### RESEARCH ARTICLE

# **Clusterin contributes to early stage of Alzheimer's disease pathogenesis**

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#### Keywords

Alzheimer's disease, amyloid pathology, apolipoprotein J, Aß oligomers, chaperone protein

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#### Abstract

While clusterin is reportedly involved in Alzheimer's disease (AD) pathogenesis, how clusterin interacts with amyloid- $\beta$  (AB) to cause AB neurotoxicity remains unclear in vivo. Using 5×FAD transgenic mice, which develop robust AD pathology and memory deficits when very young, we detected interactions between clusterin and AB in the mouse brains. The two proteins were concurrently upregulated and bound or colocalized with each other in the same complexes or in amyloid plaques. Neuropathology and cognitive performance were assessed in the progeny of clusterinnull mice crossed with 5×FAD mice, yielding  $clu^{-/-}$ ;5×FAD and  $clu^{+/+}$ ;5×FAD. We found far less of the various pools of AB proteins, most strikingly soluble AB oligomers and amyloid plaques in clu<sup>-/-</sup>;5×FAD mice at 5 months of age. At that age, those mice also had higher levels of neuronal and synaptic proteins and better motor coordination, spatial learning and memory than age-matched  $clu^{+/+}$ ;5×FAD mice. However, at 10 months of age, these differences disappeared, with AB and plaque deposition, neuronal and synaptic proteins and impairment of behavioral and cognitive performance similar in both groups. These findings demonstrate that clusterin is necessarily involved in early stages of AD pathogenesis by enhancing toxic AB pools to cause AB-directed neurodegeneration and behavioral and cognitive impairments, but not in late stage.

#### INTRODUCTION

Alzheimer's disease (AD) is a progressive neurodegenerative disease and the most common type of dementia in the elderly. Early investigation of the cause of AD supported the amyloid hypothesis of pathogenesis in which accumulation of amyloid- $\beta$  (A $\beta$ ), derived from proteolytic cleavage of amyloid precursor protein (APP), leads to neurodegeneration. Recent efforts have aimed to find risk factors other than A $\beta$  that may contribute to late-onset AD, which is the phenotype in most cases of the disease.

Clusterin, also known as apolipoprotein J (Apo J), has been identified as an important risk factor for late-onset AD by large-scale GWAS (17,24). Clusterin is a ubiquitously expressed glycoprotein and is present in very high levels in the brain. In patients with AD, clusterin levels are even higher in the brain (14,33,40), serum and cerebrospinal fluid (CSF) (8,13). Clusterin is found to colocalize with A $\beta$  in the amyloid deposits of brains of AD patients (8,22,34).

Clusterin, the first reported extracellular mammalian chaperone protein (20,42), binds to biologically active hydrophobic residues exposed on misfolded or aggregated Aß proteins (43,44), including toxic oligomers, to prevent further aggregation, render them susceptible to degradation, or suppress their toxicity. Accordingly, clusterin was shown to decrease A<sub>β</sub>-induced neurotoxicity in rodent hippocampal neurons (4) and attenuate the toxic effects of  $A\beta$  oligomers in a novel in vivo model of Caenorhabditis elegans (2). Other studies have demonstrated that Aβ-bound clusterin facilitates the clearance of Aß across the blood-brain barrier (BBB) (3,57). Aβ oligomers pre-incubated with clusterin induced less neuronal death and learning and memory impairment in rats compared with Aß oligomers without clusterin (7). These findings support protective mechanisms by which clusterin may inhibit AD pathogenesis.

However, increasing evidence suggests that clusterin also specifically interacts with  $A\beta$  oligomers to exacerbate their toxicity. When clusterin binds to toxic  $A\beta$  oligomers, it sequesters and stabilizes the oligomeric structures, interfering with their further aggregation into less toxic fibril aggregates (41). Clusterin binding has been shown to shield A<sub>β</sub> oligomers from enzymatic degradation (32,37,38). Clusterin augmented Aß toxicity in PC12 cells or organotypic mouse brain slice cultures in parallel with the formation of clusterin-induced and -stabilized Aß oligomers (25,40,41). Knockdown of clusterin in primary neurons reduced A $\beta$  toxicity (23), and the genetic ablation of clusterin  $(clu^{-/-})$  in PDAPP mice that develop AD pathology caused significantly fewer amyloid deposits and less neuritic dystrophy with no alteration of the total AB deposition (10). More interestingly, a clinical longitudinal study (11) found significant associations between levels of clusterin and  $A\beta$  in CSF and entorhinal cortex atrophy seen on MRI, implying that clusterin-A $\beta$  interaction is important in the earliest stages of AD pathogenesis.

In the present study, we evaluated the effect of genetic deletion of clusterin on amyloid pathogenesis and motor and cognitive behavioral performance in  $5 \times FAD$  transgenic mice, which co-express five familial hAPP/PS1 mutations and develop severe AD pathology at an early age (39). The results provide the first *in vivo* evidence that clusterin is an important endogenous factor facilitating the early stage of AD pathogenesis, but its influence is attenuated or lost in the late stage when amyloid deposition is already quite advanced.

#### **MATERIALS AND METHODS**

#### **Animal studies**

This animal study was approved by the Institutional Animal Care and Use Committee of the Asan Institute for Life Sciences, Asan Medical Center (Seoul, Korea).

Clusterin-deficient  $(clu^{-/-};5\times FAD)$  and their littermate  $5\times FAD$  mice  $(clu^{+/+};5\times FAD)$  were obtained by breeding clusterin-null mice  $(clu^{-/-})$  (10,35) with  $5\times FAD$  transgenic mice, which co-express hAPP<sub>695</sub> with three familial AD mutations (K670N/M671L, I716V and V717I) and presenilin (PS1) with two familial mutations (M146L and L286V) (39). Experiments were performed with 5- and 10-monthold male and female mice. All animals were bred and housed with free access to food and water at 22.0°C ± 1°C and with a 12-h light–dark cycle.

# Tissue preparation and histologic or immunohistochemical examination

Shortly after completing the behavioral tests described below, the animals were killed under deep anesthesia. The cerebral hemispheres were divided, snap-frozen in liquid nitrogen and immediately stored at  $-80^{\circ}$ C. Sagittal 12-µm thick sections of the left hemisphere were serially mounted on 1% poly-L-lysine-coated glass slides using a cryostat (HM550; Microm, Walldorf, Germany).

Brain A $\beta$  deposition was evaluated by immunohistochemistry using an anti-A $\beta$ (17–24) antibody (4G8; BioLegend, San Diego, CA, USA). The sagittal sections were fixed with 4% paraformaldehyde and briefly incubated in 70% formic acid (FA). Endogenous peroxidase activity in the tissue was depleted by treatment with 3% hydrogen peroxide, and the sections were blocked with 3% normal serum and 0.3% Triton X-100 in phosphate buffered saline (PBS) and reacted with the primary antibody (4G8; dilution, 1:200) and then with a biotinylated secondary antibody (Vector Laboratories, Burlingame, CA, USA). Finally, the sections were treated with avidin horseradish peroxidase (Vector Laboratories) and developed with 0.015% 3,3'-diaminobenzidine (DAB) and 0.001% hydrogen peroxide (Vector Laboratories). For immunofluorescent examination, the 4G8-immunoreacted tissues were visualized with Alexa Fluor-conjugated secondary antibodies (Thermo Fisher Scientific, Waltham, MA, USA).

To detect colocalization of clusterin and A $\beta$  in amyloid deposits, the brain tissues were immunologically co-stained with the primary antibodies, anti-A $\beta$  (4G8) and anti-clusterin/Apo J (Novus Biologicals, Littleton, CO, USA; dilution, 1:100) antibodies and then with two corresponding fluorescent secondary antibodies, Alexa Fluor 488- and 555-conjugated antibodies.

To examine congophilic compact amyloid plaques, brain sections were stained with Hematoxylin QS (a modification of Mayer's hematoxylin; Vector Laboratories) and Accustain<sup>®</sup> Congo red amyloid staining solution (Sigma-Aldrich, St. Louis, MO, USA) (26). After washing with absolute ethanol, the stained tissue sections were examined for congophilic plaques using a light or polarized microscope (Eclipse 800i; Nikon, Tokyo, Japan). For further analysis, microscopic images of stained tissues were captured with a DS-Fil digital camera connected to NIS-Elements image processing software (Nikon).

Because compact amyloid plaques emit blue fluorescence *in situ* under UV illumination (26), we examined blue fluorescent amyloid deposits in the brain tissues under a fluorescence microscope (Eclipse 800i) with a UV-2A filter (dichroic, 400 nm; excitation, 330–380 nm; barrier, 420 nm).

The load of amyloid deposits or plaques in the brain was expressed as the per cent area of 4G8-immunoreactive deposits or the number of congophilic plaques per mm<sup>2</sup> in regions of interest from the hippocampal, cortical and thalamic areas.

### Quantitative real-time polymerase chain reaction (PCR) assay

Total RNAs were extracted from the right hemispheres using the RNeasy Mini Kit (QIAGEN, Valencia, CA, USA) and cDNAs were generated using the iScript<sup>™</sup> cDNA synthesis kit (Bio-Rad, Hercules, CA, USA). The quantitative RT-PCR was conducted with the Bio-Rad CFX connect Real-time PCR machine using sense and antisense primers (250 nM each; Macrogen, Seoul, Korea) with 5× HOT FIREPol<sup>®</sup> EvaGreen<sup>®</sup> qPCR Supermix (Solis BioDyne, Tartu, Estonia). The sequences of sense and anti-sense primers used were as follows: for clusterin (Apo J; NM\_013492), 5'-CAG TTC CCA GAC GTG GAT TT-3' and 5'-TCC TGG CAC TTT TCA CAC TG-3', for Apo E (BC083351), 5'-TGT CCT GCA ACA ACA TCC AT-3' and 5'-TAT TAA GCA AGG GCC ACC AG-3', for Dickkopf Wnt signaling pathway inhibitor 1 (DKK-1; NM\_010051), 5'-CCT TCG GAG ATG ATG GTT GT-3' and 5'-GTT CTT GAT CGC GTT GGA AT-3' and for GAPDH (NM\_008084), 5'-AGG TCG GTG TGA ACG GAT TTG-3' and 5'-TGT AGA CCA TGT AGT TGA GGT CA-3', respectively. Finally, mRNA expression levels were calculated relative to that of GAPDH using the  $\Delta\Delta$ Ct method (27).

# Quantitative enzyme-linked immunosorbent assay (ELISA)

After the right brain hemispheres were weighed, they were homogenized in PBS (pH 7.4; 1:10, w/v) containing protease inhibitors (cOmplete<sup>TM</sup> Protease Inhibitor Cocktail; 1 tablet/50 mL; Roche Diagnostics, Mannheim, Germany) and centrifuged at 100 000 × g; the supernatant was then collected (the PBS-soluble fraction). The pellets were sonicated in 2% SDS in PBS, and the SDS-soluble fraction was collected by centrifugation at 100 000 × g. Finally, the SDS-insoluble pellets were solubilized in 70% FA and the FA-soluble fraction was collected by centrifugation at 100 000 × g.

We subjected the PBS-, SDS- and FA-soluble fractions to sandwich ELISA for quantification of human A $\beta$ 40 and A $\beta$ 42 (Thermo Fisher Scientific). Analysis of FA-soluble fractions was preceded by neutralization with 1 M Tris (adjusted to pH 11.0 using 5 M NaOH). In addition, the quantification of oligomeric A $\beta$ s (oA $\beta$ ) in PBS- and SDSsoluble fractions and A $\beta$  aggregates (aA $\beta$ ) in SDS extracts was performed using an Oligomeric A $\beta$  ELISA kit (BioSensis, Thebarton, SA, Australia) and an A $\beta$  aggregate ELISA kit (Thermo Fisher Scientific), respectively.

We also measured the amount of clusterin in the brain tissue (SDS extracts) using a SimpleStep<sup>®</sup> ELISA kit for mouse clusterin (Abcam, Cambridge, UK).

All measurements were taken at 450 nm using the Synergy H1 Hybrid microplate reader (BioTek, Winooski, VT, USA).

# Immunoblotting and co-immunoprecipitation analysis

The following primary antibodies were used for immunoblotting: anti-AB(1-16) (6E10; BioLegend; dilution, 1:2000), anti-clusterin/Apo J (Novus Biologicals; 1:1000), anti-apolipoprotein E (Apo E; Santa Cruz Biotechnology, Santa Cruz, CA, USA; 1:1000), anti-DKK-1 (Dickkopf-1; Boster, Pleasanton, CA, USA: 1:1000), anti-NeuN (Merck Millipore, Darmstadt, Germany; 1:500), anti-nonphosphorylated neurofilament protein SMI-32 (BioLegend; 1:2000), anti-synaptophysin (Santa Cruz Biotechnology; 1:5000), anti-ankyrin G (Santa Cruz Biotechnology; 1:2000), anti-PSD95 (postsynaptic density protein 95; Applied Biological Materials, Richmond, Canada; 1:2000), anti-VAMP

(vesicle-associated membrane protein)/synaptobrevin (Santa Cruz Biotechnology; 1:1000), anti-ProSAP2 (postsynaptic scaffold protein)/Shank 3 (Santa Cruz Biotechnology; 1:500) and anti-B-actin (Sigma-Aldrich; 1:2000) antibodies.

Brain hemispheres were homogenized in PBS (pH 7.4; 1:10, w/v) containing protease inhibitors (cOmplete<sup>TM</sup> Protease Inhibitor Cocktail), and the lysates were collected by centrifugation at 15 000 × g. The amount of protein in the lysate was measured using a bicinchoninic acid assay (Bio-Rad).

For Western blot analysis, proteins were prepared in sample buffer [62.5 mM Tris (pH 6.8), 2% SDS, 10% glycerol, 0.01% bromophenol blue, 5% mercaptoethanol and 50 mM dithiothreitol], and then separated by SDSpolyacrylamide gel electrophoresis and transferred onto polyvinylidene difluoride (PVDF) membranes (Merck Millipore) using semidry blotters (TE70 PWR; Amersham Biosciences, Uppsala, Sweden). To detect AB proteins, protein lysates in tricine sample buffer [200 mM Tris-HCl (pH 6.8), 40% glycerol, 2% SDS and 0.04% Coomassie Blue G-250; Bio-Rad] were electrophoresed on 10%-20% Mini-PROTEAN<sup>™</sup> Tris-Tricine Precast Gel (Bio-Rad) under nonreducing conditions. After blocking with 5% skim milk and 1% BSA in TBS-T buffer (25 mM Tris, 150 mM NaCl, 0.1% Tween 20; pH 7.4), the blot was reacted with a primary antibody and subsequently with a horseradish peroxidase-conjugated secondary antibody (Santa Cruz Biotechnology).

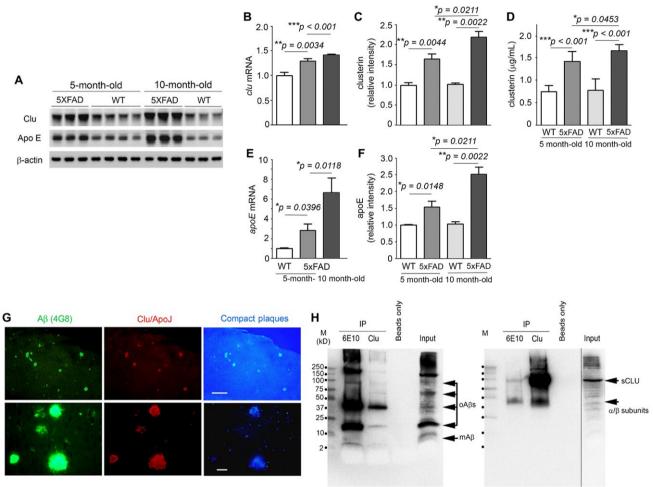
To determine interactive complexes of AB peptides and clusterin in the brain, we immunoprecipitated brain lysates (containing 800  $\mu$ g of total protein) with anti-AB(1–16) antibody (6E10) plus Protein G-sepharose beads (GE Healthcare, Buckinghamshire, UK) or anti-clusterin/Apo J antibody (Novus Biologicals) plus Protein G-sepharose beads. After washing in PBS, the protein/antibody-bound beads were eluted in tricine sample buffer [200 mM Tris-HCl (pH 6.8), 40% glycerol, 2% SDS and 0.04% Coomassie Blue G-250]. Shortly thereafter, the supernatant protein samples were collected by brief centrifugation and fractionated by electrophoresis on 10%–20% Tris-Tricine Precast Gel (Bio-Rad) under nonreducing conditions. Finally, the proteins were detected by Western blot using anti-AB antibody (6E10) or anti-clusterin/Apo J antibody.

Immunoreactive proteins were visualized using  $ECL^{\mathbb{M}}$ Prime Western Blotting Detection Reagent (GE Healthcare) with the Davinch-Western<sup>TM</sup> Chemiluminescence Imaging System (CAS-400SM; Davinch-K, Seoul, Korea). The band intensities were measured using ImageJ software (National Institutes of Health, Bethesda, MD, USA), and protein levels were normalized to those of  $\beta$ -actin.

#### Cognitive and motor behavioral analysis

Five- or 10-month-old mice were tested for spatial learning and memory in a Morris water maze (MWM) consisting of a circular plastic pool (120-cm in diameter) with a cylindrical escape platform (10-cm in diameter) in the northwest quadrant of the pool, 0.5-cm under the surface of opaque water (maintained at  $21.0^{\circ}$ C  $\pm 1.0^{\circ}$ C) and with three visual cues on the wall. For 5 consecutive days, the animals were given daily training to swim and find the hidden escape platform. Each day, they had three 60-s trials, each starting at a different quadrant (northeast, southeast and southwest quadrant). Three hours after the final training, the mice performed a probe trial for 60-s starting from the southeast quadrant with the escape platform removed. The time, path and distance spent swimming in the pool as well as the frequency the target zone was crossed trying to locate the platform were monitored with SMART Video Tracking System (Harvard Apparatus, Holliston, MA, USA).

For the following 3 days, the same mice were given a balance beam task to evaluate balance and motor coordination. The beam was a flat wooden bar (1.2-cm wide, 80-cm long) connected to two 40-cm high support columns. A dark escape chamber was located at the opposite end



W.B.: 6E10

W.B.: Clu

**Figure 1.** Interactions between clusterin and Aß in the brains of  $5\times$ FAD mice. **A**. Representative Western blots show expression or deposition of clusterin (Clu), Apo E and ß-actin (as a loading control) in the brains of 5- or 10-month-old  $5\times$ FAD mice and their wild-type littermates (WT). **B** and **E**. Clusterin (**B**) or Apo E (**E**) mRNA levels were measured by quantitative real time PCR. Values were calculated relative to that of GAPDH and bars are presented as fold changes relative to control values (5-month-old wild-type mice) (n = 6–10 for all groups). **C** and **F**. Graphs show densitometric quantification of clusterin (**C**) or Apo E (**F**) from representative blots (**A**). After normalization with ß-actin, the fold of change is presented as compared to control (5 month-old wildtype mice) (n = 3–4 for all groups). **D**. ELISA quantification of mouse clusterin from SDS extracts (n = 6–9 for all groups). Data represent mean ± SEM of three independent experiments. \**P* < 0.05, \*\**P* < 0.01, or \*\*\**P* < 0.001. **G**. Fluorescence microphotographs show

colocalization of Aß (green) and clusterin (red) in compact amyloid plaques (blue). After double immunofluorescence staining with anti-Aß(17–24) (4G8) and anti-clusterin (Clu)/Apo J antibodies, the brain tissues were examined at lower (×40, top row) and higher (×400, bottom row) magnifications under the appropriate fluorescence filters. Scale bars, 500  $\mu$ m (top row) and 50  $\mu$ m (bottom row). H. Brain lysates (800  $\mu$ g) were immunoprecipitated (IP) using either anti-Aß(1–16) (6E10) or anti-clusterin (Clu) antibody, and the precipitates were subjected to Western blot (W.B.) with anti-Aß(1–16) (6E10) (left panel) or anti-clusterin antibody (Clu; right panel). Beads only and input samples (50  $\mu$ g as total protein) show negative and positive signals, respectively. The numbers along the 1st lane (M, molecular marker) of each blot denote the molecular sizes (kD). mAß, Aß monomers; oAßs, Aß oligomers; sCLU, secretory clusterin;  $\alpha/\beta$  subunits,  $\alpha$  or  $\beta$  subunits of clusterin.

from where the mice were placed on the beam. The test was conducted in dim light with brighter illumination condensed on the starting point of the beam. The mice were encouraged to move from the starting point to the escape chamber on the other end. They were trained with three 90-s trials per day with a 15-min interval between trials. The time it took for the mice to traverse the beam and enter the chamber was recorded. If a mouse fell off the beam, we excluded that trial and conducted another trial. If a mouse stayed on the beam but did not reach the chamber within 90-s, the maximum time was recorded as 90-s.

#### Statistics

Data are presented as the means  $\pm$  standard errors of the mean. Statistical comparisons between the groups were performed with an unpaired *t*-test or one-way ANOVA with Student–Newman–Keuls post hoc test, using Prism (GraphPad Software, La Jolla, CA, USA). Differences were considered significant at P < 0.05.

#### RESULTS

# Interactive expression of Aß and clusterin in the brains of 5×FAD mice

We evaluated the expression levels of clusterin in the brains of 5×FAD mice and their wild-type littermates. In qRT-PCR or Western blot analysis (Figure 1A-C), we found that 5×FAD mice expressed higher levels of clusterin than their littermates at the same age (5 or 10 months of age) and had a significant increase in clusterin levels from 5 to 10 months of age. In the wild-type littermates, there was no significant difference in clusterin levels between 5 and 10 months of age. The increases in clusterin levels in the 5×FAD mice were also detectable by quantitative ELISA measurement of mouse clusterin in the brain lysates (Figure 1D). In addition, there were similar patterns of elevation of apolipoprotein E (Apo E) (Figure 1A, E and F) and DKK-1 (Supplementary Figure S1) in 5×FAD mice. These findings indicate that AB deposition leads to the increased expression of clusterin, Apo E or DKK-1 in the brain.

 $5 \times FAD$  mice not only produce abundant levels of Aß but also develop Aß deposits or amyloid plaques in their brain at a relatively young age (39). Aß-specific 4G8-immunohistochemistry detected a number of amyloid plaques throughout the brain tissues of 5-month-old  $5 \times FAD$ mice, primarily identified as compact core amyloid plaques by their *in situ* emission of blue fluorescence under ultraviolet (UV) illumination (26) (Figure 1G). In line with studies of human brains affected with AD (6,8,22), we found close colocalization of clusterin and Aß immunoreactivities in compact core amyloid plaques (Figure 1G).

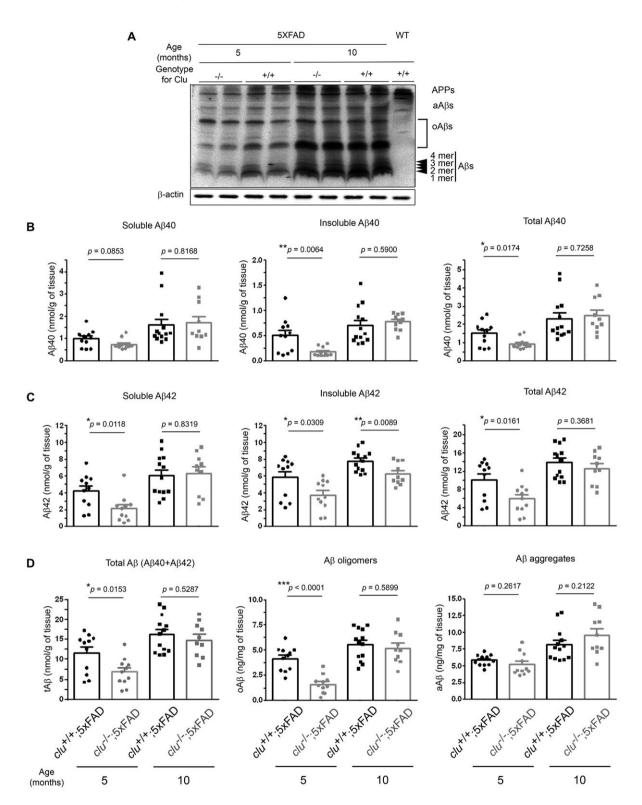
To explore *in vivo* interactions between clusterin and AB, we performed co-IP experiments with brain lysates of  $5 \times FAD$  mice (Figure 1H). When proteins

co-immunoprecipitated with anti-Aß antibody (6E10) were fractionated under nonreducing conditions and then immunoblotted with anti-clusterin antibody, we found two sizes of clusterin, 40-kD  $\alpha/\beta$  subunits and 80-kD mature secretory clusterin (sCLU). Additionally, reverse co-IP, which used anti-clusterin antibody and anti-Aß antibody (6E10) for IP and Western blot, respectively, detected multiple forms of Aß (monomers, dimers, oligomers and highmolecular-weight Aß aggregates). These results thus suggest that clusterin and Aß interact with each other *in vivo*, where clusterin interacts or binds with different Aß structures.

# Effects of genetic deletion of clusterin on amyloid pathogenesis

To investigate whether clusterin is implicated in amyloid pathogenesis, we compared the deposition of Aß in the brains of  $clu^{+/+}$ ;5×FAD and  $clu^{-/-}$ ;5×FAD mice at 5 or 10 months of age. Western blot analysis revealed that Aß was prominently upregulated with age in both clu genotypes of 5×FAD mice as their entire blot signals became more intense at 10 than at 5 months of age (Figure 2A). In contrast, the Aß signals were generally attenuated in  $clu^{-/-}$ ;5×FAD mice compared with those in  $clu^{+/+}$ ;5×FAD mice at 5 months of age, whereas the clu genotype-related differences in Aß intensities were undetectable at 10 months of age. Notably, significant differences among the groups were observed in band intensities of Aß monomers, dimers and oligomers (with molecular sizes ranging from approximately 4.0 kD to 50.0 kD).

Next, to further elaborate the differences in AB deposition, we measured the amounts of various AB peptides in the brains of  $clu^{+/+}$ ;5×FAD and  $clu^{-/-}$ ;5×FAD mice. We used ELISA to quantify the levels of soluble AB40 or AB42 and AB oligomers in PBS- and SDS-soluble fractions and insoluble AB40 or AB42 in FA-soluble fractions in serially fractionated brain lysates. Total Aβ40 or Aβ42 amounts were calculated as the sum of the soluble and insoluble fractions of AB40 or AB42. AB aggregates were also analyzed in SDS extracts collected by brief centrifugation at 10 000 × g but without ultracentrifugal fractionation. The levels of soluble, insoluble and total AB40 or AB42 were lower in 5-month-old  $clu^{-/-}$ ;5×FAD mice than in  $clu^{+/+}$ ;5×FAD mice (by 24.4%–63.8%; P = 0.0064– 0.0853) (Figure 2B and C), demonstrating that total AB (sum of AB40 and AB42) level was significantly lower in  $clu^{-/-}$ ;5×FAD mice (by 41.0%; P = 0.0153) (Figure 2D). Differences in AB42 levels were even more striking, with soluble AB42 levels in  $clu^{-/-}$ ;5×FAD mice 49.5% lower than those in  $clu^{+/+}$ ;5×FAD mice (P = 0.0118) and insoluble AB42 35.7% lower (P = 0.0309) (Figure 2C). Soluble AB oligomers were also markedly lower in  $clu^{-/-}$ ;5×FAD mice (by 60.9%; P < 0.0001), but Aß aggregates were only slightly lower (by 11.1%; P = 0.2617) (Figure 2D). From 5 months to 10 months of age, the levels of all pools of AB proteins (total, soluble and insoluble AB40 or AB42, AB oligomers and AB aggregates) were significantly or slightly increased



**Figure 2.** Deposition of various pools of brain Aß in clu<sup>+/+</sup>;5×FAD and clu<sup>-/-</sup>;5×FAD mice. **A**. Representative Western blot shows expression or deposition of multiple sizes of Aß proteins in the brains of *clu<sup>+/+</sup>*;5×FAD and *clu<sup>-/-</sup>*;5×FAD mice at 5 or 10 months of age. After immunoblotting with anti-Aß(1–16) antibody (6E10) (top panel), the blot was deprobed and reacted again with anti-ß-actin antibody as a loading control (bottom panel). Blot signals from a wild-type littermate mouse are shown on the right end of the blot (lane 9). oAßs, Aß oligomers; aAßs, Aß aggregates. B–D. ELISA quantification shows the deposition of various pools of Aß proteins. Soluble (1st column) and insoluble (2nd column) Aß40 (B) or Aß42 (C) were analyzed from PBS- and SDS-soluble fractions, and FA-soluble fractions of brain lysates, respectively.

in both *clu* genotypes of 5×FAD mice (Figure 2B–D), resulting in no differences in the levels of all pools of AB proteins between *clu<sup>-/-</sup>*;5×FAD and *clu<sup>+/+</sup>*;5×FAD mice at 10 months of age (Figure 2B–D), except that insoluble AB42 was 20.0% lower in *clu<sup>-/-</sup>*;5×FAD mice (P = 0.0089) (Figure 2C).

From these results for AB, we calculated the proportion of each pool or form of AB within the total amount of AB (sum of AB40 and AB42). First, we found neither clu genotype-related nor age-related differences in the ratio of total soluble or insoluble AB to total AB (Supplementary Figure S2A and S2B). However, compared with  $clu^{+/+}$ ;5×FAD mice,  $clu^{-/-}$ ;5×FAD mice at 5 months of age had a significantly lower ratio of soluble AB oligomers to total AB or to AB aggregates (by 34.6% or 52.9%; P = 0.0145 or 0.0002), whereas those ratios in both *clu* genotypes were not significantly different at 10 months of age (Supplementary Figure S2C and S2E). The proportion of Aß oligomers in the soluble fraction of total Aß (sum of AB40 and AB42) was also marginally lower in clu<sup>-/-</sup>;5×FAD mice at 5 months of age (by 27.3%; P = 0.0994) (Supplementary Figure S2D). However, the ratio of AB aggregates to total AB tended to be higher in  $clu^{-/-}$ ;5×FAD mice than that in  $clu^{+/+}$ ;5×FAD mice at each age (Supplementary Figure S2F). Further, when we compared the levels of AB40 and AB42, the ratio of total AB40 to total AB42 was somewhat higher in  $clu^{-/-}$ ;5×FAD mice than that in  $clu^{+/+}$ ;5×FAD mice at each age (by 32.7% or 23.7%; P = 0.1603 or 0.0695) (Supplementary Figure S2G), whereas the ratio of soluble AB40 to soluble AB42 was about twice higher in  $clu^{-/-}$ ;5×FAD mice at 5 months of age (by 91.0%, P = 0.0229), with the lack of *clu* genotyperelated difference at 10 months (by 5.1%, P = 0.7502) (Supplementary Figure S2H).

We also measured the loads of 4G8-immunoreactive or Congo red-positive amyloid deposits or plaques (Figure 3A). At 5 months of age, compared with  $clu^{+/+}$ ;5×FAD mice,  $clu^{-/-}$ ;5×FAD mice had a lower load of 4G8-immunoreactive amyloid deposits and fewer Congo red-positive amyloid plaques (by 40.4% and 43.1%; P = 0.0154 and 0.0004, respectively), which reflect the deposition of all forms of APP and AB, and the fraction of insoluble AB fibrils, respectively (Figure 3B). Thereafter, the loads substantially increased in the  $clu^{-/-}$ ;5×FAD mice so that by 10 months of age, there were no clu genotype-related differences Total levels of Aß40 (**B**, 3rd column), Aß42 (**C**, 3rd column) and Aß (**D**, 1st column) were calculated as the sum of soluble and insoluble Aß40, Aß42 and total Aß40 and total Aß42, respectively. Soluble Aß oligomers (oAß) were quantified from PBS- and SDS-soluble fractions (**D**, 2nd column), whereas Aß aggregates (aAß) were analyzed from SDS extracts without ultracentrifugal fractionation (**D**, 3rd column). All ELISA analyses were performed in duplicate. Numbers (males and females) of animals included in each group were 11 (5 and 6; for 5-month-old *clu<sup>+/+</sup>;*5×FAD), 11 (5 and 6; for 5-month-old *clu<sup>-/-</sup>;*5×FAD), 13 (6 and 7; for 10-month-old *clu<sup>+/+</sup>;*5×FAD) or 10 (5 and 5; for 10-month-old *clu<sup>-/-</sup>;*5×FAD mice). \**P* < 0.05, \*\**P* < 0.01, or \*\*\**P* < 0.001.

(Figure 3B). Finally, the ratios of Congo red-positive amyloid plaques to 4G8-immunoreactive amyloid deposits did not significantly differ between the two *clu* genotypes at 5 or 10 months of age. Instead, the ratios were significantly decreased with age in both *clu* genotypes (P = 0.0015 for *clu*<sup>-/-</sup>;5×FAD mice or P = 0.0004 for *clu*<sup>+/+</sup>;5×FAD mice) (Supplementary Figure S2I).

Considered together, these quantitative results indicate that clusterin expression is correlated with amyloid deposition at the early age in 5×FAD mice, but the effects decline or disappear with age.

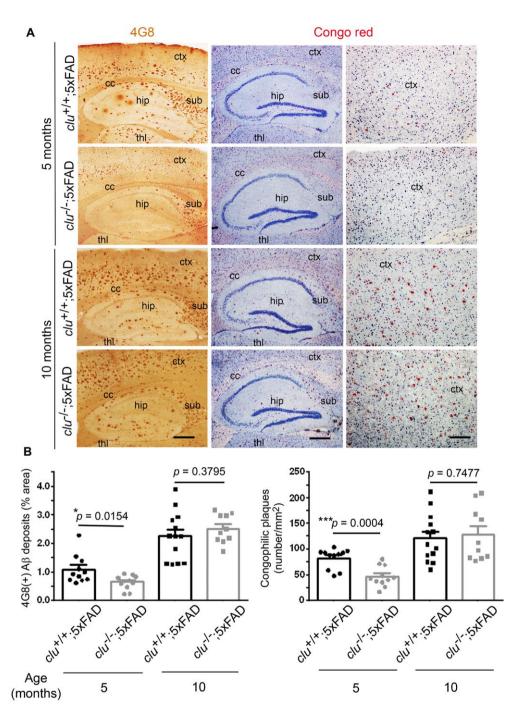
## Effects of clusterin deficiency on Aß-induced neuronal or synaptic loss

The neuropathology of AD includes loss of neurons and synapses as well as AB deposition. Thus, to determine the involvement of clusterin in AB-induced neuronal or synaptic degeneration, we compared the expression of neuronal or synaptic marker proteins using Western blot analysis (Figure 4). The proteins with striking changes in expression included neuronal specific NeuN, nonphosphorylated neurofilament protein SMI-32 (52), presynaptic synaptophysin, axon initial segment-specific and voltagegated sodium channels-associated ankyrin G (47), postsynaptic PSD-95, vesicle-associated membrane protein (VAMP; synaptobrevin) and postsynaptic scaffold protein ProSAP2/Shank 3 (15). The levels of the neuron- and synapse-specific proteins were significantly lower in 5-month-old  $clu^{+/+}$ ;5×FAD mice than those in nontransgenic wild-type mice, but were significantly increased in the age-matched clu<sup>-/-</sup>;5×FAD mice. However, the protein expressions in clu<sup>-/-</sup>;5×FAD mice were decreased by 10 months of age to the levels as those in  $clu^{+/+}$ ;5×FAD mice.

#### Cognitive and motor performance in clusterinnull 5×FAD mice

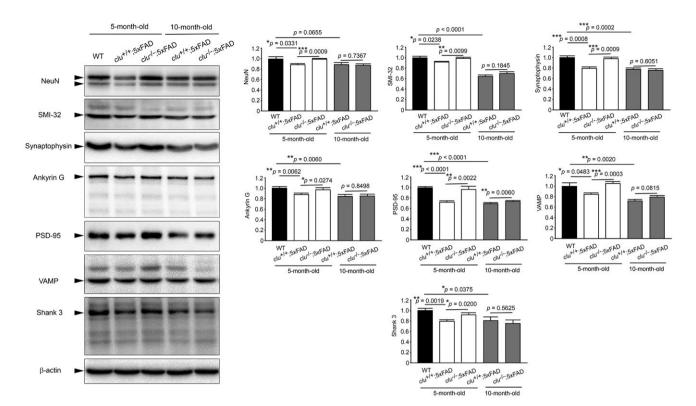
Finally, to determine whether clusterin is involved in cognitive and behavioral impairment during the development of AD, we evaluated the effect of genetic deletion of clusterin on cognitive and motor performance in  $5 \times FAD$  mice.

To evaluate the effects of clusterin on spatial learning and memory performance, the mice were trained in a Morris water maze (MWM) with 3 trials every day for



**Figure 3.** Quantification of amyloid deposits or plaques by immunohistochemical or Congo red staining. **A.** Sagittal brain sections of 5- or 10-month-old  $clu^{+/+}$ ;5×FAD and  $clu^{-/-}$ ;5×FAD mice were stained with anti-A $\beta$ (17–24) antibody (4G8) or Congo red dye to detect pan-amyloid precursor protein (APP)/amyloid- $\beta$  (A $\beta$ )-immunoreactive deposits (1st column, brown dots), or compact amyloid plaques (2nd and 3rd column, red dots). hip, hippocampus; ctx, cortex; thl, thalamus; cc, corpus callosum; sub, subiculum. Scale bars, 500 µm (1st and 2nd column) or 200 µm (3rd column). **B.** The

per cent area of 4G8-immunoreactive amyloid deposits (left graph) or the total number of congophilic amyloid plaques (right graph) in the same brain region of interest was measured in five brain sections per animal, taken every 200 µm lateral from the midline. Numbers (males and females) of animals in each group were 11 (5 and 6; for 5-month-old *clu<sup>+/+</sup>*;5×FAD), 11 (5 and 6; for 5-month-old *clu<sup>-/-</sup>*;5×FAD), 13 (6 and 7; for 10-month-old *clu<sup>+/+</sup>*;5×FAD), or 10 (5 and 5; for 10-month-old *clu<sup>-/-</sup>*;5×FAD mice). \**P* < 0.05, \*\**P* < 0.01, or \*\*\**P* < 0.001. at wileyonlinelibrary.com]



**Figure 4.** Effect of clusterin and age on expression of neuronal or synaptic proteins in the brain of 5×FAD mouse. **A.** Representative Western blots show expression of neuronal or synaptic proteins in 5- and 10-month-old *clu++;*5×FAD and *clu-+/-;*5×FAD mice and 5-month-old nontransgenic wild-type (WT) mice, representing only results with significant differences in expression; neuronal specific NeuN (molecular size, 46-48 kD), nonphosphorylated neurofilament SMI-32 (200 kD), presynaptic synaptophysin (38 kD), axon initial segment-specific and voltage-gated sodium channels-associated ankyrin G (190 kD), postsynaptic PSD-95 (95 kD), vesicle-associated membrane protein

5 days (Figure 5A). Unexpectedly, both 5-month-old  $clu^{-/-}$ ;5×FAD mice and nontransgenic wild-type mice found the platform significantly more quickly than the clu+/+;5×FAD mice of same age on the first day of training (P = 0.0021 and 0.0183, respectively). Thereafter, the 5-day repeated trainings led to significant improvement in locating the hidden platform for every group of 5-monthold mice, although the nontransgenic wild-type mice and  $clu^{-/-}$ ;5×FAD mice were consistently and significantly faster than  $clu^{+/+}$ ;5×FAD mice. On the last trial day,  $clu^{-/-}$ ;5×FAD mice were able to locate the platform more rapidly than  $clu^{+/+}$ ;5×FAD mice (P = 0.0004), indicating that clusterin deficiency promoted better spatial learning performance in 5-month-old 5×FAD mice. Three hours later, the animals underwent a free swim probe test to evaluate recall for the previous location of the target platform, a measure of spatial reference memory retention in trained animals (Figure 5B-F). Comparing the speed or the mean distance in reaching the target zone where the escape platform had been located, the number of times the target zone was crossed, and the time spent in the target quadrant,

(VAMP, synaptobrevin; 16 kD), postsynaptic scaffold protein ProSAP2/ Shank3 (180 kD) and ß-actin (42 kD). **B–H**. Densitometric quantification of indicated proteins after normalization to ß-actin. Graphs show significant differences in the levels of the indicated proteins (arrows in blots) between nontransgenic wild-type and 5×FAD mice and between 5-month-old  $clu^{+/+}$ ;5×FAD and  $clu^{-/-}$ ;5×FAD mice. Protein levels are shown as values relative to a control (black bars, wild-type mice). Bars denote mean ± SEM of three independent experiments. Each group includes 6 mice (3 males and 3 females). \*P < 0.05, \*\*P < 0.01, or \*\*\*P < 0.001.

we found that 5-month-old  $clu^{+/+}$ ;5×FAD mice had impaired performance relative to their nontransgenic wild-type littermates. Age-matched  $clu^{-/-}$ ;5×FAD mice had significantly better spatial memory performance than the  $clu^{+/+}$ ;5×FAD mice. However, this good level of performance did not persist till 10 months of age in the mice with genetic deletion of clusterin. There were no differences between 10-month-old  $clu^{-/-}$ ;5×FAD and  $clu^{+/+}$ ;5×FAD mice in performance of MWM and probe trials (Figure 5A–F).

After completion of the MWM test, the animals underwent 3 days of training on a balance beam task (Figure 6). On the first trial day, nontransgenic wild-type littermates more rapidly reached the other end, followed by  $clu^{-/-}$ ;5×FAD mice, with the  $clu^{+/+}$ ;5×FAD mice being the slowest; these differences, however, were not statistically significant. However, by the third day, the 5-month-old wild-type mice and  $clu^{-/-}$ ;5×FAD mice moved into the escape chamber increasingly quickly (P = 0.0363 and 0.0036, respectively), spending a significantly shorter time on the beam compared with the age-matched  $clu^{+/+}$ ;5×FAD mice (P = 0.0011 and 0.0008, respectively). In contrast, at 10

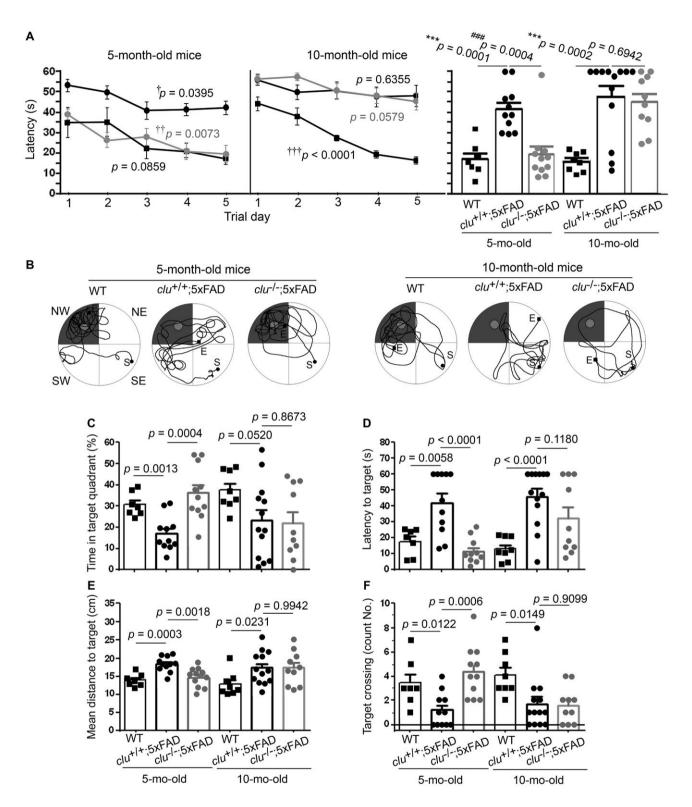
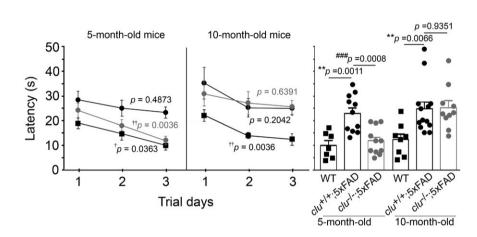


Figure 5. Effect of clusterin on performance of spatial learning and memory in 5×FAD mice. A. Animals were trained daily to find a hidden platform in a Morris water maze (MWM). The time taken to find the platform was significantly longer over 5 days in *clu*<sup>+/+</sup>;5×FAD mice (black circles) compared with their nontransgenic wild-type littermates (rectangles). The *clu-/-*;5×FAD mice (gray circles) were significantly faster than *clu*<sup>+/+</sup>;5×FAD mice at 5 months of age (left graph), but at 10 months of age, they performed no better than clu+/+;5×FAD mice (right graph). Crosses (†) denote the effect of 5 days-trainings on the time to reach the target platform in each group (by one-way ANOVA with Student-Newman-Keuls post hoc test).  $^{+}P < 0.05$ ,  $^{++}P < 0.01$ , or  $^{+++}P < 0.001$ . Vertical bars on the right panel denote the values of latency time taken by each animal group on the last trial day. B-F. After the training, 60 s-probe trials were performed to evaluate the entire path taken (B; from points "S" to "E") and the percentage of time spent in the target zone (northwest shadow

quadrant in B) searching for the platform that had been removed (small circular area in target guadrant in B) (C) as well as time (D) and mean distance (E) taken to get the first touch to the target location, and the frequency at which the target location was crossed (F). The clu+/+;5×FAD mice performed poorly on all evaluations compared with nontransgenic wild-type mice. The *clu-/-*;5×FAD mice at 5 months of age performed as well as the wild-type littermates at 5 months of age, but at 10 months of age, their performance was as poor as the *clu*<sup>+/+</sup>;5×FAD mice. Numbers (males and females) of animals in each group were 11 (5 and 6; for 5-month-old *clu<sup>+/+</sup>*;5×FAD), 11 (5 and 6; for 5-month-old *clu*<sup>-/-</sup>;5×FAD), 13 (6 and 7; for 10-monthold clu+/+;5×FAD), or 10 (5 and 5; for 10-month-old clu-/-;5×FAD mice). Asterisk (\*) and pound (#) denote the comparisons between clu+/+;5×FAD and nontransgenic wild-type mice and between clu+/+;5×FAD and clu-/-;5×FAD mice, respectively (by an unpaired ttest).  $^{*,\#}P < 0.05$ ,  $^{**,\#\#}P < 0.01$ , or  $^{***,\#\#}P < 0.001$ .



**Figure 6.** Effect of clusterin on balance and motor coordination in  $5\times$ FAD mice. At 5 months of age,  $clu^{-/-};5\times$ FAD mice (gray circles) and wild-type mice (rectangles) reached the escape chamber at the opposite end of the beam significantly faster over 3 days of trials than  $clu^{+/+};5\times$ FAD mice (black circles) (left graph). However, by 10 months of age, the  $clu^{-/-};5\times$ FAD mice took significantly longer than before and were now as slow as  $clu^{+/+};5\times$ FAD mice (right graph), both genotypes of  $5\times$ FAD mice being significantly slower than 10-month-old wild-type mice. Vertical bars on the right panel denote the values of latency time taken by each animal group on the last trial day. Numbers (males and

months of age,  $clu^{-/-}$ ;5×FAD and  $clu^{+/+}$ ;5×FAD mice both failed to acquire the capability to run along the beam with the balance and motor coordination achieved by nontransgenic wild-type mice (P = 0.6391, 0.2042 and 0.0036, respectively). Thus, there were no differences between the two *clu* genotypes of 5×FAD mice on the third day of balance beam task (P = 0.9351) (Figure 6).

#### DISCUSSION

Since clusterin can maintain Aß solubility by preventing Aß aggregation (32) and promote its clearance from brain by mediating BBB transport (3,57), early interest in clusterin was centered on defining its protective role against AD. Experimental support for this possibility came from

females) of animals in each group were 11 (5 and 6; for 5-month-old  $clu^{+/+};5\times$ FAD), 11 (5 and 6; for 5-month-old  $clu^{-/-};5\times$ FAD), 13 (6 and 7; for 10-month-old  $clu^{+/+};5\times$ FAD), or 10 (5 and 5; for 10-month-old  $clu^{-/-};5\times$ FAD mice). Asterisk (\*) and pound (#) signs denote comparisons between  $clu^{+/+};5\times$ FAD and wild-type mice and between  $clu^{+/+};5\times$ FAD and  $clu^{-/-};5\times$ FAD mice, respectively (by an unpaired *t*-test). \*,#*P* < 0.05, \*\*,##*P* < 0.01, or \*\*\*,###*P* < 0.001. Crosses (†) denote the effect of 3 days of training on the time to reach the escape chamber in each group (by one-way ANOVA with Student–Newman–Keuls post hoc test). \**P* < 0.05 or \*†*P* < 0.01.

findings that clusterin alleviated AB toxicity in neuronal cells (4,29,56). However, growing evidence suggests that clusterin interacts with AB to enhance its neurotoxicity and cause neurodegeneration (23,25,40,41). A previous study with hAPP transgenic PDAPP mice (10) provided the first evidence of an *in vivo* functional role for clusterin, with its absence leading to fewer amyloid plaques and lesser neuritic dystrophy despite a consistent level in total AB. Similarly with our present study of 5×FAD mice, a recent study (55) also found a marked decrease in total amyloid deposition containing soluble and insoluble AB as well as parenchymal amyloid plaques in clusterin-null APP/PS1 mice. Although the different findings concerning the deposition of total AB likely came from the different background features of hAPP transgenic mice used (55), it is important

that clusterin deficiency prevented amyloid deposition and  $A\beta$ -induced neurodegeneration in the hAPP transgenic mouse models of AD.

In the present study, we observed interactions between clusterin and AB proteins in the brains of 5×FAD mice. Both mRNA and protein levels of clusterin were higher in 5×FAD mice than in age-matched wild-type littermates, a finding similar to that in the brains of human patients with AD compared with controls without dementia (14,33,40). While clusterin levels did not significantly change with age in wild-type mice, they substantially increased with age in 5×FAD mice. In addition, the expression of DKK-1, a Wnt antagonist mediating clusterin-driven Aß toxicity (23,45), was also significantly higher and increased with age in 5×FAD mice. These findings indicate that AB deposition causes increased expressions of clusterin and DKK-1 in the hAPP-transgenic mouse models (46). Next, we detected colocalization of AB and clusterin in nearly all compact amyloid plaques in 5×FAD mice, which is similar to that found in the brains of AD patients (8,22,34). Furthermore, we found that AB and clusterin were in the same complex as they were co-immunoprecipitated with each other in brain lysates of 5×FAD mice. Because clusterin was detected together with AB proteins of various sizes in our co-IP experiment, it is thought that clusterin has the propensity to bind or interact with different forms of AB, such as monomers, oligomers, or fibril aggregates in vivo. Using purified or synthetic proteins, many studies have shown that clusterin binds to  $A\beta$  proteins with high affinity (2.13.31.37).

This study provides the first *in vivo* evidence that clusterin-directed Aß deposition and oligomerization augment Aß neurotoxicity and underlie AD pathogenesis in hAPP transgenic mice. Although knockdown of clusterin was previously shown to reduce A $\beta$  toxicity in primary neurons (23), we further elucidated how genetic loss of clusterin influences AD pathology in 5×FAD mice. Clusterin deficiency led to less Aß (total, soluble and insoluble Aß40 and Aß42) and fewer amyloid plaques in the brains of 5×FAD mice at 5 months of age. Clusterin binding could render Aß resistant to proteolytic degradation and maintain its stability (32,37); therefore,  $clu^{+/+}$ ;5×FAD mice would exhibit higher amyloid deposition than  $clu^{-/-}$ ;5×FAD mice. Alternatively, the reduced level of Aß likely is due to the decrease in Aß generation in  $clu^{-/-}$ ;5×FAD mice.

However, we specifically paid attention to AB oligomers and AB aggregates because AB neurotoxicity is primarily attributed to its oligomer forms rather than its monomer or fibril forms (25,40,41). AB42 is considered to be more neurotoxic than AB40 because of its propensity to more readily aggregate into oligomers (16,30,48,53). Clusterin, then, could sequester or stabilize toxic AB oligomers so that they are less degradable, or inhibit further aggregation of AB oligomers into nontoxic fibrils (32,37). When compared the results of ELISA quantification in  $clu^{-/-}$ ;5×FAD mice with those in  $clu^{+/+}$ ;5×FAD mice, level of soluble AB42, in which include AB42 oligomers, was more reduced than that of soluble AB40 or insoluble AB42, with the most dramatic reduction in soluble AB oligomers. The

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difference in AB aggregates composed of AB oligomers, protofibrils and fibrils, was not statistically significant. Next, we found no *clu* genotype-related difference in the ratio of soluble  $A\beta$ /total  $A\beta$  or insoluble  $A\beta$ /total  $A\beta$ , whereas the ratio of soluble AB oligomers/total AB, soluble AB oligomers/total soluble AB, or soluble AB oligomers/AB aggregates was markedly lower in *clu*<sup>-/-</sup>;5×FAD mice. The ratio of AB aggregates/total AB became rather somewhat higher. We further found that lack of clusterin caused lower loads of 4G8-immunoreactive plaques and fewer Congo red-positive fibrillar amyloid plaques, which are representative of total AB deposition and insoluble AB aggregates, respectively. The ratios of Congo red-positive plaques/4G8immunoreactive plaques were similar in both *clu* genotypes of 5×FAD mice. Finally, 5-month-old clu-/-;5×FAD mice had less neuronal and synaptic loss as evaluated by the expression of neuronal or synaptic proteins. These mice also had significantly better motor and cognitive performances. Collectively, these results suggest that clusterin is involved in amyloid pathogenesis responsible for neurodegeneration and behavioral impairment seen in the 5×FAD mouse model of early AD, perhaps by facilitating AB production and deposition or by shifting AB composition toward more toxic soluble AB oligomers.

However, it should be noted that any influence of clusterin deficiency on AD pathology declined or disappeared with age. By 10 months of age, all AB pools in  $clu^{-/-}$ ;5×FAD mice had significantly increased to levels similar to those in  $clu^{+/+}$ ;5×FAD mice. There were no detectable differences in amyloid plaque deposition, neuronal and synaptic loss, or motor and cognitive impairment between the two clu genotypes of 5×FAD mice. These findings reminded us of the previous human brain studies that demonstrated an association of clusterin with AD pathology in early stage. In longitudinal cohort analyses combining volumetric MRI scans with quantification of clusterin and Aß proteins, plasma clusterin levels were associated with atrophy in brain regions affected in AD (21,50). More importantly, an interaction between Aβ42 and clusterin has been exclusively related with atrophy of the entorhinal cortex, a region that is selectively affected early in the pathogenesis of AD (5,51). In cognitively normal older individuals and those with mild cognitive impairment (MCI), decreased levels of CSF Aβ42 (related to increased brain A642 deposition) and elevated CSF clusterin levels were associated with higher entorhinal cortex volume loss but not with hippocampal atrophy (11). The A\beta-clusterin association with entorhinal cortical atrophy was stronger in those individuals than in patients with advanced AD. Together with these human cohort studies, our findings that clusterin deficiency was associated with a significantly lower degree of amyloid pathology, neuronal and synaptic loss and behavioral and cognitive impairment in 5- monthold mice but not in 10- month-old mice suggest that clusterin may facilitate amyloid pathogenesis and induce Aβ-directed neurodegeneration in the early stages of AD (11, 28, 50).

Currently, it is unclear how clusterin can have disparate effects on the early and late stages of AD pathogenesis.

One possible explanation is that the effect of clusterin on Aß neurotoxicity may depend upon a substoichiometric ratio of clusterin and AB (54). Earlier in vitro studies showed that clusterin modulates amyloid pathogenesis in a biphasic manner to promote or inhibit its neurotoxicity. A higher proportion of clusterin in an AB-clusterin complex could cause relatively less AB aggregation and toxicity, whereas a lower proportion could facilitate AB aggregation and increase toxicity (4,19,41,56). Our present study supports that hypothesis because we observed that, with age, levels of AB increased to a greater extent than those of clusterin. In the brains of 10-month-old 5×FAD mice, total AB (sum of AB40 and AB42) and AB42 were elevated by 39.6% and 37.4%, respectively, relative to values in 5-monthold 5×FAD mice. While clusterin levels also increased from 5 to 10 months of age, the 17.3% increase was relatively smaller than that of AB proteins. The resulting decline in the ratio of clusterin to AB may affect amyloid pathogenesis, thereby aggravating neurodegeneration and behavioral or cognitive impairment (36). A further possibility is that as AD progresses into the later stages with a rapid increase in AB deposition, the clusterin content might become insufficient to mediate amyloid pathogenesis and other more amyloidogenic factors such as phosphorylated tau (p-tau) or Apo E may become more prominent. Corroborating this hypothesis, we noticed that, by 10 months of age, clu-/-;5×FAD mice had levels of amyloid pathology, neuronal and synaptic loss and motor and cognitive impairment comparable to those of  $clu^{+/+}$ :5×FAD mice. A prior study in Apo E and clusterin double knockout PDAPP mice (9) showed that the concurrent absence of Apo E and clusterin caused earlier onset and larger amyloid deposition, suggesting that clusterin can modify the structure, accumulation, deposition and toxicity of AB in cooperation with Apo E. In addition, a human brain study (11) found that interaction only between clusterin and  $A\beta$ was associated with volume loss in the entorhinal cortex in cognitively normal elderly people, whereas in those with MCI, atrophy was also related to the interaction of p-tau and AB. Moreover, they showed that interaction between A $\beta$  and p-tau was significantly associated with later atrophy in the hippocampus or other temporal lobe regions, which was not the case with the interaction of clusterin and  $A\beta$ . Therefore, these findings support the contention that other factors such as Apo E or p-tau have a stronger influence than clusterin on AD pathology in the later stages.

A recent study is of note because it investigated the effect of clusterin deficiency on amyloid pathology in APP/ PS1 transgenic mice (55), describing the role of clusterin in A $\beta$  transport across the BBB via low-density lipoprotein receptor-related protein-2 (LRP2) (3). They found a marked decrease in amyloid deposition in the brain parenchyma but a parallel increase in cerebral amyloid angiopathy (CAA) in *clu<sup>-/-</sup>*;APP/PS1 mice. As CAA is likely due to amyloid deposition on the cerebral vessel walls along the perivascular drainage pathways (1), it could be indicative of an abnormal A $\beta$  clearance in AD brain. Mounting evidence suggests that A $\beta$ 40 has a stronger propensity to promote CAA relative to A $\beta$ 42, and thus the higher A $\beta$ 40/ Aβ42 ratio favors toward CAA development (12,18). When we briefly compared the levels of Aβ40 and Aβ42 in our 5×FAD mice, the Aβ40/Aβ42 ratio was significantly increased in the soluble fractions of 5 months-old  $clu^{-/-}$ ;5×FAD mice, supporting the previous findings (49,55). However, there was a statistically nonsignificant increase in the ratio of total Aβ40/Aβ42, and the ratio differences disappeared at 10 months. While these differences in Aβ40/ Aβ42 ratio might represent the effect of clusterin deficiency on CAA in 5×FAD mice, it would be warranted to explore the expression of CAA in clusterin-wildtype or -null 5×FAD mice in order to define the roles of clusterin in Aβ clearance or transport *in vivo*.

In conclusion, clusterin preferentially augments the toxic A $\beta$  pools to facilitate the early stage of AD pathogenesis, and thereby promotes A $\beta$ -directed neurodegeneration and behavioral and cognitive impairments. However, as the pathology progresses, its influences decline, suggesting that substances other than clusterin may be more critical in the later stages. Therefore, a strategy to reduce the level of clusterin or to inhibit clusterin-A $\beta$  interactions may provide potential therapeutic opportunities for AD in early stage, but it should be considered whether such a treatment can consistently retard the disease pathogenesis in later advanced stages.

### ACKNOWLEDGMENTS

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### DECLARATIONS

The authors declare no conflict of interest.

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### SUPPORTING INFORMATION

Additional supporting information may be found in the online version of this article at the publisher's web site:

**Figure S1.** Expressions of mRNA and protein of DKK-1 in the brains of 5-month-old wild-type (WT), and 5- or 10-month-old 5×FAD mice.

**Figure S2.** Relative proportions of various pools of AB proteins in the brains of  $clu^{+/+}$ ;5×FAD and  $clu^{-/-}$ ;5×FAD mice.