, with only rare CP13-positive dystrophic neurites seen in association with the dense cored plaques in the oldest mice.

To determine if endogenous mouse  $A\beta$  was present in the deposits, a monoclonal antibody raised against aggregated mouse  $A\beta42$  (32.4.1) that specifically recognizes an epitope in mouse  $A\beta1-16$  and does not cross-react with human  $A\beta$  by ELISA (data not shown) was used to stain brain sections from human AD and BRI-A $\beta42$  mice. This antibody did not detect plaques from human AD tissue (Figures 4A and 4B). In BRI-A $\beta42$  mice, the antibody labeled a subset of the A $\beta$  plaque cores, particularly in the cerebellum, and some diffuse plaques in older mice (Figures 4C and 4D). These data indicate that endogenous mouse A $\beta$  is present in the plaques. We also immunostained sections from the BRI-A $\beta42$  mice with an antibody specific for the N-terminal truncated and modified pyroglutamate A $\beta$  species A $\beta$ [N3pe]. A proportion of the cored plaques and, to a lesser extent, the diffuse amyloid were immuno-positive for A $\beta$ [N3pe], showing that this N-terminal modified pyroglutamate A $\beta$  species was present in the plaques in the BRI-A $\beta$  mice (Figures 4E and 4F).

#### Enhanced Pathology in BRI-Aβ42 × Tg2576 Bigenic Mice

Hemizygous BRI-A $\beta$ 42A mice were crossed with hemizygous APP<sub>swe</sub>Tg2576 mice, and the amount of Aβ deposition in the bigenic BRI-Aβ42/Tg2576 offspring, age-matched singly transgenic BRI-Aβ42A, and Tg2576 and nontransgenic littermates was analyzed by biochemical and immunohistochemical methods. Amyloid deposition initially develops in the cerebellum of BRI-A $\beta$ 42 mice and is extensive by 12 months of age, whereas, forebrain pathology only consistently develops after 12 months of age. In contrast, amyloid deposition in Tg2576 mice only occurs in the forebrain after 6 months of age, and no cerebellar deposits are routinely observed. For this reason, we analyzed the hindbrain (including cerebellum) and forebrain A $\beta$  levels independently at 6, 10, and 14.5 months (Figure 5). Anti-A $\beta$ immunohistochemsitry showed marked augmentation of A $\beta$  deposition throughout the cortex, hippocampus, and cerebellum. Representative sections of the entorhinal/piriform cortical areas from 14.5-month-old BRI-Aβ42A, Tg2576, and bigenic BRI-Aβ42/Tg2576 litter-mate mice are shown in Figures 5A-5C. RIPA-soluble and RIPA-insoluble FA A $\beta$  levels in both the forebrain and hindbrain were analyzed by AB ELISA. At 6 months of age, the levels of RIPAsoluble A $\beta$  in the forebrain (Figure S5A), hindbrain (data not shown), and plasma (Figure S5B) of the crossed mice were consistent with an additive effect of the two transgenes with respect to A $\beta$  production, providing assurance that alteration in transgene expression would not confound these studies. A marked increase in both RIPA-insoluble FA extracted A $\beta$ 40 and A $\beta$ 42 was seen in the forebrain of bigenic mice compared to single transgenic BRI-A $\beta$ 42 or Tg2576 littermates (Figure 5D). The increase in FA A $\beta$  levels was greater than that expected for a simple additive effect of A $\beta$  levels in BRI-A $\beta$ 42 and Tg2576 (~30-to 40-fold increase at 10 months of age in the fore-brain). Additionally, although A $\beta$  deposition does not occur in the cerebellum of Tg2576, A $\beta$ 40 and A $\beta$ 42 levels were substantially elevated in the hindbrain of the bigenic mice compared to the single transgenic littermates (Figure 5E). Taken together these data suggest that A $\beta$ 42, by itself, accelerates amyloid deposition in the Tg2576 mice.

# Discussion

We have developed transgenic mice expressing BRI-A $\beta$ 40 or BRI-A $\beta$ 42 fusion proteins. Biochemical analyses of these mice demonstrate that these transgenes selectively express the encoded A $\beta$  peptide in vivo. Since there is no pathological phenotype in mice over-expressing the BRI wild-type protein under control of the MoPrP, any resultant phenotype is likely to be attributed to the specific AB peptide expressed. BRI-AB40 mice express 2- to 3-fold higher levels of A $\beta$ 1-40 than Tg2576 mice, but do not develop A $\beta$  deposits by 24 months of age. In contrast, the BRI-A $\beta$ 42 lines, which have ~5- to 10-fold lower transgene expression than BRI-A $\beta$ 40 mice, develop robust parenchymal and cerberovascular A $\beta$  deposits as early as 3 months of age. Reactive gliosis was observed in the BRI-Aβ42 mice, but the mice had neither obvious neuronal cell loss nor prominent neurofibrillary pathology. The lack of a more robust AD-like phenotype in the BRI-A $\beta$ 42 mice suggests that the absence of a more prominent AD phenotype in mutant APP mice cannot be attributed to protection conferred by overexpression of other APP derivatives (e.g., sAPP $\alpha$ ) (Mattson, 2004), although this can only be definitely tested by crossing the BRI-Aβ42 mice into an APP knockout background. Bigenic BRI-Aβ42/Tg2576 mice show a marked increase in the extent of A $\beta$  deposition and validate the physiologic relevance of selectively increasing A $\beta$ 42 via the BRI fusion proteins. Such data are entirely consistent with the increases in deposition produced by APP and PS mutant transgenes that shift  $\gamma$ -secretase cleavage to augment A $\beta$ 42 production and accelerate A $\beta$  deposition (Borchelt et al., 1997; Holcomb et al., 1998).

The pattern of amyloid deposition in BRI-A $\beta$ 42 mice generally followed the expression pattern of the PrP transgene (Borchelt et al., 1996) and was similar to Tg2576 pathology except for the presence of a large number of plaques in the cerebellum of BRI-A $\beta$ 42 mice. The extensive A $\beta$ 42 deposition in the cerebellum and severe CAA in the BRI-A $\beta$ 42 mice mimics the pattern of amyloid deposition seen in FBD (Ghiso et al., 2000; Holton et al., 2001; Vidal et al., 1999). Though high levels of transgene expression are seen in the cerebellum of both Tg2576 and BRI-A $\beta$ 42 mice, we do find that the level of RIPA-soluble A $\beta$  in the nondepositing Tg2576 forebrain is ~50% higher than that found in the hind-brain and that BRI-A $\beta$ 42 mice generate significantly higher A $\beta$  levels in their hindbrain compared to Tg2576 mice. This may indicate that the level of enzymatic processing of BRI-A $\beta$ 42 and APP varies in different brain regions and might in turn partially account for the difference in the distribution of amlyoid deposition in Tg2576 and BRI-A $\beta$ 42 mice and between AD and FBD. However, the data from the BRI mice are potentially confounded by the fact that cerebellar A $\beta$  levels prior to deposition.

As has been reported from biochemical data in one APP transgenic line (Pype et al., 2003), we find immunohistochemical evidence for codeposition of mouse  $A\beta$  within a subset of plaques. Such data are not surprising given that in vitro data demonstrate that human and mouse  $A\beta$  can coassemble into aggregates (Fung et al., 2004). We also find evidence for truncation and modification of the deposited  $A\beta$ ; a subset of plaques are labeled by an antibody recognizing  $A\beta$ [N3pe]. The most prominent staining with this antibody was seen in the compact cerebellar

plaques of old (18 month) BRI-A $\beta$ 42 mice. Since the cerebellum is the initial site of deposition in these mice, such data would be consistent with postdeposition modification of the deposited A $\beta$ .

A comparison of the various transgenic models of  $A\beta$  deposition reveals that there is a great deal of variability with respect to the age at which deposition begins, the nature and composition of the A $\beta$  deposits (compact versus diffuse; proportion of A $\beta$ 40 versus A $\beta$ 42), and the extent of CAA (Borchelt et al., 1997; Chishti et al., 2001; Games et al., 1995; Herzig et al., 2004; Holcomb et al., 1998; Hsiao et al., 1996; Mucke et al., 2000; Sturchler-Pierrat et al., 1997). One factor that clearly regulates A $\beta$  deposition is the relative levels of A $\beta$ 42, which can be manipulated by either expressing APP transgenes with FAD-linked mutations (e.g., APP V717I/F mutations) or with coexpression of mutant presentiin transgenes that shift  $\gamma$ -secretase cleavage to favor increased production of Aβ42. Increasing relative Aβ42 levels lowers the age when deposition first occurs, increases the amount of diffuse AB, and promotes parenchymal versus cerebrovascular deposition. In contrast, a high ratio of A $\beta$ 40 to A $\beta$ 42 was associated with severe CAA in the absence of parenchymal amyloid (Herzig et al., 2004). However, because these mutations have other physiological effects, it was possible that other alterations in function contributed to the phenotype. Our data critically extend these observations and provide unequivocal in vivo evidence that, in the absence of mutations within the A $\beta$  sequence, A $\beta$ 42 expression appears to be necessary for both parenchymal and cerebrovascular amyloid deposition. Thus, although the  $A\beta 40/A\beta 42$  ratio influences the extent of vascular versus parenchymal deposition, A $\beta$ 42 appears to be required to "seed" both types of deposits.

In vitro biophysical studies of synthetic  $A\beta$  demonstrate that  $A\beta42$  aggregates more readily than  $A\beta40$ , presumably because  $A\beta42$  nucleates more efficiently. Following nucleation, there seems to be little kinetic or thermodynamic differences between elongation of fibrils composed of either  $A\beta40$  or  $A\beta42$  (Hasegawa et al., 1999; Jarrett et al., 1993; Jarrett and Lansbury, 1993). Although numerous *trans*-acting factors such as ApoE (Bales et al., 1999; Holtzman et al., 2000) and  $A\beta$  degrading enzymes can modulate  $A\beta$  deposition (Farris et al., 2003; Ohno et al., 2004), alterations in these factors do not result in complete reversal of the deposition phenotype. Thus, the simplest explanation for the complete lack of pathology in the BRI- $A\beta40$  mice would be the relative inability of  $A\beta40$  to initiate nucleation events capable of promoting amyloid deposition. Similar in vivo results have been obtained in a *Drosophila* model using a different fusion protein strategy (Ijjima et al., 2004). Expression of  $A\beta1-42$ , but not  $A\beta1-40$ , promoted peptide accumulation of diffuse intracellular and extracellular deposits in the transgenic flies. Significantly, such  $A\beta42$  accumulation was associated with neurodegeneration.

These findings raise an important question regarding the possible role of intracellular  $A\beta$  in Alzheimer's disease. Previous studies with transgenic mice expressing  $A\beta$  minigenes that are not efficiently secreted suggest that intracellular  $A\beta$  can be neurotoxic (LaFerla et al., 1995), as do studies of direct injection of aggregated  $A\beta$  into cultured primary neurons (Zhang et al., 2002). However, none of these studies prove or disprove a role of intracellular  $A\beta$  in AD pathogenesis. The BRI- $A\beta$  mice that we have developed are clearly distinct from these other "minigene" models in that they efficiently secrete  $A\beta$ . To date we have little evidence for accumulation of intracellular  $A\beta$  in these mice; thus, they provide an excellent tool to study the effects of secreted  $A\beta$  independently of APP transgenes.

Collectively our data provide further support for the hypothesis that  $A\beta 42$  is the initiating molecule in the pathogenesis of AD (Hardy and Selkoe, 2002; Younkin, 1998). These data provide definitive in vivo confirmation of predictions inferred from (1) in vitro studies of  $A\beta$  aggregation, (2) effects of PS and certain APP mutations on  $A\beta$  production and deposition,

and (3) analyses of  $A\beta$  peptides in the human AD brain (Gravina et al., 1995; Iwatsubo et al., 1994). Although our studies show that  $A\beta42$  is necessary to initiate aggregation of  $A\beta$ , they do not demonstrate that aggregation of  $A\beta42$ , by itself, is sufficient to induce all the hallmark pathologies of AD in mice, including neurofibrillary tangles and cell loss. This may reflect a simple limitation of the mouse models that we and others employ, or could point to some more complex aggregation phenomena that may require interaction of  $A\beta42$  with  $A\beta40$  or other  $A\beta$  species to drive AD pathogenesis. In any case, the BRI-A $\beta$  transgenic mice that we have developed provide models for further studies of the potential physiologic and pathologic roles of  $A\beta40$  and  $A\beta42$  in vivo.

## **Experimental Procedures**

#### **Generation of Transgenic Mice**

The BRI-A $\beta$ 40, BRI-A $\beta$ 42 fusion constructs (Lewis et al., 2001), or BRI wild-type constructs were subcloned into the MoPrP/bluescript vector (Borchelt et al., 1996) at the XhoI restriction site. The transgene was linearized by NotI restriction digest, gel purified, and digested with  $\beta$ -agarase according to the manufacturer's conditions (New England Biolabs). The linearized construct was injected into the pronuclei of single-cell embryos from C3B6F1 × B6 mice and implanted into pseudopregnant females. Founder pups were identified by PCR for the fusion construct or wild-type construct against an internal PS2 control PCR. Founder mice were back-crossed to C3B6F1 mice, and the transgenic lines were maintained on a C3B6 hybrid strain. All animal procedures were conducted with Mayo Clinic's animal care and use committee approval.

#### Bigenic BRI-Aβ42 × Tg2576 Mice

Hemizygous BRI-A $\beta$ 42A mice were mated with Tg2576 mice expressing mutant APP<sub>swe</sub>under control of the hamster prion promoter (Hsiao et al., 1996), generating bigenic BRI-A $\beta$ 42/Tg2576, BRI-A $\beta$ 42, Tg2576, and nontransgenic offspring. Tg2576 mice were maintained on a B6/SJL background. For all studies, age- and strain-matched littermates were used.

#### In Situ Hybridization

Sagital cryostat sections (15  $\mu$ m) were fixed in 4% paraformaldehyde, dehydrated, and hybridized with a transgene-specific oligomer (5'-

ACACAGTCGATAAATAAAGAAACCCAGGTGATCAATG-3') 3'-end labeled with  $\alpha^{35}$ S dATP. Sections were hybridized at 37°C overnight in buffer containing 4× SSC, 1× Denhardt's solution, 50% w/v deionized formamide, 10% w/v dextran sulfate, 200 mg/µl herring sperm DNA, and 0.03% β-mercaptoethanol. Control sections were hybridized in the presence of a 50- to 100-fold molar excess of unlabeled oligonucleotide. After hybridization, the sections were stringently washed (1× SSC at 50°C), dehydrated, and exposed to  $\beta$  max hyperfilm (Amersham) for 7-10 days.

#### **Northern Blotting**

Total RNA was extracted from the brains of male and female BRI-A $\beta$  mice using TRIzol reagent (Invitrogen) according to the manufacturer's instructions. Nontransgenic and mutant BRI transgenic mice were used as controls. Total RNA (15 µg) was run on a MOPS-formaldehyde denaturing gel and transferred to nitrocellulose membranes by capillary action. The blot was probed with a <sup>32</sup>Pradiolabeled oligonucleotide probe specific for both mouse and human BRI, stripped, and reprobed with a <sup>32</sup>P-radiolabeled APP cDNA probe and subsequently with a <sup>32</sup>P-radiolabeled histone probe. Band densities were quantified using phosphorimage screens (Kodak) and Imagequant software (Molecular Dynamics).

#### Western Analysis

Proteins were extracted from brain tissues by sonication in a 2% SDS/1% protease inhibitor cocktail solution. The resulting homogenate was centrifuged at 100,000 × g for 1 hr at 4°C. The supernatant containing the SDS-soluble fraction was aliquoted and stored at -80°C. Total protein concentrations were determined using the "BCA Protein Assay kit" (Pierce). Protein extracts from BRI-A $\beta$  mice were run on a SDS denaturing gel and transferred to Immobilon P membranes. Tg2576 and wild-type BRI transgenic lines were included as controls. Blots were probed with monoclonal antibodies to human A $\beta$  (6E10 and BAM10) followed by anti-mouse HRP or anti-rabbit HRP conjugated secondary antibody, respectively. Blots were stripped and reprobed with anti- $\beta$ -actin or  $\alpha$ -tubulin as a loading control.

#### Immunoprecipitation Western Analysis

Brain samples were homogenized in 2% SDS and diluted in RIPA buffer. Protein G was added and the sample mixed at 4°C for 1 hr prior to centrifugation for 60 s at 13000 rpm. The rabbit polycolonal A $\beta$  antibody, 3160, with Protein G (Pierce) was used to immuno-precipitate the A $\beta$  peptide at 4°C overnight. Then samples were centrifuged and the supernatants run on SDS denaturing gels (Biorad) and transferred to 0.2 µm nitrocellose membranes. Blots were boiled for 10 min in 0.1 M PBS and probed with the monoclonal anti-A $\beta$ , 6E10, followed by antimouse HRP. Tg2576 and nontransgenic samples and A $\beta$  peptides were run as controls.

#### Mass Spectrometry

Six-month-old brains from the BRI-A $\beta$ 42, BRI-A $\beta$ 40, Tg2567, wild-type BRI lines, and nontransgenic littermates were homogenized in PBS plus protease inhibitor solution to extract soluble proteins. Human A $\beta$  peptides were isolated by immunoprecipitation using two monoclonal anti-A $\beta$  antibodies (6E10 and 4G8) and analyzed by a MALDI-TOF mass spectrometry (Wang et al., 1996).

#### Aβ Sandwich ELISAs

Mouse hemibrains (excluding the olfactory bulb) for BRI-A $\beta$ 40 and BRI-A $\beta$ 42 studies or forebrain and hindbrain for BRI-A $\beta$ 42 × Tg2576 studies were homogenized in RIPA buffer and ultracentrifuged to separate RIPA-soluble and -insoluble fractions. The insoluble proteins were extracted using 70% formic acid. Formic acid samples were neutralized with Tris base buffer, and all samples were appropriately diluted (from 10- to 5000-fold). For plasma analysis of A $\beta$ , blood samples were centrifuged at 3000 rpm for 10 min at 4°C and the plasma decanted and diluted appropriately. A $\beta$  levels were determined by end-specific sandwich ELISAs (Kawarabayashi et al., 2001) using Ab9 (anti A $\beta$ 1-16 of A $\beta$ ) as the capture antibody and 21.3.1 (anti A $\beta$  35-42) or 13.1.1 (anti A $\beta$  35-40) as the detection antibodies for A $\beta$ 42 and A $\beta$ 40, respectively (n = 3-7 mice per age point per group). Nontransgenic mice (n = 13) and Tg2576 mice at 3, 6, 12 and 18 months (n = 3 at each time point) were included on each ELISA plate as controls.

#### Immunohistochemistry

Paraffin-embedded tissue sections (5  $\mu$ m) from BRI-A $\beta$ 40, BRI-A $\beta$ 42, and wild-type BRI mice (3-22 months of age, minimum of n = 3 per age group) were immunostained with A $\beta$  antibodies, including anti-total A $\beta$  (Bam 10, 1:50,000, Sigma; 6E10, 1:10,000, Signet; 33.1.1, 1:1,000; T. Golde; 82E1, 1:1000, IBL), anti-A $\beta$ 40 (13.1.1, 1:1,000, T. Golde), anti-A $\beta$ 42 (21.3.1, 1:1,000, T. Golde), 4G8 (A $\beta$ 17-24; 1:5,000, Signet), anti-A $\beta$ [n3pE] (8E1, 3rd N-terminal residue converted to pyroglutamate, 1:1000, IBL), and anti-rodent A $\beta$  (32.4.1. 1:1000, C. Eckman/T. Golde) on a DAKO autostainer using standard avidin or streptavidin biotin peroxidase methods. Additional sections were immunostained with thioflavin S, anti-tau antibodies (CP13, 1:100, P. Davies), and glial markers (GFAP, 1:100, Biogenex).

#### **Electron Microscopy**

BRI-A $\beta$  mice were perfused with saline followed by 2.5% glutaraldehyde/2% paraformaldehyde/0.1 M cacodylate buffer. The entorhinal cortex was removed, cut into 2 mm<sup>3</sup>pieces, postfixed in 1% osmium tetroxide and 1% uranyl acetate, dehydrated in a series of alcohol and propylene oxide, infiltrated, and embedded in EPON 812. Thin sections were stained with uranyl acetate and lead citrate prior to examination with a Philips 208S electron microscope.

### **Supplementary Material**

Refer to Web version on PubMed Central for supplementary material.

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#### Figure 1.

Biochemical Analyses of  $A\beta$  Levels in BRI- $A\beta$  Mice (A) Schematic of the BRI- $A\beta$  fusion proteins. BRI- $A\beta$  fusion constructs were designed to take advantage of the BRI protein that is cleaved by furin or a furin-like protease near the COOH-terminus to release a soluble 23 amino acid peptide in the wild-type protein.  $A\beta$ 1-40 or  $A\beta$ 1-42 was fused to the C terminus of the BRI protein at the furin cleavage site. This cleavage releases  $A\beta$  into the lumen or extracellular space, resulting in efficient secretion. (B) IP/MS analysis of  $A\beta$  isoforms in BRI- $A\beta$  mice. The spectra from the BRI- $A\beta$ 42 lines had single peaks detected with a mass/charge ratio (m/z) of 4515 corresponding to human  $A\beta$ 1-42. In BRI-40 mice, one peak was detected at m/z 4330 corresponding to human  $A\beta$ 1-40. The shoulder of the 4330 peak corresponds to the  $A\beta$ 40

sodium adduct ions. Peaks corresponding to human A $\beta$ 1-38, A $\beta$ 1-40, and A $\beta$ 1-42 were detected in extracts from Tg2576 and were absent in nontransgenic mice and BRI wild-type mice. (C) RIPA-soluble (RIPA) and (D) RIPA-insoluble formic acid extractable (FA) A $\beta$  levels in hemibrains (n = 3-7 mice per age point per group) from BRI-A $\beta$  mice were quantified by A $\beta$  sandwich ELISAs. FA A $\beta$ 42 increased dramatically with age in the BRI-A $\beta$ 42 mice. There was no evidence for accumulation of insoluble A $\beta$ 40 in the BRI-A $\beta$ 40 mice with age.



#### Figure 2.

Amyloid Pathology in BRI-Aß Mice Progression of amyloid deposition (total Aß immunostaining with 33.1.1 raised against A\beta1-16) in the cerebellum of BRI-A\beta42A mice at 3 months (A), 6 months (B), and 16.5 months (C). BRI-Aβ42 mice develop both diffuse (arrow in [C]) and compact thioflavin S-positive deposits (D). Plaques were immunoreactive with Aβ42 end-specific antibodies (E). No pathology was evident even in 16.5-month-old BRI-Aβ40 mice, immuno-stained with a panel of Aβ antibodies or thioflavin S (F). Old (19 month) BRI-Aβ42 mice show widespread forebrain Aβ deposits, particularly in the molecular layer of the hippocampus (G) and the entorhinal/piriform cortex (H). The entorhinal  $A\beta$  deposits were largely diffuse, and neurons were often observed within these diffuse deposits (I). Increased gliosis was associated with plaque deposition (J). Sections were immunostained with anti-GFAP (brown reaction product) followed by immunostaining for AB using the antibody BAM-10 (black reaction product). CAA was progressive and severe in old mice (K); vessels were stained with thioflavin S. A dense cored plaque is also visible. The vascular deposits were fibrillar, as demonstrated by electron microscopy (see Figure 3). (L) wild-type BRI mice had no amyloid pathology (total A $\beta$  immunohistochemistry) at ages up to 24 months of age. Scale bars: 60 µm (A—F), 200 µm (G), 300 µm (H), 20 µm (I), 30 µm (J).

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#### Figure 3.

Electron Microscopy of Congo-philic Amyloid Angiopathy and A $\beta$  Plaques in BRI-A $\beta$ 42 Mice CAA was pronounced in BRI-A $\beta$ 42 mice. (A) Amyloid deposit (amy) at the outer basal lamina (marked with \*) of an arteriole found at the cortical surface. An interendothelial junction is indicated by the double arrowhead. The boxed area in (A) is enlarged in (B). (B) Note the disruption of basal lamina (bl) between the astrocyte (ast) and smooth muscle cells (smc) by bundles of amyloid fibrils (amy). Arrowhead points to surface vesicles of SMC. Many of the extracellular amyloid plaques in the BRI-A $\beta$  mice had dense amlyoid aggregates with radiating fibrils (C). There were many dystrophic neurites (arrow-heads) present at the periphery of these plaques. vl, lumen; e, endothelium; SMC, smooth muscle cell; Ast, astrocytes. Scale bars: 1  $\mu$ m (A), 30 nm (B), and 5  $\mu$ m (C).



#### Figure 4.

Mouse A $\beta$  and Modified A $\beta$  Peptides Are Present in the Plaques of BRI-A $\beta$ 42 Mice (A) Senile plaques in the temporal gyrus from an AD brain, Braak stage VI, immunostained with anti-A $\beta$ 1-16 (33.1.1). (B) A serial section of the AD tissue in (A) immunostained with anti-rodent A $\beta$  that does not detect human A $\beta$  deposited in AD. (C) Total A $\beta$  immunostaining using antibody 33.1.1, in the cerebellum of a 17-month-old BRI-A $\beta$  mouse shows multiple cored plaques, a subset of these plaque cores are immunopositive for rodent A $\beta$  in a serial section (D). Arrows in (C) and (D) indicate the same plaques immunostained with 33.1.1 and rodent A $\beta$ , respectively. (E) Diffuse and dense cored plaques in the cerebellum were immunopositive with an anti-A $\beta$ 1-16 (82E1, IBL), a subset of these plaques were stained by an antibody that recognizes a truncated N-terminal modified pyrogluta-mate form of A $\beta$  (N3pE A $\beta$ ) (F). Arrows in (E) and (F) indicate the same plaques immunostained with anti-A $\beta$ 1-16 and anti-N3pE A $\beta$ , respectively. Scale bars: 200 µm (A—D) and 60 µm (E and F). McGowan et al.



#### Figure 5.

Enhanced Pathology in BRI-A $\beta$ 42 × Tg2576 Bigenic Mice Transgenic mice expressing both BRI-A $\beta$ 42 and mutant APP<sub>swe</sub>(Tg2576) have enhanced senile plaque pathology in the forebrain compared with their singly transgenic littermates and significantly elevated insoluble A $\beta$  levels in the forebrain and hindbrain. Entorhinal/piriform cortex of 14.5-month-old BRI-A $\beta$ 42A mouse (A), 14.5-month-old Tg2576 (B), 14.5-month-old BRI-A $\beta$ 42 × Tg2576 bigenic mouse (C) immunostained with anti-total A $\beta$  (33.1.1). All mice were littermates. FA A $\beta$  levels in the forebrain (D) and hindbrain (E) of an aging series of bigenic BRI-A $\beta$ /Tg2576 compared with their single transgenic littermates. Scale bars: 300 µm (A—C).