Neuronal heparan sulfates promote amyloid pathology by modulating brain amyloid-β clearance and aggregation in Alzheimer’s disease

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Accumulation of amyloid-β (Aβ) peptide in the brain is the first critical step in the pathogenesis of Alzheimer’s disease (AD). Studies in humans suggest that Aβ clearance from the brain is frequently impaired in late-onset AD. Aβ accumulation leads to the formation of Aβ aggregates, which injure synapses and contribute to eventual neurodegeneration. Cell surface heparan sulfates (HSs), expressed on all cell types including neurons, have been implicated in several features in the pathogenesis of AD including its colocalization with amyloid plaques and modulatory role in Aβ aggregation. We show that removal of neuronal HS by conditional deletion of the Ext1 gene, which encodes an essential glycosyltransferase for HS biosynthesis, in postnatal neurons of amyloid model APP/PS1 mice led to a reduction in both Aβ oligomerization and the deposition of amyloid plaques. In vivo microdialysis experiments also detected an accelerated rate of Aβ clearance in the brain interstitial fluid, suggesting that neuronal HS either inhibited or represented an inefficient pathway for Aβ clearance. We found that the amounts of various HS proteoglycans (HSPGs) were increased in postmortem human brain tissues from AD patients, suggesting that this pathway may contribute directly to amyloid pathogenesis. Our findings have implications for AD pathogenesis and provide insight into therapeutic interventions targeting Aβ-HSPG interactions.

INTRODUCTION

Alzheimer’s disease (AD) is the most common form of dementia in which amyloid plaques and neurofibrillary tangles are the major pathological hallmarks. Mounting evidence suggests that the accumulation and aggregation of amyloid-β (Aβ), the major component of amyloid plaques in the brain, is a key initiating event in the pathogenesis of AD (1, 2). Aβ is generated from proteolytic processing of amyloid precursor protein (APP) by β- and γ-secretases (3, 4). Genetic and biochemical studies showed that early-onset forms of AD (<1%) are caused by the inheritance of autosomal-dominant mutations that affect APP processing, leading to increases in Aβ production or the propensity for Aβ to aggregate (5). However, much less is known about the pathological mechanisms that modulate Aβ accumulation in the late-onset forms of AD, which account for more than 99% of cases. The amount of Aβ in the brain represents the net balance of Aβ production and clearance (6). Excess Aβ rapidly aggregates to form oligomers, which have been shown to impair synaptic function and eventual cognitive deficits (7). Aβ has a relatively short half-life in the brain, with ~1 to 2 hours in mouse interstitial fluid (ISF) and ~8 hours in human cerebrospinal fluid (CSF) (8, 9). Increasing evidence indicates that impaired Aβ clearance is a common prelude to late-onset AD (10). Thus, understanding the factors that regulate Aβ clearance is critical to illuminate the pathogenic pathways of AD and to design effective therapies for treating AD.

Heparan sulfate proteoglycans (HSPGs), consisting of heparan sulfate (HS) chains covalently attached to a specific protein core, are abundant cell surface and extracellular molecules that interact with a spectrum of ligands (11, 12). A critical step in the biosynthesis of HS is the elongation of linear polysaccharides composed of alternating glucuronic acid (GlcA) and N-acetylgalactosamine (GlcNAc) catalyzed by the EXT1 family of molecules (13). Studies have shown that the biosynthesis of HS is disrupted upon EXT1 deficiency, indicating that the glycosyltransferase activity associated with EXT1 is indispensable for HS assembly (14, 15). HSPGs are ubiquitously expressed in almost all mammalian cell types and regulate a wide variety of biological processes, including embryonic development, growth factor signaling, cell proliferation, adhesion and migration, and homeostasis (12, 13). HSPGs are present in senile plaques and cerebral amyloid angiopathy (CAA) (16–19). HSPGs have been shown to bind to Aβ and accelerate its oligomerization and aggregation (20–22). Also, HS mediates cellular Aβ uptake, neurotoxicity, and inflammatory responses induced by Aβ (23–25). These findings suggest that HSPGs might play important roles in Aβ metabolism and the pathogenesis of AD.

A recent study reported that overexpression of heparanase reduces the amyloid burden in a mouse model, likely by modulating Aβ deposition (26). Here, using a conditional mouse model deficient in neuronal HS in adult brain, we bypassed the developmental effects of HS and investigated the cell type–specific roles of HS in amyloid pathogenesis. Our findings provide in vivo evidence that neuronal HS inhibits its brain Aβ clearance and promotes amyloid plaque deposition. Specifically, by using in vivo microdialysis to measure the clearance rate of soluble Aβ from the brain ISF, we found that deficiency of HS in neurons facilitated Aβ clearance and reduced Aβ aggregation, resulting in a reduction in amyloid plaque deposition. In addition, we showed that several HSPG species were up-regulated in postmortem human brain tissues from AD patients. Together, our results shed light on the mechanisms by which HSPGs modulate brain Aβ metabolism.
and deposition and suggest that targeting Aβ-HSPG interactions might be a promising strategy to treat AD.

RESULTS

Neuronal HS deficiency decreases Aβ and ameliorates amyloid pathology in an amyloid mouse model

The amount of Aβ in the brain is a net balance of Aβ production and clearance; thus, it is important to analyze the overall impact of altered HSPG expression on brain Aβ and Aβ pathology. To investigate the in vivo role of HSPG in brain Aβ metabolism, we conditionally disrupted the expression of the Ext1 gene encoding the HS polymerizing enzyme in neurons by crossing Ext1<sup>fl<sub>ox</sub>/fl<sub>ox</sub></sup> mice (27) with α-calcium/cammodulin-dependent protein kinase II (CaMKII-Cre) mice (28), and then, we further bred these mice to the APP<sup>swe</sup>/PS1ΔE9 (APP/PS1) amyloid mouse (29). The use of CaMKII-Cre allowed us to bypass the effects of Ext1 inactivation on embryonic brain development and to selectively study the role of HSPGs in the forebrain neurons of adult mice, which is one of the most vulnerable brain regions in AD. These mice developed normally without detectable morphological changes in the brain, consistent with previous findings (30). We found that Ext1 expression was decreased in the cortex of knockout mice at 12 months of age, but not in the cerebellum where CaMKII-Cre was not active (fig. S1A). Biochemical analysis showed that HS was significantly reduced in regions where CaMKII-Cre was active, such as cortex and hippocampus, but was unaltered in the cerebellum (fig. S1B). The residual HS observed likely represents the expression within glial cells or brain vasculature (17, 31, 32). As APP/PS1 mice develop amyloid plaques at 5 to 6 months of age, which continue to increase up to 12 months of age (33), we analyzed the burden of amyloid deposition in these mice at 12 months of age. The brain sections of the compound mutants (APP/PS1; Ext1<sup>fl<sub>ox</sub>/fl<sub>ox</sub></sup>; CaMKII-Cre; thereafter, APP/PS1; nExt1<sup>CKO</sup>) and their littermate controls (APP/PS1) were immunostained using an anti-Aβ antibody (34), and the extent of Aβ deposition in the cortical and hippocampal regions was quantified. The amyloid deposition was markedly decreased in APP/PS1; nExt1<sup>CKO</sup> mice (fig. 1A). Quantification revealed that Aβ burden in 12-month-old APP/PS1; nExt1<sup>CKO</sup> mice was about one-third of that in APP/PS1 littermate control mice (fig. 1A). Previous studies demonstrated that HS and HSPGs colocalize with amyloid plaques, in particular in the core of amyloid deposits and blood vessels (26, 32, 35, 36). We thus examined the association of HS or HSPG core proteins with amyloid plaques in our mouse model. Immunohistochemical staining showed that HS was colocalized with Congo red-positive amyloid plaques in the brain parenchyma and leptomeningeal arteries in APP/PS1 mice with heparinase III treatment abolishing HS immunoreactivity (fig. S1, C and D). As expected, the association between HS and amyloid plaques was markedly decreased in APP/PS1; nExt1<sup>CKO</sup> mice in the cortex, but was not altered in the brain vessels where CaMKII-Cre is not active (fig. S1, C and D). Deficiency of neuronal HS side chains did not alter the mRNA expression of HSPG protein backbones, including transmembrane syndecans, glycosyl-phosphatidylinositol (GPI)–linked glypicans, and extracellular matrix HSPGs (perlecan and agrin), in the brains of APP/PS1; nExt1<sup>CKO</sup> mice (fig. S2). About 20 to 40% of plaques were positive for glypican-1, syndecan-3, and agrin, whereas the colocalization between perlecan and Aβ was minimal (fig. S3A). We observed a reduction in the percentage of amyloid plaques that were positive for various HSPG subtypes (glypican-1, syndecan-3, and agrin) in the cerebral cortex of APP/PS1; nExt1<sup>CKO</sup> mice compared to control mice (fig. S3, A and B). Agrin and perlecan, the major HSPGs in the basement membrane of the cerebral vasculature (35, 37), and syndecan-3 were also colocalized with Aβ in the leptomeningeal arteries of APP/PS1 and APP/PS1; nExt1<sup>CKO</sup> mice (fig. S3C).

To further analyze how HS affects the dynamic pools of Aβ, we fractionated cortical and hippocampal mouse brain tissues into tris-buffered saline (TBS)–soluble, detergent-soluble (TBSX), and detergent-insoluble (guanidine-HCl, GDN) fractions (38), and quantified Aβ by enzyme-linked immunosorbent assay (ELISA). Although the majority of Aβ in these mice at 12 months of age was distributed in the amyloid plaques and was fractionated in the detergent-insoluble fractions, Aβ40 and Aβ42 in the TBSX-soluble fractions were also lower in the brains of APP/PS1; nExt1<sup>CKO</sup> mice compared with APP/PS1 controls (P < 0.01 in the cortex; P < 0.05 for Aβ40 and P < 0.01 for Aβ42 in the hippocampus) (Fig. 1, B and C; table S2). Consistent with decreased Aβ deposition, the concentrations of insoluble Aβ40 and Aβ42 in the guanidine fractions were very high in the APP/PS1 mice in the cortex and hippocampus and were reduced in both brain regions of APP/PS1; nExt1<sup>CKO</sup> mice (P < 0.01; fig. 1, D and E; table S2). Oligomeric Aβ has been implicated as the deleterious form of Aβ peptide (39); thus, we examined the amount of soluble oligomeric Aβ in the TBS-soluble fraction using an ELISA specific to oligomeric Aβ (40) in the cortex of these mice. We found that Aβ oligomers were significantly decreased in APP/PS1; nExt1<sup>CKO</sup> mice compared to APP/PS1 control mice (P < 0.01; fig. 1F). To investigate whether deficiency of HS affected Aβ oligomerization, we further assessed the ratio of Aβ oligomers versus total Aβ in the TBS-soluble fraction of the cortex. To quantify total Aβ, the same antibody used in oligomeric Aβ ELISA was used to capture Aβ followed by an antibody that recognizes the middle region of Aβ. The ratio of Aβ oligomers/total Aβ was significantly reduced in APP/PS1; nExt1<sup>CKO</sup> mice (P < 0.01; Fig. 1G, indicating a decrease of Aβ oligomerization in APP/PS1; nExt1<sup>CKO</sup> mouse brain. This result suggests that neuronal HS may promote Aβ oligomerization. HSPGs have been shown to promote Aβ aggregation and fibril formation (16, 26, 41). Thus, we next characterized amyloid plaque load in APP/PS1 and APP/PS1; nExt1<sup>CKO</sup> mice by staining brain sections with thioflavin S, a fluorescent dye that binds to amyloid fibrils. Consistent with the Aβ immunostaining pattern, we found that deficiency of neuronal HS led to a significant reduction in fibrillar plaques in the cortex (P < 0.01) and hippocampus (P < 0.05) of APP/PS1; nExt1<sup>CKO</sup> mice compared to APP/PS1 controls (fig. 1H). Together, our results demonstrate that the deficiency of HS in neurons reduced Aβ accumulation and aggregation, resulting in a decrease in amyloid plaque deposition in APP/PS1; nExt1<sup>CKO</sup> mice.

Neuronal HS deficiency decreases neuroinflammation in an amyloid mouse model

Abnormal activation of astrocytes and microglia has been observed in the brains of AD patients and APP transgenic mouse models (42). To examine the extent of astrogliosis in these mice, we immunostained the brain sections with anti–glial fibrillary acidic protein (GFAP) antibody followed by quantification. The immunostaining for GFAP clearly demonstrated that activation of astrocytes was suppressed in the brains of APP/PS1; nExt1<sup>CKO</sup> mice compared with those of APP/PS1 mice at 12 months of age (P < 0.01; Fig. 2, A to C, and fig. S4).
Brain sections were also immunostained with anti-ionizing calcium-binding adaptor molecule 1 (Iba1) antibody to examine the extent of microgliosis. The dystrophic microglial reactivity was more evident in APP/PS1 control mice, in which a stronger intensity and activated morphology of microglia were observed ($P < 0.01$; Fig. 2, D to F, and fig. S4). These results indicated that elimination of HS in neurons led to a reduction in plaque-associated microglial activation. Western blot analysis also confirmed that GFAP was significantly decreased in the cortex ($P < 0.01$) and hippocampus ($P < 0.05$) of APP/PS1; nExt1CKO mice (Fig. 2G).

Fig. 1. Neuronal HS deficiency reduces amyloid deposition. (A) Brain sections from control (APP/PS1) and neuronal HS-deficient (APP/PS1; nExt1CKO) mice ($n = 6$ to $10$ per group) at 12 months of age were immunostained with a pan-Ab antibody. Scale bar, 1 mm. The percentage of area covered by plaques was quantified, and the plaque load was normalized to that of APP/PS1 mice. ctrl, control. (B to E) The cortical and hippocampal brain tissues of APP/PS1 and APP/PS1; nExt1CKO mice ($n = 7$ to 12 per group) at 12 months of age were fractionated into TBS-soluble, detergent-soluble (TBSX), and detergent-insoluble (guanidine-HCl, GDN) fractions. The amount of Ab40 and Ab42 in TBS (B), TBSX (C), and GDN (D and E) fractions was quantified by ELISA. (F) Soluble oligomeric Ab in the cortex of APP/PS1 and APP/PS1; nExt1CKO mice ($n = 10$ to 12 per group). (G) Ratio of soluble oligomeric Ab versus total Ab in the TBS-soluble fraction in the cortex of APP/PS1 and APP/PS1; nExt1CKO mice ($n = 10$ to 12 per group). (H) Quantification of thioflavin S–positive amyloid plaques in the cortex and hippocampus of APP/PS1 and APP/PS1; nExt1CKO mice ($n = 13$ per group) at 12 months of age. Scale bar, 100 μm. Values represent means ± SEM. N.S., not significant; *$P < 0.05$; **$P < 0.01$. Statistical analysis was performed using Student’s t test.
Neuronal HS deficiency increases Aβ clearance in the hippocampus of amyloid model mice

To investigate the mechanism underlying the reduction of Aβ pathology, we used in vivo microdialysis to dynamically assess ISF Aβ metabolism in the hippocampus of APP/PS1; nExt1<sup>1CKO</sup> mice and APP/PS1 littermates at 3 to 4 months of age. Soluble Aβ in ISF has been shown to reflect total soluble Aβ in extracellular pools and is significantly correlated with the amount of Aβ that eventually is deposited in the extracellular space of the brain (8, 44, 45). Hippocampal ISF was sampled in APP/PS1; nExt1<sup>1CKO</sup> mice and APP/PS1 littermates for a stable baseline period, during which mice were able to freely move throughout the experiment. To understand whether the decrease in Aβ in APP/PS1; nExt1<sup>1CKO</sup> mice was the result of altered Aβ clearance from the ISF, we infused a potent γ-secretase inhibitor directly into the mice to rapidly block Aβ production, thus allowing sensitive measurement of the elimination rate of Aβ from the ISF (Fig. 3A and fig. S5A). ISF Aβ<sub>40</sub> and Aβ<sub>42</sub> concentration gradually decreased in a time-dependent manner after treatment with the γ-secretase inhibitor, with APP/PS1; nExt1<sup>1CKO</sup> mice showing a faster decline compared with control APP/PS1 mice (Fig. 3B and fig. S5B). The half-life (t<sub>1/2</sub>) of ISF Aβ<sub>40</sub> (P < 0.01) and Aβ<sub>42</sub> (P < 0.05) in the hippocampus of APP/PS1; nExt1<sup>1CKO</sup> mice was significantly shorter compared to that of APP/PS1 mice (Fig. 3C and fig. S5C). These results indicate that elimination of HS in neurons enhanced the clearance of soluble Aβ from the ISF.

Fig. 2. Deficiency of neuronal HS leads to reduced neuroinflammation. (A to C) Brain sections from APP/PS1 and APP/PS1; nExt1<sup>1CKO</sup> mice at 12 months of age were immunostained with GFAP antibody. Scale bar, 1 mm. (B) Representative images of GFAP staining in the cortex and hippocampal CA1 region. Scale bar, 100 µm. (C) Stained sections were scanned on the Aperio slide scanner and analyzed using the ImageScope software. The percentage of areas covered by GFAP staining in the cortex (n = 11 to 13 per group) and hippocampus (n = 7 to 10 per group) was quantified. (D to F) Brain sections from APP/PS1 and APP/PS1; nExt1<sup>1CKO</sup> mice at 12 months of age were immunostained with Iba1 antibody. Scale bar, 1 mm. (E) Representative images of Iba1 staining in the cortex and hippocampal CA1 region. Scale bar, 100 µm. (F) The percentage of area covered by Iba1 staining was quantified (n = 4 per group). (G) Amount of GFAP in the cortex (n = 9 per group) and hippocampus (n = 8 to 9 per group) of APP/PS1 and APP/PS1; nExt1<sup>1CKO</sup> mice examined by Western blot. (H) Amounts of TNF-α, IL-1β, and IL-6 in the cortex of APP/PS1 and APP/PS1; nExt1<sup>1CKO</sup> mice (n = 6 to 8 per group) evaluated by real-time PCR. Data represent means ± SEM. *P < 0.05; **P < 0.01. Statistical analysis was performed using Student’s t test.
To examine whether Ext1 inactivation in neurons affects APP processing, total APP and APP processing products in both APP/PS1 and APP/PS1; nExt1\(^{CKO}\) mice (\(n = 8\) per group) were analyzed. We found that there were no significant differences in the amount of full-length APP, soluble forms of APP (sAPP\(\alpha\) and sAPP\(\beta\)), and C-terminal fragments (CTFs) between APP/PS1 and APP/PS1; nExt1\(^{CKO}\) mice at 12 months of age examined by Western blot analysis (Fig. 3, D to F), indicating that deficiency of neuronal HS in these mice did not significantly affect APP processing.

Given that HS deficiency in neurons led to decreased insoluble A\(\beta\) and a substantial reduction in amyloid plaque deposition without affecting APP processing, we also analyzed the mRNA expression of neprilysin, insulin-degrading enzyme, matrix metalloproteinase 2 (MMP2), and MMP9, which are major A\(\beta\)-degrading enzymes in the brain. The quantitative reverse transcription polymerase chain reaction (qRT-PCR) results showed no significant differences in the mRNA expression of these enzymes between APP/PS1 and APP/PS1; nExt1\(^{CKO}\) mice (\(n = 7\) to 10 per group) at 12 months of age. Densitometric quantification is expressed as mean ± SEM. (G) mRNA of insulin-degrading enzyme (IDE), neprilysin (NEP), MMP2, and MMP9 in the cortex of APP/PS1 and APP/PS1; nExt1\(^{CKO}\) mice (\(n = 8\) per group) evaluated by real-time PCR. Statistical analysis was performed using Student’s t test.

**Deficiency of neuronal HSPGs increases the formation of CAA**

Given that A\(\beta\) clearance was enhanced in the brains of APP/PS1; nExt1\(^{CKO}\) mice, we thus examined whether deficiency of HS in neurons accelerated A\(\beta\) clearance without affecting APP processing or the expression of major A\(\beta\)-degrading enzymes in amyloid model mice. Deficiency of neuronal HSPGs increases the formation of CAA because A\(\beta\) deposition (including both A\(\beta\)40 and A\(\beta\)42) in cortical vessels manifested as CAA was increased (\(P < 0.01\)) in the APP/PS1; nExt1\(^{CKO}\) mice, even though the parenchymal amyloid
plagues were substantially decreased in these mice compared to APP/PS1 controls (Fig. 4, A and B, and fig. S6). Coo staining of α-smooth muscle actin (αSMA), a marker for vascular smooth muscle cells, and Aβ confirmed that Aβ was deposited in walls of arterioles in CAA (Fig. 4C). Mounting evidence suggests that the entrapment of brain Aβ in the ISF perivascular lymphatic drainage pathway is the main cause of CAA (46). Given that deficiency of neuronal HS increased Aβ clearance and reduced amyloid deposition in the brain parenchyma, relatively more Aβ might be diverted into perivascular drainage pathways for clearance in APP/PS1; nExt1CKO mice. Because several HSPGs have been shown to be expressed by cells of the cerebral vasculature and are involved in the pathogenesis of CAA (17, 36, 47), the increased cerebrovascular amyloid might be attributed to the HSPGs in the cerebral vasculature.

Several classes of HSPGs are increased in human AD brain tissues

Several HSPGs have been shown to colocalize with senile plaques and CAA (31, 48). To examine whether the distributions of different classes of HSPGs are altered in the pathogenesis of AD, we examined the amounts of HSPGs in the temporal cortex of human control (n = 20; average age, 85.1 ± 5.7 years) and AD (n = 20; average age, 84.4 ± 5.2 years) brain tissues (table S1). We processed the brain tissues into TBS, detergent-soluble (TBSX), and detergent-insoluble (GDN) fractions, and the amounts of syndecans, glypicans, perlecan, and agrin were examined by specific ELISAs. We found that the amount of syndecan-1 was not different between control and AD brain tissues (Fig. 5A). The amounts of syndecan-3 and syndecan-4 in both the detergent-soluble (P < 0.01 for syndecan-3; P < 0.05 for syndecan-4) and detergent-insoluble fractions (P < 0.05 for syndecan-3; P < 0.05 for syndecan-4) were significantly increased in human AD brains (Fig. 5, B and C). In addition, glypcan-3 in the GDN fraction was elevated (P < 0.05) in AD brains, although there was no change in the TBSX fraction (Fig. 5D). Glypcan-1, an abundant glypican in neurons, was trending to an increase in the GDN fractions, although this was not significant (Fig. 5E). We next examined the amounts of secreted extracellular matrix HSPGs, including agrin and perlecan, in these human brain tissues and found that the amount of agrin was increased in the GDN fractions of AD brains (Fig. 5F; P < 0.01). A previous study showed that a large fraction of the agrin in AD brains is detergent-insoluble (49). In our measurements, the amount of agrin in the TBSX fractions was under the detection limit using currently available ELISAs. Finally, the amount of perlecan was elevated in both detergent-soluble (P < 0.01) and detergent-insoluble fractions (P < 0.05) in AD brain tissues (Fig. 5G). Together, these findings demonstrate that a number of HSPGs, distributed abundantly in the insoluble compartments, were significantly increased in human AD brains.

DISCUSSION

The accumulation and deposition of Aβ in the brain have been hypothesized to drive the pathogenic cascades of AD (1, 2). Substantial amyloid buildup is present in AD brains long before the clinical onset of the disease (50, 51). Mounting studies have clearly demonstrated the causes of Aβ accumulation in early-onset familial AD patients as autosomal dominant gene mutations lead to increased Aβ42 production (52, 53). However, most AD cases are sporadic and late-onset, and the pathological mechanisms that lead to Aβ accumulation are less clear. HSPGs are glycoproteins composed of a core protein coupled to one or more HS glycosaminoglycan chains (12). Five distinct classes of HSPGs have been identified, including membrane HSPGs (such as syndecans and glypicans) and the secreted extracellular matrix HSPGs (agrin, perlecan, and type XVIII collagen) (54). HSPGs have been shown to interact with a variety of molecules because of their highly sulfated nature, thereby mediating biological activities including endocytosis and cell signaling (12, 55, 56). Here, we have provided in vivo evidence that neuronal depletion of HS leads to reduced Aβ and amyloid deposition in APP/PS1 amyloid model mice. Deficiency of neuronal HS did not affect APP processing and Aβ production but did enhance Aβ clearance. These results suggest that the association of Aβ and HSPGs on the surface of neurons likely suppresses or represents an inefficient pathway for Aβ.

![Fig. 4. Deletion of neuronal HS increases the abundance of CAA along the cerebral vasculature.](http://stm.sciencemag.org/)

(A) Aβ deposition in brain sections from control APP/PS1 and APP/PS1; nExt1CKO mice at 12 months of age was immunostained with a pan-Aβ antibody. Scale bar, 200 μm. (Inset) Immunostaining of Aβ deposition along leptomeningeal arteries as CAA in control APP/PS1 and APP/PS1; nExt1CKO mice. Scale bar, 10 μm. (B) The burden of CAA formation in leptomeningeal arteries in control APP/PS1 and APP/PS1; nExt1CKO mice (10 arteries per mouse; 10 mice per genotype) was quantified after scanning Aβ immunostaining by the Positive Pixel Count program (Aperio Technologies). Data represent means ± SEM. **P < 0.01. Statistical analysis was performed using Student’s t test. (C) Coimmunofluorescence staining of leptomeningeal arteries with αSMA (green) and Aβ (red) antibodies. Scale bar, 10 μm.

clearance (fig. S7). Also, neuronal HSPGs might facilitate Aβ oligomerization and aggregation (fig. S7). Aβ40 was shown to inhibit the heparanase-mediated degradation of HS (31, 57). Thus, the increases of Aβ and HS may result in a vicious cycle that further contributes to the persistence and stability of amyloid plaques in AD brains.

Aβ is mainly generated in neurons and eliminated from the brain through several clearance pathways (6, 46, 58–60); one of the major pathways depends on enzymatic degradation in ISF by proteases such as neprilysin and insulin-degrading enzyme (61). Thus, the interactions of Aβ with other molecules are predicted to interfere with the enzymatic degradation machineries by physically blocking their association sites.

HSPGs have been proposed to function as a chaperone that serves as a protective shield against the proteolytic degradation of Aβ; thus, the clearance of Aβ is likely hindered in the presence of cell surface HSPGs (fig. S7) (47, 62). Consistent with this notion, our results demonstrated that deficiency of neuronal HS accelerates Aβ elimination in the mouse hippocampus. We found that deletion of HS in neurons exacerbated CAA formation despite the reduction of Aβ and amyloid plaque deposition in the brain parenchyma. Given that significant portions of ISF Aβ flow along the cerebral vasculature and are cleared into the blood flow through the blood-brain barrier (BBB) or the perivascular drainage pathway (63, 64), Aβ likely is deposited on cerebrovasculature as CAA when these pathways are disturbed or the amount of drained Aβ overwhelms the clearance capacity. Our results suggest that the expression of HSPGs in various brain compartments might play an important role in regulating the distribution of Aβ in AD brains.

HSPGs have also been suggested to function as a catalyst to promote Aβ oligomer formation and aggregation (23, 62, 65). In vitro experiments

**Fig. 5. Several classes of HSPGs are elevated in human AD brain tissues.** Human temporal lobe brain tissues from control (Ctrl) (n = 20) and AD (n = 20) groups were lysed and fractionated through sequential extraction with TBS, TBSX, and GDN. (A to C) Transmembrane HSPGs, including syndecan-1, syndecan-3, and syndecan-4, in TBSX and GDN fractions of control and AD brain tissues were quantified by specific ELISAs. (D and E) The amounts of the GPI-anchored HSPGs glypican-1 and glypican-3 in TBSX and GDN fractions of control and AD brain tissue were quantified by specific ELISAs. (F and G) The amounts of two major extracellular matrix HSPGs agrin and perlecan in GDN fractions of control and AD brains were quantified by specific ELISAs. Results are presented as means ± SEM. *P < 0.05; **P < 0.01.
have shown that HS accelerates Aβ aggregation through their direct interaction (20–22). Membrane-bound glypicanc-1 has been shown to interact with oligomerized Aβ in the detergent-insoluble glycosphingolipid-enriched compartment (20). Agrin, an extracellular matrix HSPG, has been found to bind to Aβ and accelerated its fibril formation in human AD brain (16). Indeed, several HSPGs colocalize with senile plaques in AD brains (31, 36). We also demonstrated that several subtypes of HSPGs, including syndecan-3, syndecan-4, glypicanc-1, and glypicanc-3, were significantly higher in AD human brain tissues, particularly in the pools of detergent-insoluble fractions compared to those in control individuals. These results suggest that aberrant amounts of HSPGs and their altered distribution in AD brains might contribute to Aβ aggregation and plaque formation, although further studies are needed. It is also possible that amyloid plaques capture HSPGs or promote the accumulation of HSPG-expressing glial cells, resulting in an increase of HSPGs in AD brains (32).

Receptor-mediated cellular uptake and subsequent lysosomal degradation are other critical pathways for Aβ clearance (64). Whereas microglia and astrocytes actively internalize and clear Aβ, neurons also play a role in cellular Aβ clearance. Because HSPG is one of the major players that capture Aβ on the cell surface of neurons for endocytosis (64, 66), the deficiency of HS in neurons seems to be disadvantageous for this clearance pathway. However, when excess amounts of Aβ are internalized, it can aggregate in lysosomes, likely providing “seeds” to initiate further Aβ aggregation (67, 68). In addition, Aβ has been shown to self-propagating and is transmitted from one neuron to another after its cellular uptake (69–71). Thus, although the neuronal uptake of Aβ through HSPGs can lead to trafficking to the lysosomes (25), albeit less efficiently than via the low-density lipoprotein receptor–related protein 1 (LRP1) (66), such a pathway might be harmful when the capacity for lysosomal degradation of Aβ is overwhelmed and compromised (fig. S7). It will be critical to investigate whether neuronal HSPGs mediate the propagation of Aβ during AD pathogenesis in future studies. Extracellular tau and α-synuclein have been shown to propagate after being endocytosed via cell surface HSPGs on neurons (56), which is likely to be a critical step for the progression of related neurodegenerative diseases (72). Thus, blocking the association of these pathogenic molecules with neuronal HSPGs may ameliorate the progression of various neurodegenerative diseases by preventing their propagation between neurons during disease development.

HSPGs interact with myriad molecules that orchestrate important biological functions (13). One limitation of our study is that we cannot rule out the possibility that disruption of HS binding to other molecules, in addition to Aβ, may also contribute to the modulation of Aβ metabolism in the adult mouse brain. One such molecule is apolipoprotein E (apoE), whose binding to HSPGs is known to contribute to the metabolism of apoE itself (64, 73). The ε4 allele of the APOE gene is the strongest genetic risk factor for sporadic AD among the three polymorphic alleles (ε2, ε3, and ε4). ApoE is required for seeding amyloid and has been shown to regulate Aβ metabolism and amyloid deposition in an apoE isoform–dependent manner (6, 74). Whether apoE is involved in HSPG-regulated Aβ metabolism/aggregation and whether this effect depends on apoE isoforms require further investigation.

In summary, our studies using genetically altered mice have clearly demonstrated a critical role of neuronal HS in brain Aβ clearance and oligomerization. Given that neuronal HSPGs are also likely involved in the pathogenic propagation of various proteins in neurodegenerative diseases and HSPGs are elevated in human brain tissues from AD patients, our studies establish a rationale for targeting the Aβ–HSPG pathway for therapy. Toward this end, it is noteworthy that the administration of a low–molecular weight heparin reduced amyloid pathology in a mouse model of AD, likely by antagonizing Aβ–HSPG interactions (65). Given that HSPGs interact with myriad critical molecules that mediate diverse biological activities (55), it will be critical to design and identify compounds that specifically inhibit the interaction between Aβ and HSPGs for the treatment of neurodegenerative diseases. When combined with other strategies targeting Aβ, including immunotherapy, these approaches might allow for a reduction, if not elimination, of Aβ-related toxicity.

MATERIALS AND METHODS

Study design

This study aimed to investigate the role of HSPGs in regulating brain Aβ metabolism using a conditional knockout mouse model. To accomplish this aim, the Ext1 gene, which encodes an essential glycosyltransferase for HS biosynthesis, was genetically ablated in the forebrain neurons of APP/PS1 mice. The effects of neuronal HS deficiency on the amount of Aβ, amyloid deposition, and neuroinflammation were assessed by Aβ ELISAs, immunohistochemistry, real-time PCR, and Western blot analyses. The neuronal HS–deficient APP/PS1 mice exhibited marked reduction in amyloid oligomerization and deposition. Using in vivo microdialysis, we also detected an accelerated rate of Aβ clearance within the brain ISF of HS-deficient mice, suggesting that neuronal HS might represent an inefficient or inhibitory pathway for Aβ clearance. We further examined the amounts of various HSPG species in postmortem human brain tissues from control and AD patients using specific ELISAs to evaluate the potential relevance of altered HSPGs in AD pathogenesis. The mice of different genotypes were selected on the basis of availability. Sample sizes were adequately powered to observe the effects on the basis of past experience of animal studies (66, 75). Data collection and the quantification of immunoreactivity for mouse samples were performed with the investigators unaware of the sample identities until statistical analyses.

Human postmortem brain tissues

Temporal lobe cortical samples from neurologically unimpaired subjects (n = 20) and from subjects with AD (n = 20) were obtained from the University of Kentucky Alzheimer’s Disease Center (ADC) Neuropathology Core (Lexington, KY) (75). After autopsy, the samples were snap-frozen in liquid nitrogen and kept frozen at −80°C. Informed consent was obtained through ADC Neuropathology Core, and we obtained institutional review board approval to use these brain samples for research. Diagnosis of AD was confirmed by pathological and clinical criteria as described (76). The average age of subjects was 85.1 ± 5.7 years in the control group and 84.4 ± 5.2 years in the AD group (P = 0.69). Average postmortem interval was 3.2 hours and was not significantly different between the two groups (P = 0.22).

Animals

All animal procedures were approved by the Mayo Clinic Institutional Animal Care and Use Committee and were in accordance with the National Institutes of Health Guide for the Care and Use of Laboratory Animals. Ext1flox/flox mice have been described in previous studies (27). Double transgenic APPswe/PS1ΔE9 (APP/PS1) mice were purchased from...
The Jackson Laboratory and were initially generated by Jankowsky et al. as described (29).

**In vivo Aβ microdialysis**

To assess ISF Aβ in the hippocampus of awake, freely moving APP/PS1 and APP/PS1; nExt^{CKO} mice, in vivo microdialysis was performed as previously described (8, 44). Briefly, under isoflurane volatile anesthetic, guide cannula (BR style; Bioanalytical Systems) was cemented into the hippocampus (3.1 mm behind bregma, 2.5 mm lateral to midline, and 1.2 mm below dura at a 12° angle). A microdialysis probe (38-kD molecular weight cutoff membrane; Bioanalytical Systems) was inserted through the guide cannula into the brain. Artificial CSF (1.3 mM CaCl₂, 1.2 mM MgSO₄, 3 mM KCl, 0.4 mM KH₂PO₄, 25 mM NaHCO₃, and 122 mM NaCl, pH 7.35) containing 4% bovine serum albumin (BSA) (Sigma) filtered through a 0.1-μm membrane was used as microdialysis perfusion buffer. Flow rate was constant 1.0 μl/min. Samples were collected every 60 to 90 min overnight, which gets through the 4- to 6-hour recovery period, and the mean concentration of Aβ over the 6-hour preceding treatment was defined as basal concentration of ISF Aβ. Samples were collected through a refrigerated fraction collector and assessed for Aβ40 or Aβ42 by ELISAs. For each animal, all Aβ concentrations were normalized to the basal Aβ concentration. To measure Aβ elimination half-life, mice were administered a γ-secretase inhibitor, LY411575 (5 mg/kg), intraperitoneally to rapidly block the production of Aβ. Microdialysis samples were collected every 60 min for 6 hours. The half-life of ISF Aβ was calculated on the basis of the slope of the semi-log plot of percent change in Aβ versus time (8).

**ELISA quantification**

ISF Aβ concentrations were assessed using sandwich Aβ40- or Aβ42-specific ELISAs as described (44). Briefly, a mouse anti-Aβ40 antibody HJ2 (anti-Aβ35–40) or Aβ42 antibody HJ7.4 (anti-Aβ37–42) was used as capture antibody, respectively. A biotinylated antibody HJ5.1 (anti-Aβ13–28) targeting the central domain of Aβ was used as the detecting antibody, followed by ELISA development using streptavidin poly-HRP40 (Fitzgerald Industries). Synthetic human Aβ40 or Aβ42 peptide (American Peptide) was used to generate the standard curves for each assay. The ELISA assays were developed using Super Slow ELISA TMB (Sigma) with absorbance read on a BioTek plate reader. The Aβ concentration in the brain lysates was determined by ELISA (77) with end-specific monoclonal antibody (mAb) 2.1.3 (human Aβ1–42–specific) and mAb 13.1.1 (human Aβ40–specific) for capture and horseradish peroxidase (HRP)–conjugated mAb Ab5 (human Aβ1–16–specific) for detection. The ELISAs were developed using Super Slow ELISA TMB (Sigma). To detect soluble oligomeric Aβ species, the same antibody, 3D6 (to Aβ residues 1 to 5), was used for both capture and detection similarly as previously described (40). Briefly, 3D6 (10 μg/ml) antibody was coated into 96-well immunoassay plates overnight at 4°C. The plates were then aspirated and blocked with 4% BSA in phosphate-buffered saline (PBS) buffer for at least 1 hour at 37°C. The samples and standards were added to the plates and incubated overnight at 4°C. The Aβ oligomers used as standards were prepared and characterized as previously described (78, 79). The plates were washed three or more times with wash buffer between each step of the assay. The plate was incubated with the biotinylated 3D6 (0.5 μg/ml) antibody in assay buffer (1% BSA, PBS) for 90 min at 37°C. The avidin-HRP (Vector Laboratories) was added to the wells for 90 min at room temperature. The ELISAs were developed using Super Slow ELISA TMB (Sigma). To measure total Aβ amount, 3D6 (10 μg/ml) and biotinylated mAb 266 (to Aβ residues 13 to 28, 0.5 μg/ml) were used as the capture and detection antibodies, respectively. All procedures were the same as those in oligomeric Aβ ELISA described above.

**Preparation of brain homogenates**

Mouse brain tissues and human postmortem brain tissues were processed through sequential extraction as described (38). Frozen brain tissues were homogenized with TBS and centrifuged at 100,000g for 60 min at 4°C with supernatant defined as TBS-soluble fraction. Pellets were resuspended in TBS buffer containing 1% Triton X-100 (TB SX) and mixed gently by rotation at 4°C for 30 min, followed by a second centrifugation at 100,000g for 60 min with supernatant defined as TB SX-soluble fraction. The TB SX-insoluble pellet was resuspended with 5 M guanidine, mixed by rotation at room temperature overnight, and centrifuged at 16,000g for 30 min with supernatant defined as GDN-soluble fraction.

**Immunohistochemical and immunofluorescence staining and analyses**

Paraffin-embedded sections were immunostained using pan-Aβ (Aβ 33.1.1; human Aβ1–16–specific), anti-GFAP (BioGenex), anti-Aβ40 (Millipore), anti-Aβ42 (21F12, provided by E. Lilly), and anti-Iba1 (Wako) antibodies (80). Immunohistochemically stained sections were captured using the ImageScope AT2 image scanner (Aperio Technologies) and analyzed using the ImageScope software. The intensities of GFAP and Iba1 staining in the hippocampus were calculated using the Positive Pixel Count program available with the ImageScope software (Aperio Technologies) (80). The fibrillar Aβ was immunostained with thioflavin S. The images were captured by Aperio Fluorescent Scanner, and the stained areas covered by the fibrillar plaque were quantified by ImageJ. For examining the CAA in the cerebral vasculature, paraffin-embedded sections were coated with αSMA antibody (Abcam) and Aβ antibody (MOAB2), followed by Alexa Fluor 488– and Alexa Fluor 568–conjugated secondary antibodies (Invitrogen). MOAB2 antibody was a gift from M. J. LaDu (University of Illinois at Chicago). The images were acquired by a confocal laser scanning fluorescence microscope (model LSM 510 invert, Carl Zeiss).

**Statistical analysis**

All quantified data represent an average of samples. Statistical analyses were performed with Excel or GraphPad Prism software. Statistical significance was determined by two-tailed Student’s t test. Levels of significance are as follows: *P < 0.05, **P < 0.01; P < 0.05 was considered significant.

**SUPPLEMENTARY MATERIALS**

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Materials and Methods

Fig. S1. Characterization of neuronal HS-deficient mice.

Fig. S2. The mRNA expression of HSPG subfamily in APP/PS1 and APP/PS1; nExt^{CKO} mice.

Fig. S3. HSPG core proteins codeposit with amyloid plaques in the brain parenchyma and leptomeningeal arteries.

Fig. S4. Deficiency of neuronal HS leads to a decrease in amyloid-associated astrogliosis.

Fig. S5. Deficiency of neuronal HS increases the clearance of ISF Aβ42 in the hippocampus of APP/PS1 mice.
REFERENCES AND NOTES


Neuronal heparan sulfates promote amyloid pathology by modulating brain amyloid-β clearance and aggregation in Alzheimer's disease

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Neuronal HSPGs lay a trap for amyloid

The accumulation of neurotoxic amyloid-β (Aβ) in the brain is a pathological hallmark of Alzheimer's disease (AD). Heparan sulfate proteoglycans (HSPGs) are abundant cell surface receptors that colocalize with amyloid plaques. Here, Liu and colleagues show that genetically engineered mice lacking heparan sulfates in forebrain neurons were protected from amyloid deposition because of a faster clearance of Aβ and reduction in Aβ aggregation. Also, the authors found that several HSPG species were increased in human AD postmortem brain tissue. These findings suggest that targeting Aβ-HSPG interactions might be an effective strategy for AD prevention and treatment.