

Invited Mini Review

Therapeutic implication of autophagy in neurodegenerative diseases

Md. Ataur Rahman¹ & Hyewhon Rhim^{1,2,*}

¹Center for Neuroscience, Korea Institute of Science and Technology, Seoul 02792, ²Department of Neuroscience, Korea University of Science and Technology, Daejeon 34113, Korea

Autophagy, a catabolic process necessary for the maintenance of intracellular homeostasis, has recently been the focus of numerous human diseases and conditions, such as aging, cancer, development, immunity, longevity, and neurodegeneration. However, the continued presence of autophagy is essential for cell survival and dysfunctional autophagy is thought to speed up the progression of neurodegeneration. The actual molecular mechanism behind the progression of dysfunctional autophagy is not yet fully understood. Emerging evidence suggests that basal autophagy is necessary for the removal of misfolded, aggregated proteins and damaged cellular organelles through lysosomal mediated degradation. Physiologically, neurodegenerative disorders are related to the accumulation of amyloid β peptide and α-synuclein protein aggregation, as seen in patients with Alzheimer's disease and Parkinson's disease, respectively. Even though autophagy could impact several facets of human biology and disease, it generally functions as a clearance for toxic proteins in the brain, which contributes novel insight into the pathophysiological understanding of neurodegenerative disorders. In particular, several studies demonstrate that natural compounds or small molecule autophagy enhancer stimuli are essential in the clearance of amyloid β and α -synuclein deposits. Therefore, this review briefly deliberates on the recent implications of autophagy in neurodegenerative disorder control, and emphasizes the opportunities and potential therapeutic application of applied autophagy. [BMB Reports 2017; 50(7): 345-354]

INTRODUCTION

Autophagy is a cellular process which can have a wide range

*Corresponding author. Tel: +82-2-958-5923; Fax: +82-2-958-6937; E-mail: hrhim@kist.re.kr

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of effects on pathological and physiological conditions, frequently leading to the removal of damaged/injured organelles and aggregated proteins (1). In more general terms, basal autophagy is important for removing misfolded proteins and damaged organelles, and therefore plays a crucial role in maintaining cellular homeostasis. Autophagy participates in a variety of normal physiological processes such as glucose homeostasis (2), lipid metabolism (3), aging (4) and many more. However, failure of autophagy causes cellular dysfunctions which render them unable to remove defective proteins or damaged organelles. Knockdown/inhibition of autophagy often causes premature aging in Caenorhabditis elegans (5). Interestingly, research in Drosophila melanogaster suggests that lifespan extends after enhanced autophagy (6). Several other studies have shown that deletion/mutation of the autophagy-related gene (Atg) provokes an imbalance of body homeostasis, resulting in obesity (7), insulin resistance (8), diabetes (9), and some neurodegenerative disease as well (10). In addition, neurodegenerative disease occurs when aggregated proteins are not accurately degraded by autophagy; for example, accumulation of aggregated forms of amyloid-β (Aβ) peptide (11) and tau protein (12) in Alzheimer's Disease (AD), α-synuclein in Parkinson's Disease (PD) (13), and that of huntingtin in Huntington's Disease (HD) (14). Therefore, autophagy pathways could be a potential therapeutic target for the treatment of neurodegenerative diseases. Indeed, neuronal cells are easily damaged through impaired autophagy among all types of cells in physiological systems (15). Understimulated autophagy has been correlated with difficulty to maintain synaptic activity (16) and axonal homeostasis (17). Several studies have managed to demonstrate a relationship between neurodegeneration and the autophagy signaling pathway (18). This review would like to emphasize the understanding of the basic molecular mechanisms and regulation of autophagy in neurodegenerative disease control.

MOLECULAR MECHANISMS OF AUTOPHAGY

Autophagy occurs through consecutive steps in the molecular biological pathway, such as sequestration, delivery of autophagosome to lysosomes (autolysosomes), degradation and utilization of toxic components, and collectively every step

demonstrates the diverse role of autophagy. Importantly, formation of phagophores are initiated during the early stages of autophagy, the production components of which are likely derived from numerous sources, including the endoplasmic reticulum (19), the endoplasmic reticulum-golgi intermediate compartment (20), the plasma membrane (21), recycling endosomes (22), the golgi complex (23), and lipid droplets (24). In fact, multiple autophagy-related genes are involved in membrane rearrangements and autophagosome formations that consequently become attached to the lysosomes. Generally, there are two ubiquitin-like conjugation reactions involved during autophagy. Firstly, there is the Atg12 conjugation to Atg5. Secondly, Atg16L1 binds to conjugate Atg12-Atg5 to form a complex which is essential for phagophore maturation (25). However, the second conjugation of Atg8 and LC3 is necessary to complete the autolysosome formation, in that the process of LC3 being cleaved through Atg4 to obtain cytosolic LC3-I. Finally, LC3-I is coupled to phosphatidylethanolamine (PE) during the macromolecular assembly of phagophores, which appears in the form of microtubule-associated protein light chain 3 (LC3-II) proteins (Fig. 1). It has been found that LC3-II localizes to the autophagosomal membrane, which is the most important marker of autophagy in mammal cells (26).

Whereas with several adaptor proteins, such as the neighbor of BRCA1 gene 1 (NBR1), nuclear dot protein 52 (NDP52), and p62/sequestosome 1 proteins are directly bind to LC3-II (27, 28). Subsequently, misfolded proteins (29) or abnormal cellular organelles (as well as mitochondria (30), peroxisome (31) and the endoplasmic reticulum (32) are engulfed by double-membrane autophagosomes. Strikingly, the autophagosomes are ultimately combined with lysosomes to form an autolysosome, where the internal substrates are degraded to produce nutrients and metabolites (33).

CLEARENCE OF AGGREGATE-PRONE PROTEINS THROUGH AUTOPHAGY IN NEURODEGENERATION

Recently, several studies revealed that intracellular protein aggregation and misfoldings are a common phenomenon of many neurodegenerative diseases, such as AD, PD, HD, and spinocerebellar ataxias (SCA) (34). To date, there are no efficient therapeutic approaches that can cure or prevent neurodegeneration diseases in humans. Thus, the need to understand the precise molecular mechanisms of neurodegeneration is urgent. To be studied are factors that regulate expression levels of protein misfolding and aggregate-prone proteins during the

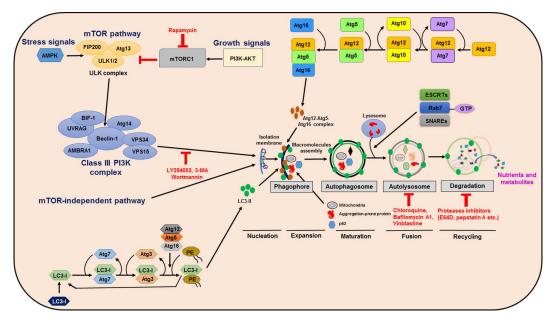


Fig. 1. Regulation of autophagy signaling pathway. Autophagy may initiate deprivation of nutrients or growth factors which activate AMPK and/or inhibition of mTORC1, leading to stimulation of ULK complex (FIP200 and ATG13). Beclin-1 become phosphorylated, leading to VPS34 activation and then initiation of phagophore formation. VPS34 complex function comprises a regulatory subunit like VPS15 (p150) and Beclin-1, their connection being with other regulatory factors e.g. AMBRA, ATG14, UVRAG, and BIF-1. Atg5-Atg12 conjugation involves Atg7 and Atg10 to form a complex, Atg12-atg5-Atg16 influences the formation of phagophores. Atg5 and Atg12 forms a complex with Atg16, which acts like an E3-function towards LC3-PE assembly (LC3-II). This has a role in the initiation of phagophore formation. LC3-II is a particular autophagy indicator marker which is eventually disrupted by autolysosomes. Autophagosome maturation also involves fusion with lysosomes which are mediated by Rab7, ESCRT and SNARE proteins, eventually leading to cargo degradation and recycling of nutrients and metabolites.

process of neurodegeneration. Usually, there are many methods to treat neurodegenerative diseases that might enhance degradation of prone-proteins. Among the most possible routes of misfolded protein clearance are the autophagy-lysosome and ubiquitin-proteasome pathways (Fig. 2). Although the ubiquitin proteasome system primarily degrades small-molecule proteins, its mechanism is not clear and it is unknown whether it is a possible therapeutic target for the clearance of aggregate-prone proteins. In addition, autophagic degradation of aggregate-prone proteins are related to decreasing protein aggregation and toxicity. Therefore, enhancing autophagy may be a promising therapeutic approach for neurodegenerative diseases, in that aggregate-prone proteins are used as autophagy substrates, such as tau (35), α-synuclein (36), mutant huntingtin (14), and mutant ataxin 3 (35), for their clearance of toxic products and having an effect on neurodegeneration control.

Autophagy in Alzheimer's disease

AD is a multifactorial, irreversible, progressive and the most common type of dementia in elderly people worldwide. It is characterized by the gradual diminishment of memory and thoughts, and ultimately the capability to carry out any cognitive function (37). Recently it has been shown that the main pathological feather of AD is comprised of intracellular neurofibrillary tangles containing hyperphosphorylated tau proteins, self-aggregating extracellular Aβ plaques, the weakening of cholinergic function, and eventually autophagy dysfunction (38, 39). However, the AB peptide derived from the successive splitting of amyloid precursor protein (APP) through the β -site APP cleavage enzyme 1 (BACE1) and γ -secretase also has been linked to neurodegeneration. Accumulation of this peptide has been shown to cause cognitive deficits and neuronal dysfunction in the cortex and hippocampus during AD pathogenesis (40). Particularly, Aβ may also be produced in autophagosomes, which seem to comprise APP and presenilin-1 enzymes that participate in the cleavage of APP to AB (41). Additionally, autophagy conditions may play an important role in AB secretion into extracellular space, where it contributes to the accumulation of plaques. Numerous studies shows that deletion of Atg7 in transgenic mice models of APP results in fewer AB extracellular secretions as well as plaque formations; therefore

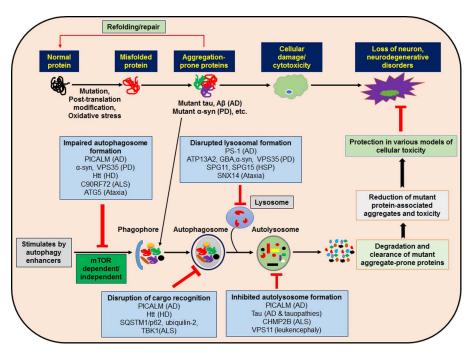


Fig. 2. Modulation of autophagy and intersections between autophagy and disease-associated genes as a therapeutic approach for neurodegenerative disorders. Impaired autophagy is involved in the pathogeneses of several neurodevelopmental and neurodegenerative disorders. Different cytosolic mutant and aggregate proteins are linked with neurodegenerative symptoms, which may predominantly be removed through autophagy. Pharmacological stimulation of small molecule autophagy enhancers may initiate autophagy via mTOR dependent/independent pathways that reduce mutant protein-associated aggregates and toxicity, leading to cytoprotection and rescue against neurodegeneration. Several genes connected with neurodegenerative disorders have now been associated with autophagy function. These genes perform a number of diverse steps throughout the autophagic cycle, from early steps of autophagosome development through autolysosome formation. Their proposed sites of action are now designated, along with the neurodegenerative disease with which they are associated.

autophagy up–regulation would simply lead to degraded $A\beta$ secretion (42). Furthermore, other research has suggested that autophagy is involved in the removal of $A\beta$ under physiological states, suggesting that maintenance of $A\beta$ homeostasis in a healthy brain is essential (43). In correlation with these findings, it has been indicated that autophagy activation is decreased in AD patient brains and animal models (44). More importantly, the direct link between autophagy reduction activity and promotion of accumulation of $A\beta$ in the brain consequently demonstrates their contribution to AD pathogenesis (45). Therefore, targeting $A\beta$ and tau protein should be considered as an important therapeutic approach against AD pathogenesis via upregulation of the autophagy process.

Perhaps most importantly, class III PI3K/beclin-1 signaling has been found to be a constructive and significant regulator in autophagy (46). For that reason, it seems that modulation of autophagy represents a feasible method for Aβ clearance in the brain (45). Moreover, substantial efforts have been made to recognize safe and effective pharmacological stimulation of autophagy for AD treatment (47). In practical terms, the existing outcomes of ongoing research have provided biochemical, epidemiological, neuropathological and genetic evidence to further motivate the design of viably therapeutic approaches against AD. Incidentally, most anti-AD research groups are now embracing the discovery of BACE1, a rate-limiting enzyme in Aβ generation, which acts as an enzymatic inhibitor (48). To understand the molecular basis of AD in autophagy, recent studies have been shown that elevated protein levels of BACE1 found in AD patients, as well as mouse models, suggest the down-regulation of BACE1 expression as a novel target for AD treatment (49, 50). Even though this approach may offer some improvement, an effective therapy to halt the development of neural damage in AD patients is still missing. Meanwhile, medications for AD have already been approved by the FDA, such as an NMDA receptor antagonist and four other acetylcholinesterase (AChE) inhibitors (51). It is possible that combined therapy with NMDA receptor antagonists along with AChE inhibitors could facilitate greater treatment efficacy. Otherwise, multifunctional agents are also being considered as possible drug candidates for AD (52). For that reason, research for multi-target compounds has been constantly growing for the treatment of AD (53).

For treatment purposes, autophagy enhancers treated in various AD mouse models have produced useful evidence related to a decrease in A β levels. For example, carbamazepine, an mTOR-independent autophagy stimulator, improves memory deficits by enhancing autophagy via decreasing A β formation in APP/PS1 double-transgenic mice (54). Notably, A β plaque formation diminishes by rapamycin in triple-transgenic PS1/APP/Tau mice and improves cognitive defection through inducing autophagy (55). Lithium treatment also suppresses tau protein in aged APP/PS1 mutant mice pathology via GSK3 β inhibition, found to disturb tau phosphorylation (56).

Interestingly, latrepirdine, an antihistamine drug, shows to increase autophagy by reducing the A β peptide through inactivating the mTOR complex 1, which has been found to recover cognition function in an AD mice models (57). Correspondingly, trehalose, a disaccharide sugar, reduces aggregation of insoluble tau, which enhances neuronal brainstem survival in the cerebral cortex via influences of autophagy (58). Moreover, rapamycin and lithium have been demonstrated to decrease the aggregation of p62 in cortical and hyperphosphorylated tau proteins (59, 60). Taken together, all of this evidence indicates that autophagy stimulation is a promising therapeutic approach for "tauopathies," as well as for A β clearance in AD mouse models.

Autophagy in Parkinson's disease

PD is a progressively degenerative disorder of the nervous system and the second most common neurodegeneration in humans. PD is a long-term degenerative brain disorder that mostly affects motor function of the central nervous system, due to the progressive degeneration of dopamine neurons in an area of the brain called the substantia nigra, where clumps of α -synuclein protein known as Lewy Bodies are formed (61). Dementia becomes more common in advanced phases of the disease, with depression and anxiety generally occurring in more than one third of people with PD (62). PD is also related with A53T and A30P point mutations of α-synuclein or leucine-rich repeat kinase 2 mutations, and sometimes other gene multiplication. PTEN-induced kinase 1 or Protein deglycase mutations in parkin have also c been associated with recessive, early-onset PD (63). The pathological feature of early-onset PD is Lewy body formation via α-synuclein aggregation (64). It has been found that α-synuclein point mutations are sometimes autophagy substrates and their excessive expression does not distress autophagy (65). For that reason, autophagy influences by lithium, trehalose, carbamazepine, calpastatin, and rilmenidine promote the removal of mutant α -synuclein (66). In contrast, wild-type α -synuclein overexpression reduces autophagosome synthesis in transgenic mice and mammalian cells through down regulation of Ras-related protein Rab-1A (65). Additionally, overexpression of α-synuclein has been proposed to prevent autophagy in the rat midbrain by repossessing transcription factor EB into combinations, thereby affecting cytoplasmic retention. Conversely, transcription factor EB overexpression activates autophagy and prevents neurotoxicity of α-synuclein-induction (67). Hence, autophagy activator protein beclin-1 ameliorates neuronal pathology in lentiviral-overexpression transgenic mice models through a decrease in α -synuclein accumulation (36). Even though the effectiveness of autophagy stimulation in vivo attempts further warrant consideration, absence of strong neurodegeneration phenotypes, together with dopaminergic neuronal susceptibility in numerous transgenic PD models, have prevented treatment trials in PD (68).

In the preclinical approach to PD, autophagy's significance

has been increased by the promising effects of several potential autophagy enhancers, such as rapamycin, lithium, metformin and trehalose - routinely used in laboratory settings to increase clearance of α-synuclein - thereby protecting neuronal cells (69). To date, the most comprehensively-tested autophagy-enhancer is rapamycin, isolated from bacterial strain of Streptomyces hygroscopicus. In particular, rapamycin blocks the upstream signaling factor mTOR by binding to its intracellular small protein FK506 binding protein 12 receptor, thus disrupting mTOR function (70). Rapamycin also has the capability to decrease α-synuclein accumulation and block α-synuclein induction of neurodegeneration in wild-type and A53T α -synuclein expressing PC12 cells (71), as well as α synuclein overexpressing mice (72) and rats (67). In addition, rapamycin decreases neuronal cell death induce via 6-OHDA (73), rotenone (74) and MPTP mouse models of PD (74). Furthermore, rapamycin exclusively improves motor function in A53T α-synuclein overexpressing mice (75) and also shows neuroprotective activity in Drosophila melanogaster mutated PTEN-induced putative kinase-1 as well as parkin (76); although it has been noted to accompany a widespread number of side-effects (77). By comparison, another preclinical trial drug, lithium, induces autophagy in an mTOR-independent manner and has a role in in vivo as well as in vitro models of neuroprotection (78). In preclinical models of PD, lithium has been shown to inhibit aggregation of α -synuclein in PC12 cells expressing A53T and A30P α-synuclein (79). In addition, lithium has also been demonstrated to prevent rotenoneinduction neurotoxicity and cell death through the stimulation of autophagy (80). When taken together, these preclinical studies via rapamycin in addition with lithium shows valuable effects on diverse models of PD-related pathological progressions, such as α -synuclein aggregation, which is associated with autophagy stimulation. These pioneering studies have demonstrated that the prospective role of autophagy-enhancing approaches in the setting of experimental drugs and medical treatments lends credence to the furtherance of research in PD-related pathology.

Autophagy in other neurodegenerative diseases

Huntington disease (HD), an autosomal dominant mutation, is a progressive neuronal genetic disorder that impacts muscle coordination and manifests in the formation of cognitive decline and dementia. Symptoms include unsteady movements, lowering of mental and emotional abilities, and finally psychiatric and behavioral problems (81). Pathologically speaking, the polyglutamine disorder group of neurodegenerative diseases are caused by the expansion of cytosine-adenineguanine repeats in DNA encoding, therefore rendering the protein aggregation-prone and pathogenic (82). For treatment of HD with CCI-779, a rapamycin analog drug improved pathology in mice models of HD by diminishing the aggregation of mutant huntingtin protein via stimulation of the autophagy pathway (14). Interestingly, numerous mTOR-

independent autophagy activators, for example Ca2+ channel blockers (loperamide, verapamil, amiodarone), inositol-lowering agents (lithium), calpain inhibitor (calpastatin), and NOS inhibitor (L-NAME), improved the removal of huntingtin as well as lowered toxicity in cellular HD models (66). Recently, a natural compound, the quaternary ammonium salt berberine (an isoquinoline alkaloid), has been found to prevent mutant Htt protein accumulation, alleviate HD symptoms, as well as activate autophagy in HD cell and mouse models (83). Another natural product, onjisaponin B (derivative of Radix Polygalae), has been found to control autophagy in PC12 cells through Atg7 as well as the AMPK-mTOR signaling pathway to enhance clearance of the Htt mutant protein and α -synuclein mutant (84). Therefore, it has been suggested that modulation of autophagy in HD might be a favorable approach to the treatment of this neurodegeneration.

Spinocerebellar ataxia (SCA), a progressive, genetic neuronal disease, is described by means of gradually growing problems associated with muscle stiffness, speech and swallowing difficulties, which also leads to involuntary eye movements. In addition, SCA is caused by the aggregation of mutant ataxin-3, thereby impairing learning and cognition function (85). Interestingly, modulation of autophagy is also involved in the removal of the abnormal ataxin-3 protein, and impaired autophagy may affect the aggregation of the mutant ataxin-3 protein that initiates cellular toxicity as well as cell death. Accordingly, targeting SCA3 through directing autophagy could become a novel therapeutic strategy (86). Likewise, trehalose, a natural alpha-linked disaccharide, is an mTORindependent autophagy inducer that could be used to control the ataxin-3 protein in SCA. Recently, it has also been found that trehalose analogs - such as lactulose and melibiosesignificantly decreased aggregation of abnormal ataxin-3 protein in a cell model, which was also associated with the stimulation of autophagy, along with a reduction in free radical production (87). Therefore, autophagy induction could potentially be used as a therapeutic strategy to control SCA.

Amyotrophic lateral sclerosis (ALS), a rare group of neurological disorders, is the effect of gradual degeneration and death of nerve cells; mainly motor neurons which control voluntary muscle function, resulting in difficulty in speaking, walking, swallowing, chewing, and finally breathing (88). To control this disease, it is necessary to induce autophagosomes through a decreasing mTOR complex 1, using a Cu-Zn superoxide dismutase mutant mice model (89). It is suspected that p62 could bind to superoxide dismutase mutations in an ubiquitin-independent manner to target its removal by autophagy (90). Conversely, mTOR-independent autophagy stimulation by lithium eases removal of mutant superoxide dismutase and hinders the disease from the start. However, treatment with trehalose has been found to be more beneficial in superoxide dismutase mutant mice through upregulation of mTOR-independent autophagy, as is found in ALS (91). Therefore, autophagy stimulation is very essential issue for in

vivo studies of ALS treatment.

THERAPEUTIC IMPLICATIONS OF AUTOPHAGY IN NEURODEGENERATION

For therapeutic purposes, it has been suggested that upregulation of autophagy through the mTOR complex 1-mediated pathway might help target the removal of aggregate protein molecules, as has been suggested by a decrease in cytotoxicity of mouse, zebrafish, and Drosophila studies as well (92). Tauopathies, α -synucleinopathies, SC 3, HD, and familial prion disease models, have all been implicated in the successful treatment of neurodegenerative disease through this strategy of stimulated autophagy. In particular, mTOR-independent autophagy inducers, rapamycin analogues such as rilmenidine and trehalose, haves been used in the treatment of those diseases (92). On the other hand, autophagy inhibitor activity increases the toxicity of proteins that lead to enhancement of neurodegeneration (92). Recently, it has been shown that the phosphatidylinositol-binding clathrin assembly protein, also known as PICALM, acts equally in the autophagosome formation and autophagosome degradation process, in AD genome-wide association studies (GWAS). Hence, altered PICALM activity in vitro as well as in vivo leads to the increase in toxicity and accumulation of tau in AD pathogenesis (93). Moreover, pharmacological inhibition of mTOR complex 1 is a demonstrated target in autophagy control in neurodegeneration (94). This also implies that rapamycin and its chemically synthesized analogues such as CCI-779, are widely used as potential activators of autophagy in yeast and mammalian cells in neurons, as well as in vivo in the mouse brain. Eventually, widespread preclinical animal model studies are required to explore the use of autophagy in the prevention of neurodegenerative disease.

Recent observations have found that inhibition or reduction of histone deacetylase 6 helps to rescue memory in 5XFAD AD mouse models and AD patients, which implies further therapeutic possibilities for AD, as well as other neurodegenerative diseases (95). In addition, it has been found that transplanted protein-induced pluripotent stem cells (iPSCs), derived from skin of mouse fibroblasts, differentiate into glial cells, resulting in the reduction of plaque depositions shown to alleviate cognitive dysfunction in 5XFAD transgenic AD mouse models. This suggests a favorable therapeutic application for AD (96). Similarly, insulin-degrading enzymes and major proteases of Aβ secreted from astrocytes have been shown to potentially alleviate AD symptoms through alternative, autophagy-based secretory pathways (97). Furthermore, statin, a class of lipid-lowering medications, has been demonstrated to induce autophagy in astrocyte cultures through AMPK-mTOR mediated pathways, and as it has been suggested that autophagy is essential in insulin-degrading enzyme secretion, modulation of autophagy could provide a possible therapeutic approach in Aß pathology by increasing clearance

of extracellular AB (98). Hence, accumulation of AB peptide participates in the pathological condition of AD, while inhibiting Aβ production or increasing Aβ removal may be implicated in slowing the progression of AD (40). In particular, the promotion of A β clearance is currently considered to be an additional therapeutic approach for AD. Thereby, autophagy has been found to have an important role in the clearance of Aβ under normal physiological conditions; for that reason it is essential to maintain $A\beta$ homeostasis in a healthy brain (43). Most importantly, our current research is considerably directed to the identification of safe and more effective pharmacological inducers of autophagy in neurodegenerative diseases. Therefore, to target AD drug development, natural compoundmediated enhancement of autophagy might represent a sustainable strategy to AB clearance. To assess this question, alternatively a natural compound could be a potential drug candidate which facilities the enhancement of autophagy in AD patients. It has recently been found that several alkaloids, such as oxoisoaporphine, significantly decreased Aβ secretion in neuroblastoma cells when overexpressed in human APPsw Swedish mutants (99). Another recent report suggests that chronic administration of the isoquinoline alkaloid berberine significantly decreases Aβ deposits while promoting Aβ clearance. As a result, this was shown to improve cognitive impairments in 3XTg AD mice by enhancing activity of autophagy via the class III PI3K/beclin-1 pathway (100).

For the treatment of PD, several natural compounds have recently gained attention due to their ability to clear α-synuclein in animal and cell models. Specifically, studies have indicated that a natural compound, curcumin, derived from the curry spice of turmeric shows low toxicity in normal cells, and also significantly decreases the accumulation of α-synuclein in A53T cell via downregulation of the mTOR signaling via increasing autophagy (101). Interestingly, oxyresveratrol, a natural antioxidant, increases mTOR dependent autophagy in human neuroblastoma cell models independent of apoptosis (102). In addition, 18α -glycyrrhetinic acid, well known a gap-junction inhibitor, induces autophagy through upregulation of beclin-1, Atg5, Atg7 and LC3, as well as downregulation of p62 in neuroblastoma cells (103). Therefore, these reviews attempt to expound upon the important observation that the pharmacological targeting of autophagyenhancing natural compounds hold a favorable implication for the development of PD therapies.

CONCLUSION

Even though a variety of autophagy-related proteins have been demonstrated to participate and control endogenous autophagy pathways (104), several studies have been performed to explore autophagy regulation through use of the active compounds of plants. Although numerous fundamental queries are essential for further address, many novel agents could one day be useful in a clinical approaches; thus the interest of

research in autophagy is developing rapidly, and clinical applications might be anticipated in the near future. Furthermore, it is now important to characterize dysfunctional autophagy in diverse stages of genetic and molecular subtypes in order to better understand their roles in neurodegeneration. It is also necessary to produce clinical results demonstrating the efficacy of downstream autophagy regulation, which would help usher in an exciting new era for the development of neurodegenerative therapeutic strategies. Consequently, additional studies are required on the physiological roles of modulation of brain autophagy processes in neurodegenerative disease. Finally, we would like to screen novel, natural compounds that modulate autophagy and identify main therapeutic targets, such as the key molecular mechanisms that underlie the pathophysiological roles of neurodegeneration, with our main concern being the discovery of potential therapeutic drug targets.

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CONFLICTS OF INTEREST

The authors have no conflicting financial interests.

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