Aβ Secretion and Plaque Formation Depend on Autophagy

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SUMMARY

Alzheimer's disease (AD) is a neurodegenerative disease biochemically characterized by aberrant protein aggregation, including amyloid beta (A β) peptide accumulation. Protein aggregates in the cell are cleared by autophagy, a mechanism impaired in AD. To investigate the role of autophagy in A β pathology in vivo, we crossed amyloid precursor protein (APP) transgenic mice with mice lacking autophagy in excitatory forebrain neurons obtained by conditional knockout of autophagy-related protein 7. Remarkably, autophagy deficiency drastically reduced extracellular Aβ plaque burden. This reduction of A β plaque load was due to inhibition of A β secretion, which led to aberrant intraneuronal $A\beta$ accumulation in the perinuclear region. Moreover, autophagy-deficiency-induced neurodegeneration was exacerbated by amyloidosis, which together severely impaired memory. Our results establish a function for autophagy in Aβ metabolism: autophagy influences secretion of A β to the extracellular space and thereby directly affects $A\beta$ plaque formation, a pathological hallmark of AD.

INTRODUCTION

Alzheimer's disease (AD) is the major form of dementia in the elderly, characterized by memory loss and cognitive decline. AD brain pathology includes intracellular aggregation of amyloid beta (A β) peptide and protein tau and extracellular A β plaques (Wirths and Bayer, 2012). A β is generated by sequential cleavage of type I transmembrane amyloid precursor protein (APP) by β - and γ -secretases (LaFerla et al., 2007). Mutations in *APP* or the γ -secretase subunits presenilin 1 (*PS1*) and *PS2* cause early-onset familial AD (FAD), thereby tightly linking AD and A β . However, FAD cases represent a few percent of all AD cases,

and therefore other cellular mechanisms that affect A β metabolism likely contribute to the pathology in sporadic AD.

In the AD brain, autophagosomes accumulate in the dystrophic neurites indicating impaired autophagy (Nixon, 2007). Macroautophagy (herein referred to as autophagy; reviewed in Mizushima and Komatsu, 2011 and Harris and Rubinsztein, 2012) controls cellular proteostasis by sequestering and delivering protein aggregates and cellular organelles to lysosomes for degradation. In AD, both the autophagy-inhibitory mammalian target-of-rapamycin (mTOR) signaling and the levels of lysosomal hydrolases are increased (Yang et al., 2011), the latter of which may reflect impaired autophagosomal-lysosomal clearance (Boland et al., 2008). Furthermore, FAD-associated mutations in PS1 disrupt lysosomal proteolysis (Lee et al., 2010a), whereas genetic deletion of the endogenous lysosomal-associated cathepsin inhibitor cystatin B restores lysosomal clearance in autophagy-deficient TgCRND8 mice (Yang et al., 2011). In addition, autophagy sustains axonal homeostasis of neurons (Komatsu et al., 2007) and its absence causes neurodegeneration (Hara et al., 2006; Inoue et al., 2012; Komatsu et al., 2006). A role for autophagy in A β metabolism has been suggested (Boland et al., 2010; Caccamo et al., 2010; Jaeger et al., 2010). For example, autophagosomes generate and contain Aß (Yu et al., 2005) and oxidative stress-induced autophagy increases Aβ generation (Zheng et al., 2011). Moreover, induction of autophagy by rapamycin in vivo lowers intracellular $A\beta$ levels and improves cognition (Caccamo et al., 2010), and long-term rapamycin treatment reduces plaque load in AD model mice (Majumder et al., 2011). Conversely, heterozygous deletion of autophagy-initiating Beclin1, which is decreased in early AD, increases both intracellular and extracellular Aß load (Pickford et al., 2008). Endocytosis of exogenous Aß in turn inhibits autophagy by increasing mTOR signaling (Caccamo et al., 2010).

Although previous studies have linked A β metabolism to the degradative function of autophagy, the ultimate effect of genetic deletion of autophagy on A β metabolism remained to be elucidated. We therefore generated forebrain excitatory neuron-specific autophagy-deficient APP transgenic mice. This was achieved by conditional knockout of autophagy-related gene 7 (*Atg7*). Unexpectedly, autophagy deficiency drastically reduced







the extracellular A β plaque load. Through our detailed examination, we have uncovered that the reduced A β burden was caused by impaired secretion of A β . These data reveal an additional role of autophagy in A β metabolism and highlight the importance of autophagy in AD.

RESULTS

Aβ Plaque Formation Is Dependent on Autophagy

To investigate the role of autophagy in A β pathology of AD, we generated neuron-specific autophagy-deficient mice by conditionally knocking out the autophagy-essential enzyme Atg7 in excitatory neurons in the mouse forebrain. This was achieved by crossbreeding $Atg7^{flox/flox}$ mice (Komatsu et al., 2005) with calcium/calmodulin-dependent protein kinase II (CaMKII)-*Cre* transgenic (Tg) mice. In agreement with previous studies (Komatsu et al., 2006; Inoue et al., 2012) autophagy deficiency led to accumulation of p62/p-S403-p62-positive and polyubiquitinated proteins that formed inclusion bodies in cornu ammonis 1 (CA1), accompanied by accumulation of quality control autophagy-associated histone deacetylase 6 (HDAC6) in CA3 and triggered astrocytosis (Figures S1, S2, and S3).

Next, $Atg7^{flox/flox}$; CamKII-*Cre* mice were crossbred with APP23 Tg AD model mouse (herein referred to as APP), and at 20 months of age, A β plaque formation was investigated in $Atg7^{flox/flox}$ × APP and $Atg7^{flox/flox}$; CamKII-*Cre* × APP littermates by A β immunostaining. As expected, $Atg7^{flox/flox}$ × APP mice exhibited heavy plaque burden at this age. In sharp contrast, A β plaque load was drastically reduced upon genetic deletion of Atg7 (Figure 1A; p < 0.005). Consistently, the levels of Tris-soluble (TS) and guanidine-HCI-soluble (GS) A β 40 and A β 42 were substantially lowered in $Atg7^{flox/flox}$; CamKII-*Cre* × APP mice (Figure 1B; p < 0.005). These findings imply that autophagy plays a critical role for A β plaque formation.

Autophagy Deficiency Leads to Intracellular A β Accumulation

The decreased A β plaque load suggested that either A β generation is decreased or that A β accumulates intracellularly in the autophagy-deficient mice. To elucidate the underlying cause, brain sections of 6-month-old *Atg7^{flox/flox}* × APP and *Atg7^{flox/flox}*; CamKII-*Cre* × APP littermates were immunostained with A β specific antibodies, a time point well before A β plaque formation starts (Figure S2E; Figure 1C). Interestingly, autophagy deficiency induced intracellular A β accumulation in CA1 and cortical pyramidal neurons (p < 0.05). Consistently, the levels of both GS-A β 40 and GS-A β 42 were slightly but significantly increased in

Atg7^{flox/flox}; CamKII-Cre mice compared to Atg7^{flox/flox} mice and in Atg7^{flox/flox}; CamKII-Cre × APP mice compared to Atg7^{flox/flox} × APP mice (Figure 1D; p < 0.05). However, the measured A β concentrations were normalized to the wet weight of the brain tissue, and, as described below, autophagy deficiency induces neurodegeneration, which reduces the brain weight by approximately 10% at 6 months of age (data not shown). Therefore, the increased GS-A β concentrations most likely reflect both increased intracellular A β levels (approximately 10%) as well as reduced brain weight. In addition, the levels of APP, APP C-terminal fragment, PS1, and β -secretase 1 were not altered in autophagy-deficient mice (Figure 1E). In conclusion, autophagy deficiency induces intracellular A β accumulation.

Autophagy Influences Secretion of A_β

The lowered extracellular A β plaque burden and the increased intracellular A β accumulation indicated that autophagy deficiency impairs secretion of A β to extracellular space either through impaired exocytic or excretory mechanisms. Indeed, genetic inhibition of autophagy in cortical/hippocampal primary neurons reduced the extracellular release of endogenous A β by 90% (Figure 2A; p < 0.0001). Supplementing the autophagy-deficient neurons with Atg7, expressed from a lentivirus to a level similar to that of endogenous Atg7 in autophagy-competent neurons, reactivated autophagy and restored the level of released A β to extracellular space to that of autophagy-competent neurons (Figures 2B and 2C; p < 0.01). In addition, lenti-Atg7 expression in autophagy-competent neurons increased autophagy and enhanced extracellular A β release.

In parallel, wild-type primary neurons were treated with pharmacological activators and inhibitors of autophagy. Low nanomolar concentrations of the mTOR inhibitor rapamycin increased the amount of autophagosomes as measured by LC3 metabolism and p70 phosphorylation and induced Aß secretion (Figure 2D; p < 0.01). In contrast, inhibition of autophagy by spautin-1 significantly reduced extracellular A β release (p < 0.05), as did inhibition of transport by exposing the neurons to the microtubule destabilizing agent vinblastine (p < 0.01). The data obtained by genetically and pharmacologically manipulating autophagy suggest a role for autophagy in intracellular transport and secretion of A β . In agreement, A β immunostaining of autophagy-deficient neurons revealed substantial accumulation of $A\beta$ in the perinuclear region, whereas significantly less AB was transported to the neurites as compared to autophagy-competent neurons (Figure 2E). From these data, we conclude that autophagy influences intracellular transport and secretion of $A\beta$,

Figure 1. A^β Plaque Formation Depends on Autophagy

Scale bars represent 500 μ m (A) and 100 μ m (C). Data are represented as mean \pm SEM. See also Figures S1, S2, and S3.

⁽A) Immunohistological analysis of A β plaque (4G8 antibody) in 20-month-old $Atg7^{flox/flox} \times APP$ and $Atg7^{flox/flox}$; CamKII-Cre × APP mouse brains. A β plaque staining was quantified (n = 4, **p < 0.005).

⁽B) A β ELISA measurements of hippocampal and cortical brain homogenates of 20-month-old $Atg7^{flox/flox} \times APP$ and $Atg7^{flox/flox}$; CamKII-Cre × APP mice (n = 6, *p < 0.05, **p < 0.05).

⁽C) Brain sections of 6-month-old $Atg \overline{\tau}^{flox/flox}$ × APP and $Atg \overline{\tau}^{flox/flox}$; CamKI-Cre × APP mice were immunostained with N1D-A_β antibody (upper panels) and A_β40 antibody (lower panels). Inset shows cortical neurons at 40× magnification. A_β staining in CA1 was quantified (n = 5, *p < 0.05).

⁽D) A BELISA measurements of hippocampal brain homogenates of 6-month-old mice with genotypes as indicated (n = 6, *p < 0.05).

⁽E) APP, APP-CTF, PS1, and BACE1 levels in hippocampal brain homogenates from 6-month-old mice with genotypes as indicated were determined by quantitative western blot (n = 5, no significant difference).





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Figure 2. Autophagy Influences Aß Secretion

(A) Release of endogenous Aβ from Atg7^{flox/flox} and Atg7^{flox/flox}; Nes-Cre cortical/hippocampal primary neurons was determined by ELISA measurements of conditioned media (n = 3, ***p < 0.0001).

(B and C) Lenti-Atg7 expression in primary neurons activates autophagy and increases release of Aβ. Autophagy activation was measured by monitoring LC3 metabolism by quantitative western blot (n = 6, *p < 0.05, **p < 0.01).

(D) Aß ELISA measurements of conditioned media from wild-type primary neurons infected with SFV-APP and treated with pharmacological compounds as indicated (n = 4, *p < 0.05, **p < 0.01). Activation and modulation of autophagy were determined by measuring LC3II/β-actin and p-p70/p70 levels, respectively, by quantitative western blot (n = 6; *p < 0.05, **p < 0.01).

(E) Atg7^{flox/flox} and Atg7^{flox/flox}; Nes-Cre primary neurons were infected with SFV-APP and stained for Aβ (Aβ40 antibody). Three representative neurons per genotype are shown.

Data are represented as mean ± SEM.









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Figure 3. Amyloidosis Inhibits Autophagy and Activates Neurodegenerative Processes

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(A) Western blot analysis of LC3 in cortical brain homogenates from Atg7^{flox/flox} and Atg7^{flox/flox} × APP mice (representative samples from two individuals per genotype are shown). LC3 immunoreactivity was quantified (n = 5, *p < 0.0005).

(B and C) Coimmunostaining of A β (4G8 antibody) and LC3 (B) and A β and p62 (C) using brain sections of 15-month-old Atg7^{flox/flox} × APP mice. (legend continued on next page) through either exocytic or excretory mechanisms, and hence plays a key role in A β plaque formation.

Aβ Amyloidosis Inhibits Autophagy and Exacerbates Autophagy-Deficiency-Induced Neurodegeneration

Given that autophagy is impaired in AD, we investigated the in vivo effects of A β amyloidosis on autophagy in Atg7^{flox/flox} × APP mice. We measured the levels of LC3I and LC3II in brain homogenates and found that LC3II levels were significantly reduced by 50% in $Atg7^{flox/flox}$ × APP mice compared to Atg7^{flox/flox} mice, indicating that amyloidosis suppresses autophagy (Figure 3A; p < 0.0005). Furthermore, LC3 staining was markedly increased in the neuronal cells immediately surrounding the Aβ plaques, suggesting a direct local inhibitory effect of amyloidosis on autophagy (Figure 3B). In addition, LC3 colocalized in certain loci with intracellular Aβ (Figure 3C). A second indication of autophagy inhibition by amyloidosis is the accumulation of p62 found both in the vicinity of and inside the A β plaque (Figure 3C) closely resembling dystrophic neurites, although p62 staining did not directly overlap with phosphorylated tau (data not shown). These data suggest, in agreement with previous studies, an inhibitory effect of amyloidosis on autophagy.

The indication of autophagy-impaired neurons in $Atg7^{flox/flox}$ × APP mice and the fact that autophagy deficiency induces neurodegeneration (Komatsu et al., 2006, Inoue et al., 2012) prompted us to investigate neurodegeneration in the mutant mice. Although no apoptotic DNA fragmentation was found by TUNEL (data not shown), cleaved caspase-3 was detected in hippocampal brain homogenate by western blot and in CA1 pyramidal neurons by immunostaining of autophagy-deficient Atg7^{flox/flox}; CaMKII-Cre mice and was further enhanced in Atg7^{flox/flox}: CaMKII-Cre × APP mice (Figures 3D and 3F; p < 0.01). In addition, Fluoro-jade C-positive staining was observed in Atg7^{flox/flox}; CamKII-Cre mice (Atg7^{flox/flox}; CamKII-Cre × APP mice were omitted from the analysis because Fluoro-jade C stains A_β) indicating activation of necrotic-mediated degeneration. Consistently, receptor-interacting serine/threonine-protein kinase 1 staining was detected in CA1 pyramidal cells of Atg7^{flox/flox}; CamKII-Cre mice and was significantly enhanced in Atg7^{flox/flox}; CamKII-Cre × APP mice (Figures 3E and 3G; p < 0.01). In summary, these data indicate that amyloidosis intensifies autophagy-deficiency-induced neurodegenerative processes.

To determine if activation of neurodegenerative processes leads to brain atrophy, we investigated the brains of the mutant mice by several means. First, a decrease in the wet weights of dissected cortical and hippocampal brain tissue was detected upon deletion of autophagy and the degeneration was further exacerbated by amyloidosis, accompanied by a significantly decreased body weight (Figure 4A; Figure S4A; p < 0.01). Consistently, hematoxylin and eosin (H&E) staining revealed a significant decrease in the size of hippocampus and cortical thickness (measured at posterior parietal associated area) in

Atg7^{flox/flox}; CamKII-Cre × APP mice as compared to Atg7^{flox/flox}; CamKII-Cre mice (Figure 4B; p < 0.05). T2 magnetic resonance imaging (MRI) measurements confirmed a 22% decrease in hippocampal volume of Atg7^{flox/flox}; CamKII-Cre × APP mice as compared to Atg7^{flox/flox} mice (Figure 4C; Figure S4B; p < 0.005) and a trend toward decreased hippocampal volume as compared to Atg7^{flox/flox}; CamKII-Cre mice. Cell counting revealed a 10% loss of p62-positive pyramidal neurons in CA1 of Atg7^{flox/flox}; CamKII-Cre × APP mice as compared to Atg7^{flox/flox}; CamKII-Cre mice (Figure 4D; p < 0.05). In conclusion, amyloidosis exacerbates the autophagy-deficiency-induced neurodegeneration and causes neuronal cell death.

Memory Impairment in Autophagy-Deficient Mice

To analyze the memory effects of intracellular Aß accumulation and amyloidosis-exacerbated autophagy-deficiency-induced neurodegeneration in Atg7^{flox/flox}; CamKII-Cre × APP mice, we subjected 15-month-old littermates to Morris water maze. Whereas Atg7^{flox/flox} mice efficiently learned to find the hidden platform, Atg7^{flox/flox}; CamKII-Cre × APP mice exhibited severe memory impairments (Figure 4E; p < 0.005) and performed significantly worse than $Atg7^{flox/flox} \times APP$ mice (p < 0.05). The performance of autophagy-deficient Atg7^{flox/flox}; CamKII-Cre mice was not significantly different from that of Atg7^{flox/flox} mice (p = 0.28). However, the improvement in learning was modest, suggesting that autophagy deficiency affects memory to some extent. In summary, these results indicate that impaired proteostasis and amyloidosis together severely affect memory either directly or by inducing neurodegeneration. Furthermore, the data highlight that extracellular Aß plagues may not be a critical factor for severe memory impairment and support previous findings that intracellular $A\beta$ is potentially neurotoxic.

DISCUSSION

In this study, we have investigated the in vivo role of autophagy, which is impaired in AD, in A β metabolism by analyzing neuron-specific autophagy-deficient APP mice. Surprisingly, and in contrast to the well-established degradative role of autophagy ending in lysosomal degradation, we found that autophagy influences the secretion of A β . These findings imply that autophagy directly affects two of the hallmarks in AD: intracellular A β accumulation and extracellular A β plaque formation (Figures 1 and 2).

Recently, a role for autophagy in protein secretion has emerged (reviewed in Deretic et al., 2012). Autophagy participates in nondegradative secretion of integral membrane proteins directly from endoplasmic reticulum (ER), through ER-to-Golgito plasma membrane (PM) secretory pathway or via secretory lysosomes. APP has previously been shown to be transported and processed to A β through the ER-Golgi-to-PM secretory pathway or transported unprocessed to PM, where A β is generated after endocytosis. Autophagosomes are sites of A β

⁽D and E) Immunostaining for cleaved caspase 3 (D) and RIPK1 (E) of 15-month-old brain sections with genotypes as indicated. The intensities were quantified (n = 5, *p < 0.01).

⁽F) Quantitative western blot analysis of cleaved caspase 3 (n = 3, *p < 0.05).

⁽G) Fluor-jade C staining of 15-month-old brain sections with indicated genotypes. Insets show 40× magnification.

Scale bar represents 50 µm (B), 25 µm (C, upper panel), 4 µm (C, lower panel), and 100 µm (D, E, and G). Data are represented as mean ± SEM.





Figure 4. Amyloidosis Exacerbates Autophagy-Deficiency-Induced Neurodegeneration

(A) Wet weights of dissected brain tissue from 3- and 15-month-old mice with genotypes as indicated (n = 5/genotype).

(B) Representative sections stained by H&E from 15-month-old mice with genotypes as indicated. The cortical and hippocampal thickness were quantified.

(legend continued on next page)

generation; it remains to be determined if any of these secretory pathways is influenced by autophagy. In this context, it is noteworthy to mention that autophagosomes are formed at the ER-mitochondria contact site (Hamasaki et al., 2013). Because autophagosomes can fuse with endosomes at the late stage of autophagy, Aβ-containing endosomes could potentially be the source of Aβ released to extracellular space. However, it cannot be excluded that extracellular release of Aβ influenced by autophagy could be part of a general excretory mechanism for cellular waste, independent of the regulated secretory pathways.

If the increased number of autophagosomes observed in AD is due to increased autophagy, then it would result in increased Aß secretion. On the other hand, if the autophagosomes accumulate due to impairment in the end stage of autophagosome clearance, intracellular Aß levels would rise. Indeed, neurons in the AD brain exhibit intraneuronal Aß accumulation, and recent data suggest that intracellular AB causes neurodegeneration by increased ER stress. We found that neurodegeneration induced by autophagy deficiency was exacerbated by amyloidosis and that these two pathologies, together with impaired proteostasis, caused severe memory impairment (Figures 3 and 4). The enhanced neurodegeneration and memory impairment could potentially be explained by the intracellular Aß accumulation, given the sparse A β plaque load in Atg7^{flox/flox}; CamKII-Cre × APP mice, which would indicate that intracellular $A\beta$ is toxic. More research is warranted to elucidate how intracellular AB mediates toxicity.

Administration of autophagy-activating rapamycin clears intracellular A β and improves cognition in the 3×Tg-AD mice (Caccamo et al., 2010), raising the question if such treatment is applicable to AD. Inducing autophagy would clear potentially neurotoxic intracellular A β at the expense of increased A β release. Hence, coadministration of an A β -lowering treatment would be needed to effectively prevent extracellular amyloidosis. In conclusion, our data establish that autophagy influences A β transport and release to the extracellular space and thereby directly affects A β plaque formation. Thus, autophagy plays a crucial role in AD pathology and could be a potential AD drug target.

EXPERIMENTAL PROCEDURES

Animals

Atg7^{flox/flox} mice (Komatsu et al., 2005) were kindly provided by Dr. Keiji Tanaka (Tokyo Metropolitan Institute of Medical Science) and CamKII-Cre mice (Tsien et al., 1996) were kindly provided by Dr. Shigeyoshi Itohara (RIKEN Brain Science Institute). APP23 mice have been described previously (Sturchler-Pierrat et al., 1997). All animal experiments were carried out according to the guidelines of RIKEN Brain Science Institute.

Immunoblot Analysis

A total of 10 μg of brain homogenates was separated by SDS-PAGE, transferred to membranes, and probed with antibodies as listed in Table S1. For detection of cleaved caspase-3, the membrane was incubated with 1% glutaraldehyde in PBS before processing.

Histochemical and Immunohistochemical Analysis

Sections 4 µm (paraffin embedded) or 10 µm (fresh frozen) thick were stained with H&E, cresyl violet, toluidine blue, or fluoro-jade C (AG325, Chemicon), or immunostained using antibodies listed in Table S1. For Aβ immunostaining, tissue sections were treated with 90% formic acid for 5 min. Quantification was performed with MetaMorph imaging software (Universal Imaging). For the cell counting experiment, PFA-fixed sections 5 µm thick were collected from bregma -0.90 to -4.40 with 50 µm intervals and stained by p62 and Hoechst. p62-positive neurons were manually counted with blinded samples.

Primary Neuron Culture

Cortical/hippocampal neurons were prepared from embryonic day 17 to 18 (E17-E18) mouse embryos as previously described (Hama et al., 2001). Embryos from Atg7^{flox/flox} ; Nes-Cre mice were separately genotyped. A total of 1.8×10^5 vital cells were plated in 24-well plates by trypan blue staining a 10 µl cell suspension and counting vital cells in a hemacytometer. After 10 days in vitro, endogenous Aß levels in conditioned media were measured by ELISA by centrifugation of the media for 1 min at 3,000 rpm. Guanidin-HCI was added to prevent aggregation (final concentration 0.5 M) or the cells were infected with 10 µl semliki forest virus (SFV) expressing APP. Then, 24 hr postinfection, conditioned media were collected followed by A β ELISA measurement. Same conditions were applied for autophagy activation/inhibition experiments with rapamycin (2.7 nM, Sigma), vinblastine (50 µM, Sigma), and Spautin-1 (10 µM, BioVision Technologies) added to the media 23 hr postinfection to assure full effect of the compounds prior to A β measurements. The media was changed 24 hr postinfection to inhibitor-containing media. Three hours later, A $\!\beta$ levels in conditioned media were measured by ELISA. The cells were collected in PBS, centrifuged, and dissolved in SDS sample buffer containing 0.1 M DTT for subsequent western blot analysis of APP expression analysis, to which $A\beta$ levels were normalized.

Immunofluorescence

Primary neurons were infected with SFV-APP for 24 hr, fixed for 10 min in cold 4% paraformaldehyde and 0.1 M phosphate buffer, washed with PBS, blocked with 5% NGS and 1% saponin, and incubated with antibodies overnight. Wash buffer and antibodies dilutions contained 1% saponin.

ELISA

TS and GS A β from cortical and hippocampal homogenates (Iwata et al., 2004) and conditioned media were determined by ELISA (Wako or IBL) according to the manufacturer's instructions.

Morris Water Maze

Fifteen-month-old littermates $(Atg7^{flox/flox}, Atg7^{flox/flox}; CaMKII-Cre, Atg7^{flox/flox} \times APP, Atg7^{flox/flox} : CaMKII-Cre \times APP, n = 10) were acclimatized to the behavioral laboratory 3 days before tests. The light condition was 12 hr:12 hr (lights on at 8:00 am). Tests were performed from 9:30 am to 3:30 pm. Each mouse was assessed in two training sessions per day. Mice had ad libitum access to food and water (except during the tests).$

MRI

Fifteen-month-old *Atg7*^{flox/flox}, *Atg7*^{flox/flox}; CaMKII-*Cre*, *Atg7*^{flox/flox} × APP, *Atg7*^{flox/flox}; CaMKII-*Cre* × APP (n = 3/genotype) mice were anesthetized with isoflurane (1.5%–2% in air) and mounted in a stereotaxic apparatus. The depth of anesthesia was monitored with a breathing sensor. MRI scans of the whole brain were performed with a vertical-bore 9.4 T Bruker AVANCE 400WB imaging spectrometer with a 250 mT m⁻¹ actively shielded imaging gradient insert (Bruker BioSpin) controlled by Paravision software. T2-weighted images were obtained with the following parameter settings: matrix dimensions = 256 × 256 × 29, TE = 53.5 ms, TR = 434.2 ms, flip angle = 180

(C) Hippocampal volume was calculated from T2 MRI data.

(D) Cell counting of p62-positive neurons in CA1 of 20-month-old Atg7^{flox/flox}; CamKII-Cre and Atg7^{flox/flox}; CamKII-Cre × APP mice.

(E) Morris water maze was performed with 15-month-old littermates with genotypes as indicated (n = 10). The performance on day 5 was analyzed by ANOVA followed by Tukey's post hoc test.

Scale bar represents 500 μ m. Data are represented as mean \pm SEM. *p < 0.05, **p < 0.01, ***p < 0.005. See also Figure S4.

degrees, FOV = 16 × 16 × 0.5. A total of 29 high-resolution coronal slices of the whole brain were collected. Total scanning time was 60 min per individual. Within the 29 MRI slices, 8 slices contained the hippocampal formation (from bregma -0.90 to -4.40), which was manually selected by the paintbrush tool and used for the volumetric calculation of hippocampus using InsightITK-Snap software version 2.2.0.

Statistical Analysis

Data were analyzed by Student's t test if not stated otherwise and are presented as average \pm SEM.

SUPPLEMENTAL INFORMATION

Supplemental Information includes four figures and one table and can be found with this article online at http://dx.doi.org/10.1016/j.celrep.2013.08.042.

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