REVIEW PAPER

Autophagy and Alzheimer's Disease

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Abstract Autophagy is an essential degradation pathway in clearing abnormal protein aggregates in mammalian cells and is responsible for protein homeostasis and neuronal health. Several studies have shown that autophagy deficits occurred in early stage of Alzheimer's disease (AD). Autophagy plays an important role in generation and metabolism of β -amyloid (A β), assembling of tau and thus its malfunction may lead to the progress of AD. By considering the above evidences, autophagy may be a new target in developing drugs for AD. So far, a number of mammalian target of rapamycin (mTOR)-dependent and independent autophagy modulators have been identified to have positive effects in AD treatment. In this review, we summarized the latest progress supporting the role for autophagy deficits in AD and the potential therapeutic effects of autophagy modulators in AD.

Keywords Autophagy · β-Amyloid · Tau · Alzheimer's disease

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Introduction

AD, which is the most prevalent neurodegenerative disease, is characterized by deficiency in memory and cognitive functions. The predominant pathological changes of AD are development of \overrightarrow{AB} plaques in specific brain areas, neurofibrillary tangles in neurons, and progressive loss of synapses and neurons (Ingelsson et al. [2004\)](#page-9-0). Although the etiology mechanisms underlying these pathological changes are not clear yet, recent studies have shown deficits in the autophagy-lysosome pathway, which is an important system to clear misfolded proteins or damaged organelles, is likely to precede the formation of $\mathbf{A}\beta$ plaques or neurofibrillary tangles (Correia et al. [2015\)](#page-9-0). Until now, three types of autophagy: macroautophagy, microautophagy, and chaperone-mediated autophagy (CMA), have been identified in mammalian cells. Among them, macroautophagy is best studied and considered most relevant to AD, CMA has also been thought to play a role in AD. To make it simple, hereon we will refer macroautophagy as ''autophagy.'' In this review, we begin with the autophagy physiological machinery, evidence of autophagy malfunction in AD, relationship between autophagy dysfunction and AD-related pathology, and finally focus on the therapeutic potential of autophagy modulators.

Autophagy Machinery

There are several stages for autophagy biogenesis, including phagophore membrane isolation, phagophore elongation and engulf of random cytoplasmic content, and autophagosome maturation and fusion with lysosome (Lee et al. [2013\)](#page-9-0) (Fig. [1](#page-1-0)). Generally, the autophagy activation is largely depended on cellular starving condition (Kuma

et al. [2004,](#page-9-0) Mizushima et al. [2004](#page-10-0)), such as low amino acids or glucose concentration. However, recent studies have also shown autophagy could be induced by diverse stimulations, such as reactive oxygen species (ROS) (Scherz-Shouval et al. [2007](#page-11-0)), hypoxia impairments (Mazure and Pouysségur 2010), subcellular organelle damages (Song et al. [2013](#page-11-0)), and protein aggregation (Liu et al. [2010](#page-10-0)). However, in all the regulators of autophagy, the mammalian target of rapamycin complex 1 (mTORC1) has been studied most thoroughly and considered to play a key role in autophagy biogenesis.

In amino acid depletion environment, mTORC1 complex is de-activated and distributed freely in cytoplasm, where it cannot inhibit autophagy initiation. However, in nutrition-rich condition, the lysosomal amino acid concentration is largely increased, which in turn stimulates v-ATPase on the lysosome membrane. v-ATPase is a lysosome membrane multimeric channel protein majorly functions as ATPase and H^+ channel, meanwhile, it also functions as a sensor for lysosomal amino acid concentration. After sensing the increase of amino acid concentration, v-ATPase and its binding partner, Ragulator, would both change their conformation and thus activate downstream signal molecule: RagA/B, by converting its conformation from GDP-bound to GTP-bound. Activated RagA/B then recruit active mTORC1 onto lysosome membrane, on where mTORC1 could be activated by Rheb, a small GTPase attached on lysosome membrane (Zoncu et al. [2011;](#page-11-0) Jewell et al. [2013;](#page-9-0) Yadav et al. [2013](#page-11-0)). After mTORC1 activation, it will negatively control autophagy genesis by blocking Unc-51-like kinase (ULK)1/2 complex, which is an important inducer for autophagy by regulating Beclin1-VPS34 complex (Russell et al. [2013](#page-10-0)). On the other hand, activated ULK1/2 could active microtubule-associated protein 1 light chain 3 beta (LC3B) by phosphatidylethanolamine (PE) (Fujita et al. [2008](#page-9-0)). LC3 is important for phagophore elongation and finally form autophagosome (Bernard and Klionsky [2014](#page-9-0)). Furthermore, activated mTORC1 could also inhibit autophagosome fusion with lysosome by phosphorylating UV radiation resistance-associated gene protein (UVRAG) (Kim et al. [2015](#page-9-0)) and thus block autophagosome maturation. Therefore, activated mTORC1 could inhibit all autophagy biogenesis required stages, and inhibition of mTOR pathway by either pharmacological compounds such as rapamycin or activation of AMP-activated protein kinase (AMPK) (Wu et al. [2015](#page-11-0)) could facilitate autophagy process.

Autophagy in Neuronal Cells

Autophagy is an essential homeostatic pathway in neurons (Maday et al. [2012](#page-10-0)). Similar to other cells, with aging process, neuronal cell will also accumulate intracellular toxicant or damaged organelles such as mitochondria that must to be cleared by autophagy to maintain an appropriate intracellular homeostasis (Mariño et al. [2011\)](#page-10-0). However, unlike other cell types, neurons are post-mitotic cell that can never dilute toxic substance by mitosis. So autophagydependent protein/organelles clearance would be more important in neurons.

Neurons are characterized by their highly polarized axons and dendritic compartments, by which they can extend over distances many times larger than their cell soma and thus connect to each other from long distance with minimized cell number. Autophagosomes in neurons majorly formed at axon tip and gradually matured when they are retrograde transporting along axons (Maday et al. [2012\)](#page-10-0). There is also evidence that autophagosome formation can be induced along the axon in order to clear damaged mitochondria (Ashrafi et al. [2014\)](#page-8-0). Furthermore, some research indicated distal biogenesis of autophagosome might be related with synaptic function (Wang et al. $2015a$, [b](#page-11-0)). The first step of autophagosome biogenesis at axon tip is recruiting Atg13 and Atg5 to double-FYVE containing protein 1 (DFCP-1), which is a phosphatidylinositol 3-phosphate (PI3P)-enriched omega-shaped ER structure that serves as a platform for autophagosome biogenesis (Maday and Holzbaur [2014\)](#page-10-0). Then in 4–6 min, lipidated LC3 was incorporated into the developing autophagosome (Maday et al. [2012\)](#page-10-0). Autophagy is more efficient in young neurons (Boland et al. [2008](#page-9-0)), as autophagy-related proteins such as beclin-1, Atg5, and Atg7 will decline with age (Lipinski et al. [2010](#page-10-0); Shibata et al. [2006](#page-11-0)), which potentially contributing to the late onset of many neurodegenerative diseases (Rubinsztein et al. [2011\)](#page-10-0).

Autophagy Malfunction in AD

There is substantial evidence that dysregulation of autophagy occurs in both AD animal models and AD patients. As early as 1967, Suzuki had found a large amount of abnormal aggregated tau protein and subcellular vesicles accumulated in the dystrophic or swollen neurites in AD patient brains (Suzuki and Terry [1967\)](#page-11-0), but the identity of these vesicles was unclear until 2005, when Nixon's group found immature autophagic vacuoles (AVs) accumulated in dystrophic neuritis in AD brains using immunogold labeling and electron microscopy (Nixon et al. [2005\)](#page-10-0). This is the first direct evidence that autophagy deficiency was involved in AD. In the same year, similar results from PS-

1/APP double transgenic mice (Yu et al. [2005](#page-11-0)) also showed AVs accumulated in neuronal dendrites and soma even before \overrightarrow{AB} plaques appeared compared to age-matched controls. In addition to these direct ultramicroscopic results, changes in the expressions of several autophagyrelated proteins also indicated autophagy dysfunction in AD. For example, Rubinsztein reported increased expression level of lysosomal protease in the early phase of AD patients (Rubinsztein et al. [2005\)](#page-10-0). A recent study found that expression of the autophagy-related genes: atg1, atg8a, and atg18 in Drosophila melanogaster were down-regulated with aging, and the subsequent reducing of autophagy activity and hyper-generation of $\mathbf{A}\boldsymbol{\beta}$ were both considered to correlate with late-onset neuronal dysfunction and AD phenotype (Omata et al. [2014\)](#page-10-0), indicating that age-induced reduction of autophagy-related gene expression is associated with late onset of AD.

There are two types of AD identified: sporadic AD (SAD, also known as late-onset AD), and familial AD (FAD, also known as early-onset AD). In SAD, little is known about the cause of the onset, but it is widely accepted that both genetic and environmental factors contribute to this pathogenesis. Apolipoprotein E4 (apoE4), a major genetic risk factor for SAD, has been found to induce autophagy malfunction. Studies from ApoE4 transgenic mice showed that overexpression of ApoE4 elevated $A\beta42$ amount in lysosome and finally led to hippocampus neurons death (Belinson et al. [2008](#page-8-0)). In addition, ApoE4 potentiates lysosomal leakage and apoptosis caused by $A\beta$ peptides in Neuro-2a cells(Ji et al. [2006](#page-9-0)). Taken together, these studies indicated that ApoE4 and \overrightarrow{AB} may work in concert to increase the susceptibility of lysosomal membranes disruption, release of lysosomal enzymes, and hence neuronal degeneration (Ji et al. [2006\)](#page-9-0).

Regarding to FAD, at least three genes, amyloid precursor protein (APP), presenilin-1 and -2(PS-1 and PS-2), have been identified as causative genes so far. FAD is caused by mutation in at least one of the three genes (Tang [2003](#page-11-0)). Research has found wildtype PS-1, but not mutation forms, is crucial for acidification of lysosome by regulating distribution of v-ATPase, and thus contributes to autophagy degradation in a gamma-secretase-independent way (Lee et al. [2010\)](#page-9-0). Meanwhile, hyper-activation of Glycogen synthase kinase-3 (GSK-3), which is also a high-risk factor for AD, could interfere lysosome acidification via similar mechanism as PS-1 (Avrahami et al. [2013\)](#page-8-0). These investigations suggested that autophagy malfunction is involved in FAD, while the mechanism(s) is still not clear.

Proper formation and degradation of autophagosome is critical for normal autophagic flux. In healthy neurons, low-basal autophagic activity was detected because of the quick subsequent degradation of autophagosome by lysosome. In hippocampus neurons of AD mice, abnormal

accumulation of immature AVs in axon was observed before synaptic and neuronal loss (Sanchez-Varo et al. [2012\)](#page-10-0). However, as either autophagosome axonal trafficking deficiency or insufficient lysosome acidification could cause AVs accumulation in axon, the actual mechanisms underlying autophagy dysfunction in AD is still need further investigate. It is also a debate whether dysfunction of autophagy is the cause or result of AD. Reports to date usually show some controversies (Peric and Annaert [2015](#page-10-0); Cho et al. [2015;](#page-9-0) Lee et al. [2014\)](#page-10-0). Many factors may lead to these differences, such as different animal models, cellular models, and experimental paradigms. Also, the different model systems are likely represent of different particular stages of AD pathogenic process.

CMA and AD

Chaperone-mediated autophagy (CMA) is another form of autophagy which functions quite distinct from macroautophagy, as CMA pathway will degrade certain proteins one by one without engulf process, and most of its targeting proteins would be cytoplasm soluble ones. This selective degradation pathway contributes to protein quality control and maintenance of cellular homeostasis particularly under stress conditions (Cuervo [2010](#page-9-0); Dice [2007](#page-9-0); Kaushik and Cuervo [2012](#page-9-0)). CMA-mediated degradation needs heat shock cognate protein of 70KD (hsc 70), which is a cytosolic chaperone, to recognize its targets. Most of hsc 70 target proteins share a reserved KFERQ sequence (Chiang et al. [1989\)](#page-9-0). After binding to its target, cytoplasmic hsc70 will then mediate target protein unfolding and deliver it onto lysosome membrane, where lysosomal-associated membrane protein 2A (LAMP-2A) will form a transient multimeric channel to allow them to translocate through lysosome membrane. lysosome-resident form of hsc70 (lys-hsc70) in lysosome luminal side will also assist this process (Cuervo and Dice [1996](#page-9-0); Cuervo et al. [1997](#page-9-0); Agarraberes et al. [1997](#page-8-0)). It has been considered that the number of lys-hsc70 positive lysosome would decide the speed of CMA-mediated degradation under physiology condition.

Similar to macroautophagy, CMA is also considered to connect with neurodegeneration diseases such as AD and PD, as CMA plays an important role in both Tau tangles and \overrightarrow{AB} plaques generation and its activity is impaired with aging process (Koga and Cuervo [2011](#page-9-0)). It has been found although Tau bears a CMA recognizing motif in its sequence, trace amount of wildtype Tau is degraded by CMA. Normally, when facing degradation, Tau will be first cleaved in cytoplasm and then remained c-terminal part will translocate into lysosome luminal either through autophagy or CMA, where the second and third cleavage will happen (Wang et al. [2009\)](#page-11-0). However, in certain Tau mutations, the first cleavage will happen in wrong amino acid position. After this cleavage, although remaining c-truncated Tau could still be recognized by hsc70 and recruited onto lysosome membrane, it cannot translocated into lysosome luminal efficiently, thus resulting in accumulation of truncated Tau on lysosome membrane. Some of these products organize as oligomeric structures at the lysosomal membrane and with time, these oligomers can promote disruption of the lysosomal membrane and the subsequent leakage of lysosomal enzymes (Koga and Cuervo [2011\)](#page-9-0).

Another recent study found APP also bears KFERQ motif which could be recognized by hsc70. Deletion of this sequence will keep APP away from lysosome and increase its secretase cleavage products, without influence its ability to bind to hsc70 (Park et al. [2016\)](#page-10-0).

Relationship Between Autophagy Dysfunction and AD-Related Pathology

Ab, Tau, and Autophagy

As mentioned above, \overrightarrow{AB} plaques and neurofibrillary tangles are the two major neuro-pathological changes occurred in AD patients. Emerging evidence has demonstrated complicated interactions among autophagy, \overrightarrow{AB} , and tau, which may contribute to the progress of AD.

Autophagy plays an important role in the metabolism of A β . First, along with A β degradation enzymes (Miners et al. [2008](#page-10-0)), autophagy is believed to be another major $\mathbf{A}\mathbf{\beta}$ clearance pathway. Autophagy facilitates the degradation and clearance of APP (Aymloid- β precursor protein) (Zhou et al. [2011\)](#page-11-0) as well as all APP cleavage products including $\text{A}\beta$ (Son et al. [2012\)](#page-11-0) and APP-CTFs (amyloid precursor protein-cleaved C-terminal fragment) (Tian et al. [2014](#page-11-0)). In microglia, $\mathbf{A}\beta$ is also degraded by autophagy through the autophagy receptor optineurin (Cho et al. [2014\)](#page-9-0).

Second, autophagy-lysosome system, which is important for \overrightarrow{AB} degradation under physiological conditions, is demonstrated to be a novel way for \overrightarrow{AB} production under pathological condition or in aging process (Yu et al. [2005](#page-11-0)). Although \overrightarrow{AB} is thought to be produced in lysosome, endoplasmic reticulum, and Golgi apparatus, emerging studies have provided evidence that $\mathbf{A}\beta$ generation could be detected in autophagic vacuoles after autophagy activation (Mizushima [2005](#page-10-0)). Thus, accumulated immature autophagic vacuoles found in AD brains and in APP/PS1 transgenic mice maybe a source of \overrightarrow{AB} generation (Nixon et al. [2005\)](#page-10-0). Immunohistochemistry staining showed that $A\beta$ 42 were stained in AEL (autophagy–endosomal–lyso-somal) vesicles in Drosophila neurons (Ling et al. [2014](#page-10-0)). Another study found in neuron that although APP and its processing enzyme BACE1 should be separated into distinct vesicles under normal condition, they showed strong co-localization and co-trafficking into autophagy–lysosome pathways under glycine/NMDA-receptor agonist/ $K+$ or GABAA antagonist Picrotoxin (PTX) stimulations (Das et al. [2013\)](#page-9-0). This result indicated under some certain pathological conditions, the convergence of APP and BACE1 in autophagosomes may act as some new $\mathbf{A}\mathbf{B}$ generation places.

Third, autophagy is also involved in the secretion of $A\beta$. Recent findings support that autophagy is responsible for extracellular release of Aβ. Measurement of extracellular $\text{A}\beta$ in autophagy-deficient mice revealed that the $\text{A}\beta$ secretion was reduced by 90 %, while restoration of autophagy enhanced Ab secretion to normal levels (Nilsson and Saido [2014](#page-10-0)). There is another recent work has found in ATG7 knockdown mice, the secretion \overrightarrow{AB} was largely reduced, which was accompanied by a significant intracellular $\mathbf{A}\beta$ accumulation (Nilsson, et al. [2015](#page-10-0)).

On the other side, \overrightarrow{AB} could also regulate autophagy. $A\beta40$ in vascular could induce autophagy in endothelial cell and impair neurovascular regeneration (Hayashi et al. [2009\)](#page-9-0). Further study showed \widehat{AB} -induced formation of AVs was regulated through the RAGE-calcium-CaMKK β -AMPK pathway (Son et al. [2012](#page-11-0)).

For the case of tau, although ubiquitin–proteasome system (UPS) was considered to be the major pathway for tau turnover, recent studies suggested that autophagy maybe another effective degradation route for tau. A number of studies have demonstrated that dysfunction of autophagy-lysosome system leads to the formation of tau oligomers and insoluble aggregates, while induction of autophagy could alleviate this aggregation (Hamano et al. [2008,](#page-9-0) Congdon et al. [2012](#page-9-0)). Moreover, autophagy may affect phosphorylation status of tau. Hyperphosphorylated tau was found to be accumulated in brains of autophagydeficient mice (Inoue et al. [2012](#page-9-0)), and this phospho-tau accumulation will be largely reduced after autophagy was restored. Researchers have found autophagic degradation of tau is regulated by nuclear factor erythroid-2-related factor 2 (Nrf2)-mediated activation of the autophagy receptor NDP52 (Jo et al. [2014](#page-9-0)). Furthermore, Phosphatidylinositol binding clathrin assembly protein (PICALM, also known as CALM) could also regulate Tau degradation by modulating SNAREs (Soluble NSF Attachment protein Receptors) endocytosis, which is critical for autophagy clearance of tau (Moreau et al. [2014](#page-10-0)). On the other hand, hyperphosphorylation of tau may result in autophagy dysfunction (Lim et al. [2001;](#page-10-0) Lin et al. [2003](#page-10-0)). Tau is well identified to facilitate the assembly and stabilization of microtubule, which is critical for autophagosome retrograde trafficking and maturation to fuse with lysosome. On contrast, hyperphosphorylated tau could lead

to the instability and disassemble of microtubule cytoskeleton, which could subsequently inhibit autophagosome retrograde trafficking and thus accumulate immature autophagosomes in axons.

Axonal Transfer and Autophagy

Axonal transport is an essential process required to maintain neuronal homeostasis. Impaired axonal transport can lead to axon degeneration and has been found in many neurodegenerative diseases including AD. In mammalian cells, newly formed autophagosomes move along with microtubule tracks (Monastyrska et al. [2009](#page-10-0)), and during this process, autophagosomes engulf long-lived protein, misfolded protein, or damaged organelles—such as mitochondria—subsequently degrade them after fused with lysosome. Deficits in axonal transport usually result in accumulation of large amounts of autophagosome. Axonal dystrophic neurites arising from neurofibrillary tangles could be easily detected in hippocampus CA3 and CA1regions (Su et al. [1993\)](#page-11-0).

Years of pathological examination in AD brains have yielded many descriptions of abnormal axonal transport in both early and late phase of AD (Bell and Claudio [2006](#page-9-0)). Meanwhile, the phosphorylated tau level affects axonal transport and degradation (Rodríguez-Martín et al. [2013](#page-10-0)). These data supported that abnormal protein aggregates disrupt axons, thereby autophagosomes could not transport to cytoplasm and thus failed to fuse with lysosomes. In contrast to this, some other studies gave opposite viewpoints that lysosomal protease abnormalities are the causative factor of axonal degeneration (Xie et al. [2015](#page-11-0)). However, the mechanism(s) underlying this transport disruption is not clear.

Autophagy and Treatment

Currently, most drugs available for AD treatment are developed based on cholinergic hypothesis. Some Chinese traditional herbs also show potential effects in AD treatment (Yu et al. [2012,](#page-11-0) Sun et al. [2012\)](#page-11-0). However, these drugs can only alleviate some symptoms of AD. Development of innovative medicines to prevent AD pathogenesis has always been a hotspot field. As AD is caused by abnormal protein aggregation, new strategies that can enhance the degradation of toxic aggregates are essential for effective therapy. Autophagy, which represents a major route for clearance of aggregated proteins and organelles, may serve as an emerging and promising therapeutic target for blocking AD pathogenesis. Although some autophagy inducers showed promising effect in other neurodegeneration disease, research is still underway to identify

selective autophagy regulator as potential drugs for AD treatment.

Pharmacological Chemicals Targeting mTOR-Dependent Autophagy Pathway

mTOR is a protein kinase that senses cellular energy availability, and regulates cellular proliferation. mTOR also serves as a major negative regulator in autophagy. In mammalian cells, mTOR could enhance the binding between ULK and ATG13, probably via phosphorylation of ULK. Previous reports have shown that mTOR signaling is hyperactive in selected regions of AD brains (An et al. [2003;](#page-8-0) Pei and Hugon [2008](#page-10-0)); genetic reduction of mTOR signaling in the brains of Tg2576 mice resulted in reduced \overrightarrow{AB} deposits and rescued memory deficits by increasing autophagy induction and restoring the hippocampus gene expression signature (Caccamo et al. [2014\)](#page-9-0). All these results demonstrated that mTOR hyperactivity could result in $\mathbf{A}\boldsymbol{\beta}$ accumulation.

As a major regulator of the autophagy, mTOR is widely considered as a pharmacological target for autophagy regulation (Fig. 2). Moreover, research conducted in recent years revealed a potential role for mTOR-related proteases as anti-AD drug targets. Until now, amounts of chemicals that can regulate mTOR have been identified and some of

them may become the potential candidates for AD treatment.

The beneficial effects of autophagy up-regulation in neurodegenerative diseases first came with the mTORC1 inhibitor rapamycin. Rapamycin is a US Food and Drug Administration (FDA)-approved antifungal antibiotic, anticancer cytostatic agent, and an immunosuppressant. In mammalian cells, rapamycin forms a complex with the immunophilin FKBP12 (FK506-binding protein of 12 kDa), thereby stabilizing raptor (Regulatory-associated protein of mTOR)-mTOR association and inhibiting mTORC1 kinase activity, which could then subsequently induce autophagy initiation. Some studies revealed a potential beneficial effect of inducing autophagy with rapamycin in AD. In two AD mouse models, rapamycin administration both resulted in reduced Ab generation, lower tau accumulation and better cognitive performance (An et al. [2003;](#page-8-0) Pei and Hugon [2008](#page-10-0)). However, although drugs that target proteins involved in mTOR signaling pathway have clinical use potential, mTOR pathway itself involved in many critical cellular processes including cell growth and gene translation, so long-term inhibition of mTOR may cause toxic side effects on patients. Therefore, rapamycin is not considered as an ideal drug for long-term use.

Temsirolimus, which is a newly developed compound approved by FDA and European Medicines Agency for

renal cell carcinoma treatment, could enhance Ab clearance in HEK293-APP695 cells and in brains of APP/PS1 mice in an autophagy-dependent manner (Jiang et al. [2014a](#page-9-0)). Furthermore, Results showed temsirolimus could enhance autophagic clearance of hyperphosphorylated tau in either okadaic acid-administrated SH-SY5Y cells or P301S transgenic mice brain (Jiang et al. [2014b](#page-9-0)). Meanwhile, behavioral tests showed temsirolimus improved spatial learning and memory abilities in both APP/PS1 mice and P301S transgenic mice. These results strongly indicated temsirolimus may use as a potential therapeutic strategy for AD treatment (Jiang et al. [2014a,](#page-9-0) [b\)](#page-9-0).

However, there are many issues to be considered and confirmed before these candidate drugs to be applied in clinical therapies. First, whether the beneficial results observed in mice and tissue cultures can be also observed in human remains unknown. Sometimes even chemicals that have beneficial effects in small clinical trials may fail in the subsequent large sample trials. For example, Latrepirdine (Dimebon), which is a mTOR targeted drug, significantly alleviated cognitive deficiency in mild to severe AD patients in a small trial, failed in a phase III trial later (Sweetlove 2012). Second, as we mentioned above, because mTOR is such an important molecule that is involved in many essential cellular functions, treatment by those mTOR inhibitors over a prolonged period might be harmful to patients. Therefore, clinical studies are needed to be conducted carefully to adjust the adverse effects of these kind of chemicals.

Pharmacological Chemicals Targeting mTOR-Independent Autophagy Pathway

In addition to the well-defined mTOR pathway, there are several mTOR-independent signaling pathways that are involved in autophagy initiation (Fig. [3\)](#page-7-0), such as inositol signaling pathway, Ca^{2+}/ca lpain pathway, cAMP/Epac/ Ins(1,4,5)P3 pathway, and JNK1/Beclin-1/PI3KC3 pathway. There are several FDA-approved drugs that have been reported to regulate autophagy through mTOR-independent pathways (Williams et al. [2008](#page-11-0)). In this section, we will highlight some examples in this category.

Inositol Signaling Pathway

The first evidence for existing a mTOR-independent pathway came from study that indicated intracellular IP_3 levels negatively regulate autophagy. IP_3 is a well-known signaling transducer produced by PIP_2 . IP₃ can be hydrolyzed into free inositol that is essential for the inositol signaling pathway. Lowering intracellular inositol or $IP₃$ levels could induce autophagy, while elevation of them could inhibit autophagy by inhibiting autophagosome synthesis.

The most well-studied chemical that can induce autophagy through inositol signaling pathway is lithium, which is a mood stabilizer that can inhibit inosito monophosphatase (IMPase) and lower inositol levels. Although it was used to treat affective disorders for more than a half-century, whether lithium could have positive effects on AD patients is still unclear. Some studies have shown longterm use of lithium is associated with a lower risk of dementia in bipolar disorder patients (Nunes et al. [2007](#page-10-0); Kessing et al. [2010\)](#page-9-0). However, contradictory evidences suggested lithium was not useful in the treatment of AD patients (Hampel et al. [2009](#page-9-0); Macdonald et al. [2008](#page-10-0)). Further studies are needed to assess the effect of lithium on AD.

Another autophagy inducer targeting this pathway is scyllo-Inositol (SI), which has shown potential benefit to AD patients in clinical trials. In addition to its inhibitory effect on aggregation and fibril formation of $\mathbf{A}\mathbf{\beta}$, SI treatment also decreased the size and the number of accumulated AVs in TgCRND8 mouse model (Lai and McLaurin [2012](#page-9-0)), suggesting beneficial effect of $SI-A\beta$ interaction may resolve autophagy deficiency in the AD brains. Other inducers include L-690,330, carbamazepine, and sodium valproate, which can enhance the clearance of mutant Huntingtin fragments and attenuate polyglutamine toxicity (Nalivaeva et al. [2009](#page-10-0), Sarkar and Rubinsztein [2008](#page-10-0)). Studies have also shown sodium valproate and carbamazepine could inhibit \overrightarrow{AB} aggregates and thus reduce cell toxicity (Nalivaeva et al. [2009,](#page-10-0) Li et al. [2013\)](#page-10-0). However, further investigations are needed to confirm whether these chemicals have positive effects on AD patients.

$Ca^{2+}/Calpain$ Pathway

Elevation in cytosolic Ca^{2+} concentration could inhibit autophagy, while decrease of it enhances autophagy. Cytosolic Ca^{2+} affects autophagy at both the formation and autophagosome–lysosome fusion stages. Several FDA-approved L-type Ca^{2+} channel antagonists such as verapamil, loperamide, amiodarone, nimodipine, and nitrendipine, have been screened as enhancer of autophagy (Williams et al. [2008\)](#page-11-0), but the effects of these antagonists on AD have not been well-studied. One potential drug is Isradipine, which is a FDA-approved dihydropyridine calcium channel blocker, attenuates $\mathbf{A}\boldsymbol{\beta}$ oligomer toxicity by suppressing calcium influx into cytoplasm and suppressing $CaV_{1,2}$ expression in vitro and in AD animal models, suggesting that isradipine exhibited bio-availability, lowered \overrightarrow{AB} plaque burden, and improved autophagy function (Anekonda and Quinn [2011](#page-8-0)).

Fig. 3 mTOR-independent autopahgy pathways

Other mTOR-Independent Pathways

There are many other kinds of drugs that target mTORindependent pathways such as histone acetyltransferase inhibitors (Eisenberg et al. [2009](#page-9-0)), AMPK, and the protein deacetylase sirtuin 1 enhancers like Cilostazol (Lee et al. [2015\)](#page-10-0). Besides, drugs that regulate transcriptional factor EB (TFEB) (Zhang and Zhao [2015](#page-11-0)), activate pro-autophagic enzymes, modify the maturation of autophagosome, or inhibit the enzymes antagonizing the formation of membrane structures (Cavieres et al. [2015,](#page-9-0) Papp et al. [2015\)](#page-10-0) may all have beneficial effects on AD patients. Emerging mTOR-independent autophagy enhancers have been screened, but the precise mechanisms of some enhancers remain unclear (Sarkar et al. [2007b\)](#page-11-0).

Combined Treatment May Have Additional Effects

Since AD is a multi-factor caused disease, the use of combination drug therapy may have more positive effects in AD treatment. Studies have shown that additive therapeutic effects and fewer side effects can be achieved at the same time through simultaneous stimulation of mTORdependent and independent routes. The promising effect of combination administration of lithium and rapamycin in Huntington's disease (HD) treatment may give us some clues. Lithium could activate autophagy through mTORindependent pathway while rapamycin activate autophagy through mTOR-dependent pathway. Combined administration of lithium and rapamycin results in stronger induction of autophagy, better protective effects on neuron survival, and more efficiency on autophagy substrates clearance such as mutant Huntingtin and α -synuclein compared with single medication alone (Sarkar et al. [2009](#page-11-0)). Similar effects were also observed in several cell lines from combination use of rapamycin with other mTOR-independent autophagy enhancers, such as trehalose, calpastatin, or SMERs (Sarkar et al. [2007a,](#page-10-0) [b,](#page-11-0) Williams et al. [2008](#page-11-0)). In addition to the better effect in autophagy induction, lower doses administration of each chemical reagent may be safer for longer treatment by minimizing dose-related side effects.

When to Deliver Drug is an Important Concern

When designing and evaluating autophagy interventional therapies, timing is always an important factor to be considered as the role of autophagy in different AD stages remains controversial. Early activation of autophagy is supposed to be beneficial, aiding neurons to clear abnormal protein aggregates and organelles. However, in the advanced phase, the activation of autophagy may show less benefit to alleviate AD pathology as more mature and stable aggregates cannot be eliminated efficiently through autophagy. Study has shown that giving rapamycin to 2-month-old 3xTg-AD mice throughout their lifespan induced autophagy and significantly reduced \overrightarrow{AB} plaques, Tau tangles, and subsequent cognitive deficits. However, inducing autophagy in 15-month-old 3xTg-AD mice which have established stable plaques, and tangles has no effect on AD-like pathology and cognitive deficits, suggesting autophagy induction exhibited beneficial effects only when they are administrated before the plaques and tangles formed (Spilman et al. [2010;](#page-11-0) Majumder et al. [2011](#page-10-0)). Ultimately, understanding the specific steps affected in the autophagy process in AD will be essential for the development of autophagy-based therapeutic drugs.

Gene Therapy for Autophagy Regulation

In addition to pharmacological chemicals, gene therapy is also a selective approach with the advantage of function in a tissue-directed manner. Several studies have shown gene therapy treatment designed to regulate autophagy via lentivirus or adenovirus-associated viral delivery had positive effects in diverse human disease including aging and neurodegenerative diseases. Although most gene therapy studies for AD-targeted NGF, recent work showed that oral vaccination with a recombinant $AAV/A\beta$ vaccine increased the clearance of $\mathbf{A}\beta$ from the brain and improved cognitive ability in AD animal models through enhanced autophagy (Wang et al. [2015a](#page-11-0), [b](#page-11-0)), suggesting modulating the autophagy pathway by viral delivery may be an important strategy for AD prevention and intervention.

BECN1 gene encoded protein beclin-1, which is a wellknown essential inducer of autophagy. Beclin-1 serves as a molecular platform assembling components which regulate the initiation of the autophagosome formation. Activation of Beclin-1 could up-regulate autophagy. In Becn $1^{+/-}$ transgenic mice, significantly reduced Beclin 1 expression lead to a decrease of neuronal autophagy and finally resulted in neurodegeneration (Pickford et al. [2008\)](#page-10-0). Consistent with this, the down-regulation of Beclin1 was also observed in the brains of AD patients especially in those brain regions which were most vulnerable to AD pathology (Pickford et al. [2008\)](#page-10-0). Administration of lentivirus encoding mouse beclin 1 to the frontal cortex and hippocampus of 6-month-old APP transgenic mice for 8 weeks resulted in prominent beclin 1 expression and significant reduced intracellular $\mathbf{A}\beta$ immunoreactivity (Pickford et al. [2008](#page-10-0)),

which suggested beclin 1 is a potential target when designing gene therapy for AD.

Up to now, few reports have been released considering enhancing autophagy through gene therapy. Continued investigations seem to hold tremendous potential for successful application of this novel approach.

Concluding Remarks and Future Perspectives

Although the etiology of AD remains unclear, and many factors including genetic mutation, environmental factors, imbalance of energy metabolism, and heavy metal ion (Yu et al. [2010;](#page-11-0) Li et al. [2010](#page-10-0)) seem to contribute to the etiology of AD, emerging advances regarding to autophagy have indicated its role as a protective factor in the early phase of AD, but an evil player in the late phase. Autophagy can influence the generation, secretion, and clearance of Ab, and it will also influence the phosphorylation status and clearance of tau. Thus, chemical modulators of autophagy as well as gene therapy targeting autophagyrelated proteins offer great potential for the treatment of AD. A number of mTOR-dependent and independent autophagy modulators have been demonstrated to have positive effects in AD animal models and patients. However, a more throughout understanding of autophagy malfunction in AD, as well as brain pharmacokinetics of autophagy modulators will be critical for designing new experiments with appropriate drug doses in any future clinical trials for AD.

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