

EGCG-mediated autophagy flux has a neuroprotection effect via a class III histone deacetylase in primary neuron cells

Ju-Hee Lee^{1,2}, Ji-Hong Moon^{1,2}, Sung-Wook Kim^{1,2}, Jae-Kyo Jeong^{1,2}, Uddin MD. Nazim^{1,2}, You-Jin Lee^{1,2}, Jae-Won Seol¹ and Sang-Youel Park^{1,2}

¹ Biosafety Research Institute, College of Veterinary Medicine, Chonbuk National University, Jeonju, Jeonbuk, South Korea

² Department of Bioactive Material Sciences and Research Center of Bioactive Materials, Chonbuk National University, Jeonju, Jeonbuk, South Korea

Correspondence to: Jae-Won Seol, email: jwsseol@chonbuk.ac.kr

Sang-Youel Park, email: sypark@chonbuk.ac.kr

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ABSTRACT

Prion diseases caused by aggregated misfolded prion protein (PrP) are transmissible neurodegenerative disorders that occur in both humans and animals. Epigallocatechin-3-gallate (EGCG) has preventive effects on prion disease; however, the mechanisms related to preventing prion diseases are unclear. We investigated whether EGCG, the main polyphenol in green tea, prevents neuron cell damage induced by the human prion protein. We also studied the neuroprotective mechanisms and proper signals mediated by EGCG. The results showed that EGCG protects the neuronal cells against human prion protein-induced damage through inhibiting Bax and cytochrome c translocation and autophagic pathways by increasing LC3-II and reducing and blocking p62 by using ATG5 small interfering (si) RNA and autophagy inhibitors. We further demonstrated that the neuroprotective effects of EGCG were exhibited by a class III histone deacetylase; sirt1 activation and the neuroprotective effects attenuated by sirt1 inactivation using sirt1 siRNA and sirtinol. We demonstrated that EGCG activated the autophagic pathways by inducing sirt1, and had protective effects against human prion protein-induced neuronal cell toxicity. These results suggest that EGCG may be a therapeutic agent for treatment of neurodegenerative disorders including prion diseases.

INTRODUCTION

Transmissible spongiform encephalopathies or prion diseases are neurodegenerative disorders that include bovine spongiform encephalopathy in cattle, chronic wasting disease in elk and deer, endemic scrapie of sheep and goats, and Creutzfeldt-Jacob disease and Gerstmann-Straussler-Scheinker syndrome in humans [1, 2]. Human prion diseases can occur fortuitously or by infection caused by mutation of the cellular prion protein (PrP^c) [3]. An important step in the disease pathophysiology is conversion of cellular PrP^c to disease-associated misfolded conformers or PrP^{Sc} [4]. These PrP^{Sc} are mainly composed of prion infectivity and are detectable in damaged brain and lympho-reticular organs [5].

The amino acid residue 106-126 of the cellular

prion protein exhibits PrP^{Sc} characteristics including neuronal toxicity. This leads to fibril accumulation and mitochondrial damage, formation of amyloid-like fibrils *in vitro*, and proteinase K resistance [6, 7]. PrP (106-126) expresses significant toxicity to cultured cells, including primary rat hippocampus neuron cells, primary retinal cells, and PC12 cells [8, 9]. Furthermore, PrP (106-126) induces the accumulation of cytosolic PrP, which leads to enriched PrP in cell membranes [10].

Epigallocatechin-3-gallate (EGCG) is a natural component of green tea representing approximately 43% of its total phenol content [11]. EGCG has attracted attention as a potential therapeutic target in different areas of study, including neuronal diseases [12, 13], cardiovascular risk factors [14, 15], cancer [16], obesity, and type 2 diabetes [17, 18]. EGCG has protective effects

against neuronal damage in diseases such as Alzheimer's disease (AD) and Parkinson's disease by blocking the formation of reactive oxygen species (ROS) and through iron chelating and anti-apoptotic functions in neurons [19-21].

Autophagy is an intracellular lysosomal process in which a double membrane vesicle (autophagosome) forms and sequesters the intracellular organelles and part of the cytoplasm. The autophagosome then fuses with a lysosome to form the autolysosome, exposing the inner compartment to lysosomal hydrolases for bulk degradation [22]. The formation of the autophagosome is mediated by the Atg12-Atg5-Atg16 complex and microtubule-associated protein light chain 3 (LC3-I)-phospholipid conjugates (LC3-II) [23, 24]. P62/SQSTM1 is well known autophagy substrate; p62 is incorporated into autophagosomes by interacting with LC3 and is efficiently degraded by autophagy. Inhibiting autophagy results in rapid accumulation of cellular p62, whereas decreased p62 levels are associated with activating autophagy; thus, p62 is employed as an indicator of autophagy flux [25].

Silent information regulator two ortholog 1 (sirt1) is a member of the sirtuin family and is a nicotinamide adenine dinucleotide-dependent histone deacetylase [26, 27]. It is the human ortholog of the yeast sir2 protein and is most likely one of the key proteins regulating caloric restriction-mediated lifespan extension [28]. Sirt1 removes acetyl groups from a variety of substrates such as the tumor suppressor protein p53, members of the FOXO family (forkhead box factors regulated by insulin/Akt), and hairy and enhancer of split 1. It is also related to physiological functions such as metabolism, cell survival, and aging [29, 30]. Sirt1 has been linked to neurodegenerative diseases such as Huntington's disease, AD, and Parkinson's disease neuropathology [29, 31]. sirt1 stimulates peroxisome proliferator-activated receptor gamma coactivator (PGC-1 α) signaling [32] and AMPK activity by modulating deacetylation, activating LKB1 [33], and controlling the interaction between MnSOD and FoxO3a [34]. All of these key target proteins activated by sirt1 are essential for attenuating mitochondrial damage and oxidative stress [32, 35-37]. Furthermore, sirt1 protects against neurodegenerative diseases such as prion diseases by regulating mitochondrial homeostasis [38].

In some reports, EGCG stimulates the activity of the sirt1 regulatory protein [39]. Under stabilizing conditions, deacetylation of sirt1 is enhanced by diverse polyphenols such as EGCG, epicatechingallate, and myricetin (3.19-fold) [40]. A previous report showed that EGCG is an antagonist that suppresses oxidative stress induced by 1-methyl-4-phenyl-pyridine (MPP⁺) by modulating the sirt1/PGC-1 α signaling pathway in PC12 cells [41]. Protective mechanisms of EGCG against cell damage by MPP⁺ and Cd²⁺-induced mitochondrial damage and neurodegenerative diseases such as Parkinson's disease have been suggested [41, 42]. The

role of EGCG as an activator of autophagy in various cells such as macrophages [43], vascular endothelial cells [44], and hepatocytes [45] has been reported. In many cells, EGCG has health-associated benefits by activating autophagy against various diseases, including cardiovascular disorders, cancer, and atherosclerosis disorders. Although some reports have demonstrated that EGCG induces autophagy [44] or the sirt1 pathway [41] respectively, EGCG controlling autophagy via sirt1 has not been reported. Thus, this present study focused on the induction of autophagy and activation of the sirt1 pathway by EGCG on PrP (106-126)-induced neurotoxicity and mitochondrial damage in SH-SY5Y cells. The data support the hypothesis that EGCG-mediated autophagy via sirt1 expression and activation may be a therapeutic implication for prion disease and mitochondrial damage caused by prions.

RESULTS

EGCG has neuroprotective effects against PrP (106-126)-induced neurotoxicity

Many studies have shown that EGCG limits brain inflammation and reduces neuronal damage [19]. In this study, the neuroprotective effect of EGCG on PrP (106-126)-induced neuronal cell death (Figure 1) was confirmed. The primary neuron cells were pretreated with EGCG at concentration of 10 μ M for 1 hr and then exposed to 50 μ M PrP (106-126) for 36 hr. The primary neuron cells were measured for cell viability using Annexin V assay (Figure 1A, 1B). Without EGCG, only PrP (106-126)-treated cells exhibits cytotoxicity, and EGCG-treated cells appeared increased cell viability. To evaluate whether EGCG prevents apoptotic signaling, we confirmed the cleaved caspases 3 protein levels. Increased cleaved caspases 3 by treating PrP (106-126) was restored similarly to control group.

SH-SY5Y cells were pre-incubated with EGCG at concentrations of 1 and 10 μ M for 1 hr and then exposed to 50 μ M PrP (106-126) for 36 hr. When neuronal cells were treated with PrP (106-126) without EGCG, it resulted in neurotoxicity as measured by Annexin V (Figure 1D, 1E). Treatment with EGCG prevented PrP (106-126)-induced apoptosis in a dose-dependent manner. We also measured cell viability against scrambled PrP (106-126) as a control condition for PrP 106-126. As a result, cell death caused by scrambled PrP (106-126) did not occur. LDH release levels showed that EGCG inhibited PrP (106-126)-induced apoptosis in a dose-dependent manner but scrambled PrP did not lead to cell death (Figure 1F), consistent with previous results. Cleaved caspase 3 increased by PrP (106-126) was attenuated by EGCG treatment in a dose-dependent manner (Figure 1G).

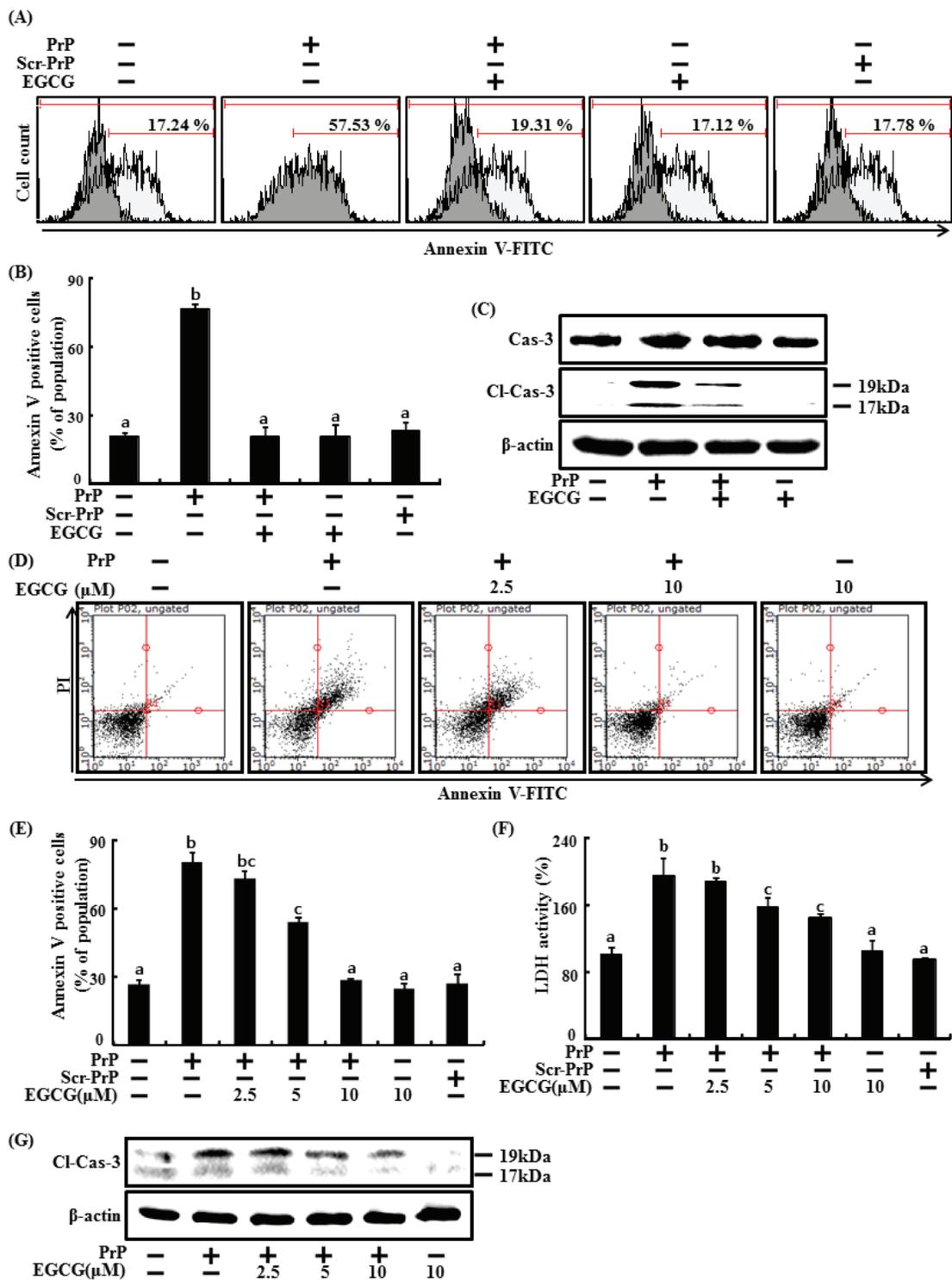


Figure 1: EGCG prevents neuronal cells from PrP (106-126)-induced cell death. The primary neuron cells were pretreated with 10 μ M of EGCG for 1 hr and then exposed to 50 μ M PrP (106-126) or scrambled PrP (106-126) (scr-PrP) for 36 hr. Cell viability was measured by the Annexin V assay A, B. Western blot for cleaved caspases 3 protein was conducted with primary neuron cells. Beta-actin was used as the loading control C. SH-SY5Y cells were pretreated with 2.5, 5, or 10 μ M of EGCG for 1 hr and then exposed to 50 μ M PrP (106-126) or scrambled PrP (106-126) (scr-PrP) for 36 hr. Cell viability was measured by the Annexin V assay D, E. Lactate dehydrogenase (LDH) assay was used to quantify LDH released into the medium F. Western blot for cleaved caspases 3 protein was conducted with SH-SY5Y cells. Beta-actin was used as the loading control G. Bars indicate mean \pm standard error ($n = 4$). The data were analyzed using ANOVA and Tukey multiple range tests ($P < 0.01$). Means sharing a common alphabetical symbol did not significantly differ. Cl-Cas-3, Cleaved caspase-3.

EGCG inhibits the PrP (106-126)-induced mitochondrial apoptotic pathway

Mitochondrial dysfunction is related to neuronal cell death caused by prions [46]. This study also examined whether an increase in Sirt1 activity causes prion-mediated mitochondrial impairment. Treatment of 50 μ M PrP (106-126) increased green fluorescence indicating a lower MTP, whereas decreased PrP (106-126)-induced green fluorescence in cells pre-treated with EGCG (10 μ M) (Figure 2A). These data were reconfirmed by measuring JC-1 using fluoroscopy (Figure 2B), PrP (106-126) treatment induced green fluorescence indicating a lower MTP, and PrP (106-126) treatment changed red fluorescence indicating a normal MTP following EGCG pretreatment (Figure 2B and 2C). In the same manner, PrP (106-126)-treated cells led to Bax translocation in the mitochondria, which increased cytochrome *c* release into the cytosol, whereas PrP (106-126)-induced

Bax translocation and cytochrome *c* release decreased following EGCG treatment in SH-SY5Y cells. EGCG prevented PrP (106-126)-induced Bax translocation and cytochrome *c* release (Figure 2D and 2E). These data suggest that EGCG treatment inhibits the PrP (106-126)-mediated mitochondrial apoptotic pathway.

EGCG inhibits the PrP (106-126)-induced apoptotic pathway by autophagy

We used autophagy as a survival strategy for prion-mediated neurotoxicity.

First, we examined whether EGCG induces autophagy in primary neuron cells. LC3-II and p62 expression levels were detected using a specific antibody in a Western blot analysis. The primary neuron cells were treated with EGCG at concentrations of 1, 2.5, 5, or 10 μ M for 24hr. EGCG increased LC3-II expression levels but decreased p62 levels (Figure 3A)

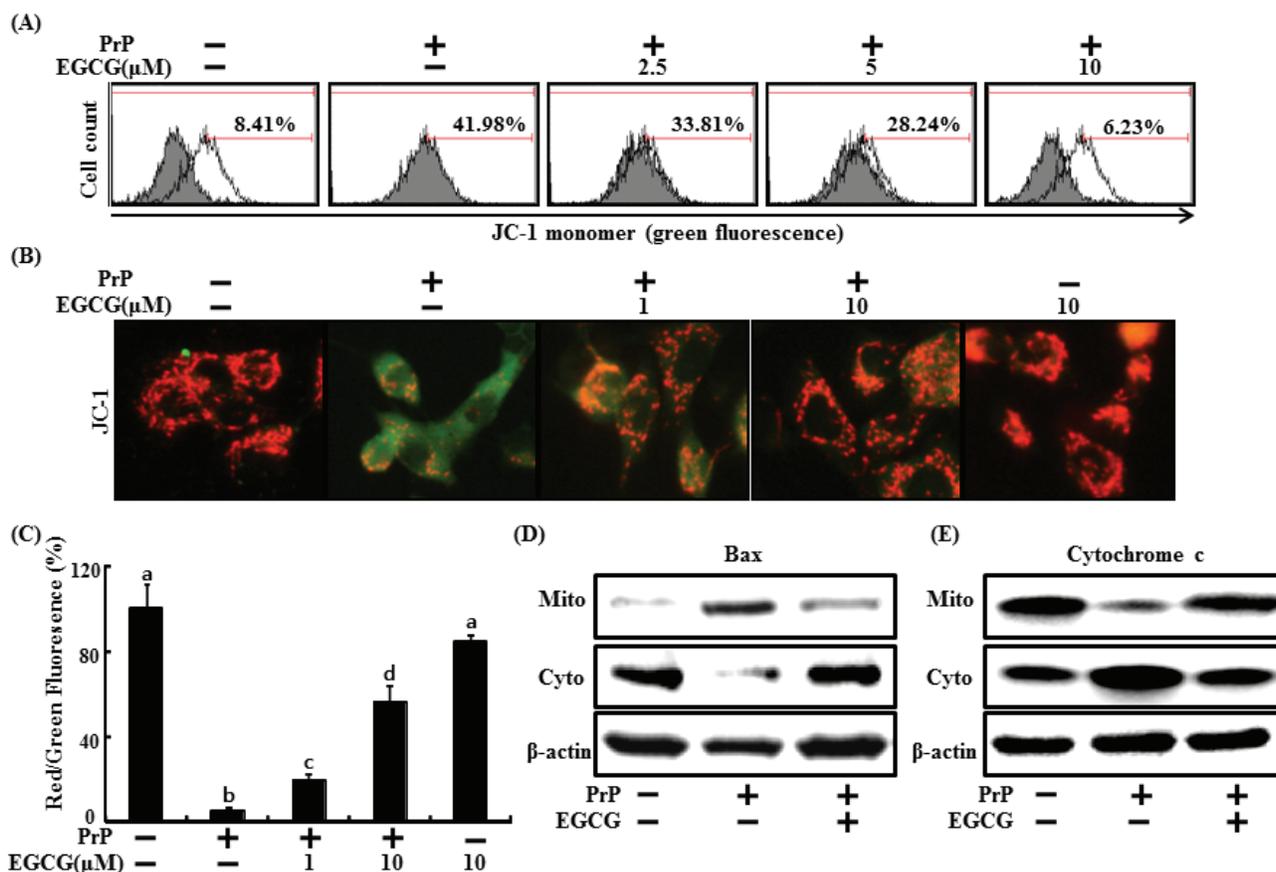


Figure 2: EGCG protects PrP (106-126)-induced mitochondrial damage. SH-SY5Y cells were pretreated with 2.5, 5, or 10 μ M of EGCG for 1 hr and then exposed to 50 μ M PrP (106-126) for 36 hr. The treated cells were measured for JC-1 mono form (green) by flow cytometry. Percent values in the histogram represent the population of JC-1 monomeric cells. **A.** The treated cells were measured for JC-1 aggregates form (red) and mono form (green) by confocal microscopy analysis. Scale bar = 50 μ m **B.** Separation of the cytosol and mitochondrial extracts was analyzed by Western blot using antibodies, to cytochrome *c* and Bax protein **D, E.** Bars indicate mean \pm standard error ($n = 4$). The data were analyzed using ANOVA and Tukey multiple range tests ($P < 0.01$). Means sharing a common alphabetical symbol did not significantly differ.

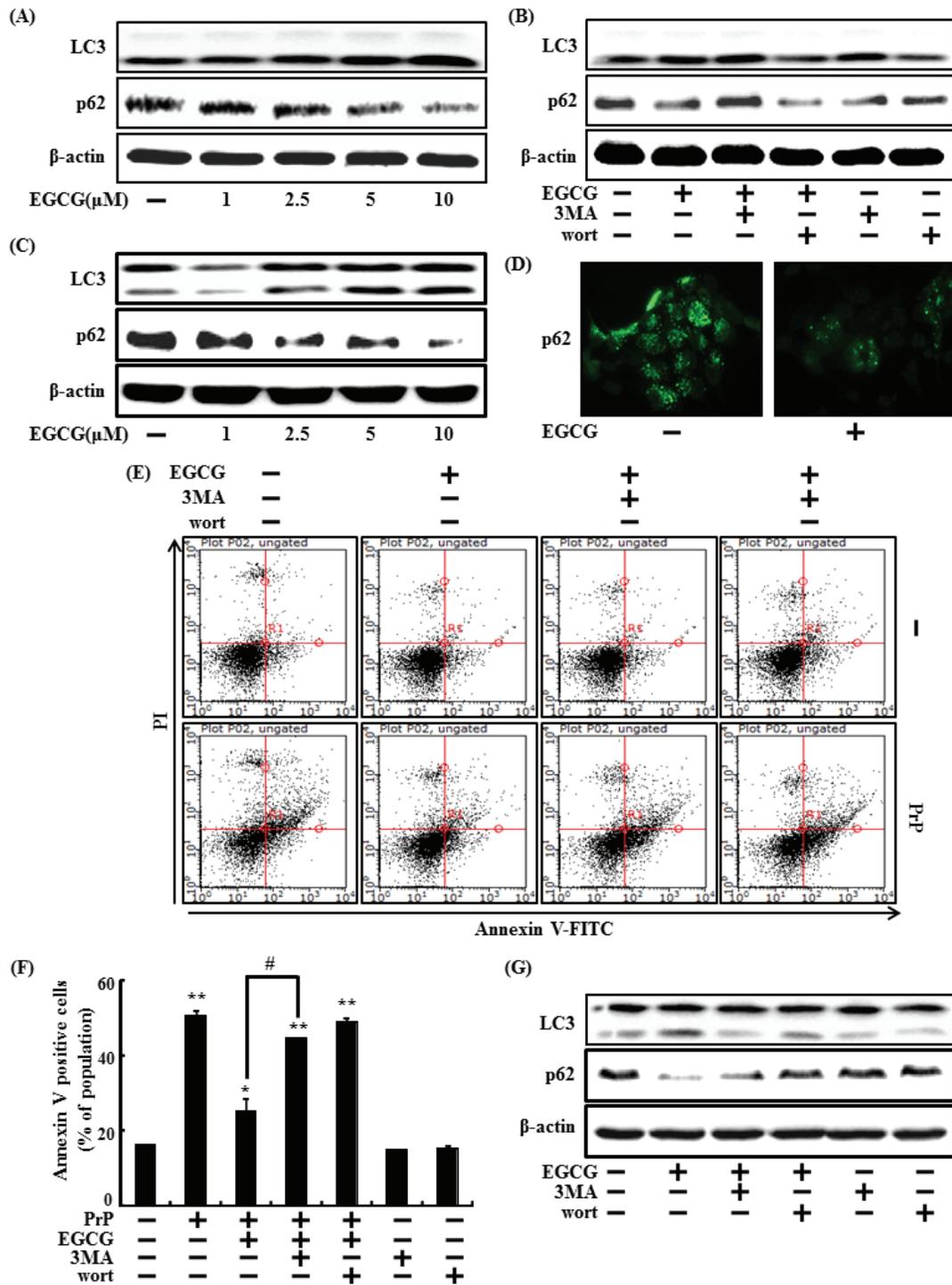


Figure 3: EGCG increases the induction of autophagy. The primary neuron cells were treated with 1, 2.5, 5, 10 μ M of EGCG for 30 h. Western blot for LC3-II, and p62 proteins was analyzed from primary neuron cells. Beta-actin was used as a loading control **A**. The cells were immunostained with p62 antibody (green) and observed in fluorescent view. Primary neuron cells were pretreated with 10 μ M of EGCG in presence of autophagy inhibitor (3MA and wortmannin) for 30 h and western blot for LC3-II and p62 proteins was analyzed **B**. SH-SY5Y cells were treated with 2.5, 5, or 10 μ M EGCG for 30 hr. Western blot for LC3-II, and p62 proteins was conducted with SH-SY5Y cells. Beta-actin was used as the loading control **C**. SH-SY5Y cells were analyzed by immunocytochemistry for p62 **D**. The cells were immunostained with p62 antibody (green) and observed in fluorescent view. SH-SY5Y cells were pretreated with 10 μ M of EGCG in the presence of autophagy inhibitors (3MA or wortmannin) for 1 hr and then exposed to 50 μ M PrP (106-126) for 36 h. Cell viability was measured by the Annexin V assay **E**, **F**. SH-SY5Y cells were treated as described in Figure 3A and Western blot for the LC3-II and p62 proteins was analyzed **G**. Bars indicate mean \pm standard error ($n = 4$). * $p < 0.05$, ** $p < 0.01$, significant differences between control and each treatment group, and # $p < 0.01$; significantly different when compared with PrP (106-126)-treated group.

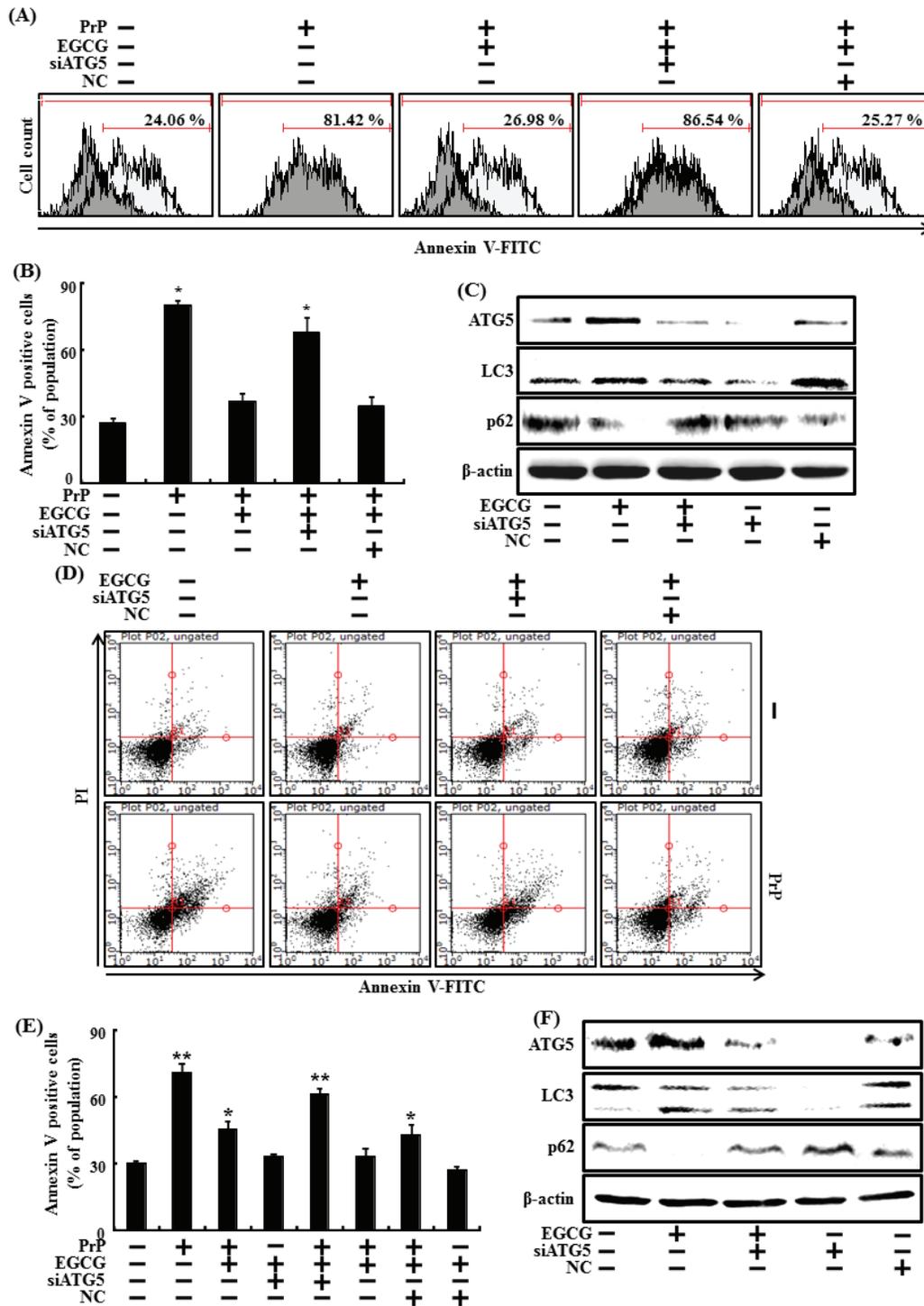


Figure 4: Inhibiting autophagy with ATG5 siRNA reduced the increase in autophagy caused by EGCG. ATG5 small interfering RNA (siATG5) or negative control siRNA (NC) transfected primary neuron cells were incubated with 50 μ M PrP (106-126) for 36 h in the presence of EGCG. Cell viability was measured by annexin V assay **A, B**. siATG5 or NC transfected primary neuron cells were incubated with EGCG (10 μ M) for 30 h. Western blot for ATG5, LC3 and p62 proteins was analyzed from SH-SY5Y cells. β -actin was used as loading control **C**. siATG5 or NC transfected SH-SY5Y cells were incubated with 50 μ M PrP (106-126) for 36 hr in the presence of EGCG. Cell viability was measured by Annexin V assay **D, E**. siATG5 or NC transfected SH-SY5Y cells were incubated with EGCG (10 μ M) for 30 hr. Western blot for ATG5, LC3-II and p62 proteins was analyzed from SH-SY5Y cells. Beta-actin was used as the loading control **F**. Bars indicate mean \pm standard error ($n = 4$). * $p < 0.05$, ** $p < 0.01$, significant differences between control and each treatment group, and # $p < 0.01$; significantly different when compared with PrP (106-126)-treated group.

And also we confirmed the effects of EGCG in the SH-SY5Y cells. SH-SY5Y cells were treated with EGCG (1, 2.5, 5, or 10 μ M, respectively) for 24 hr. EGCG increased LC3-II expression levels but decreased p62 levels (Figure 3C), and ICC also showed the reduction of p62 (Figure 3D). These data suggest that EGCG induced autophagy in SH-SY5Y cells.

We further tested to see whether inhibiting autophagy using autophagy inhibitors (3MA and wortmannin) could decrease the EGCG-induced neuroprotective effects against PrP (106-126). SH-SY5Y cells were pre-treated with autophagy inhibitors for 1 hr and then exposed to EGCG 10 μ M in presence of PrP (106-126). As shown in Figure 3E and 3F, the neuroprotective effects of the EGCG treatment decreased following application of the autophagy inhibitors. And also, in protein levels in the primary neuron cells and SH-SY5Y cells, the autophagy inhibitors restored LC3-protein II expression levels and decreased p62 by EGCG treatment, compared to those in the control (Figure 3B and 3G).

Knockdown of ATG5 using ATG5 small interfering RNA (ATG5 siRNA) decreased EGCG-induced autophagy (Figure 4C), as well as attenuated the neuroprotective effects caused by EGCG in both primary neuron cells (Figure 4A and 4B) and SH-SY5Y cells (Figure 4D and 4E).

EGCG inhibits the PrP (106-126)-induced apoptotic pathway by sirt1

sirt1 regulates key genes in neuronal survival [47] and stimulation of sirt1 is strongly affected by EGCG stability and metabolism [40]. To determine whether increased sirt1 was expressed, SH-SY5Y cells were treated with EGCG for 24 hr (Figure 5A). Sirt1 protein expression increased in a dose-dependent manner, and the result was reconfirmed by ICC (Figure 5B). Sirt1 binds with diverse transcription factors such as tumor suppressor p53, NF- κ B p65, and FOXOs. Because p53 is an important regulator of apoptosis induced by prion fragments [48, 49], it was thought that these factors may be related to EGCG-induced neuroprotection against the prion protein. p53 is mainly used as a DNA damage marker in the neuroblastoma cell line [50, 51]. Additionally, deacetylation of p53 results in a decrease in apoptotic cell death in neurons [52, 53]. These studies demonstrate that sirt1 plays an important role in neuroprotection by deacetylating and inhibiting p53-induced and NF κ B-induced inflammatory and apoptotic pathways. Sirt1 knockdown increased acetylated p53 level, leading to apoptosis in neurons. Hence, we examined whether EGCG contributed to deacetylation of p53 (Figure 5A). Acetylation of the p53 protein decreased in EGCG-treated cells. Consequently, EGCG up-regulated sirt1 activity, and treatment with PrP (106-126) resulted in sirt1 inactivation (Figure 5C). These results suggest that EGCG treatment is involved in both sirt1 expression and

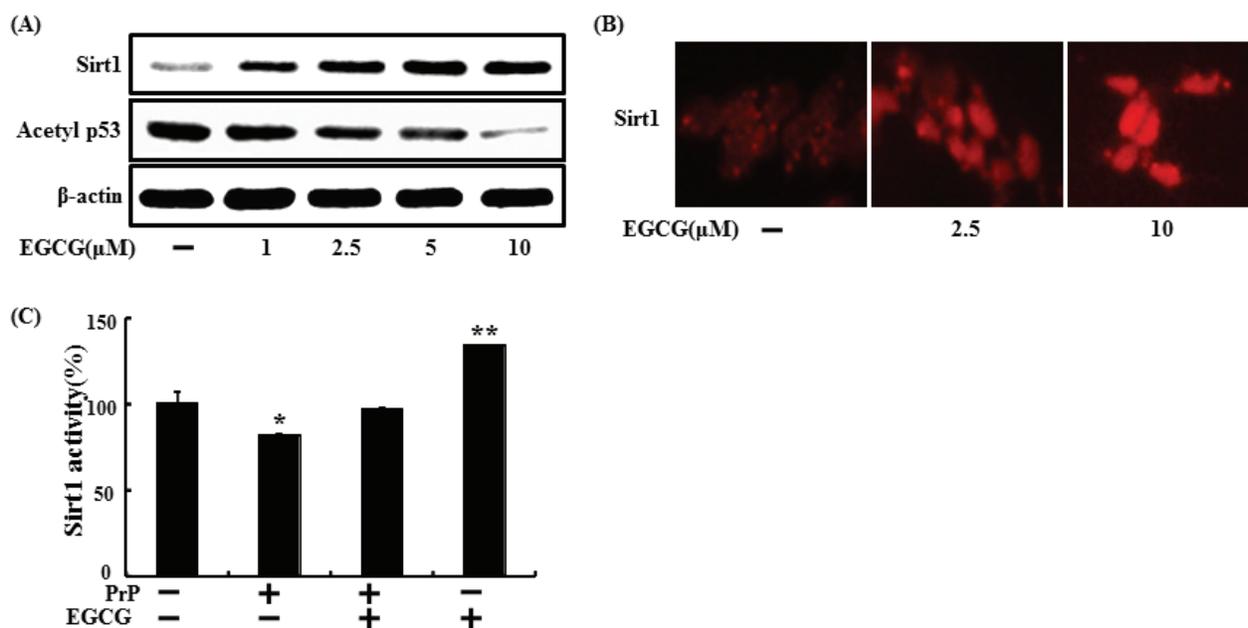


Figure 5: EGCG upregulates sirt1 expression and activation. SH-SY5Y cells were treated with 2.5, 5, or 10 μ M of EGCG for 30 hr. Western blot for sirt1 and acetyl-p53 proteins was analyzed from SH-SY5Y cells. Beta-actin was used as the loading control **A**. SH-SY5Y cells were analyzed by immunocytochemistry for sirt1 **B**. The cells were immunostained with sirt1 antibody (red) and observed in fluorescent view. SH-SY5Y cells were pretreated with 10 μ M EGCG for 1 hr and then exposed to 50 μ M PrP (106-126) for 36 hr. Sirt1 deacetylase activities in were analyzed in the nuclei of SH-SY5Y cells **C**. Bars indicate mean \pm standard error ($n = 4$). * $p < 0.05$, ** $p < 0.01$ significant differences between control and each treatment group.

activation.

Furthermore, sirt1 siRNA was used to knockdown sirt1 gene expression, to confirm that EGCG protects against prion-induced apoptosis by activating sirt1 and increasing sirt1 expression. We found that knockdown of sirt1 expression blocked the neuronal protective effects caused by EGCG, which inhibited neurotoxicity by PrP (106-126) (Figure 6A and 6B). No EGCG

mediated-neuroprotective effects were observed when sirt1 activation was inhibited by sirtinol (sirt1 activation inhibitor) (Figure 6C). These results show that the protective effect on the MTP by activating sirt1 was inhibited by transfection with sirt1 siRNA. The SH-SY5Y cells pre-treated with EGCG 10 μ M and exposed to 50 μ M PrP (106-126) decreased PrP (106-126)-induced JC-1 monomer fluorescence intensity, whereas cells transfected

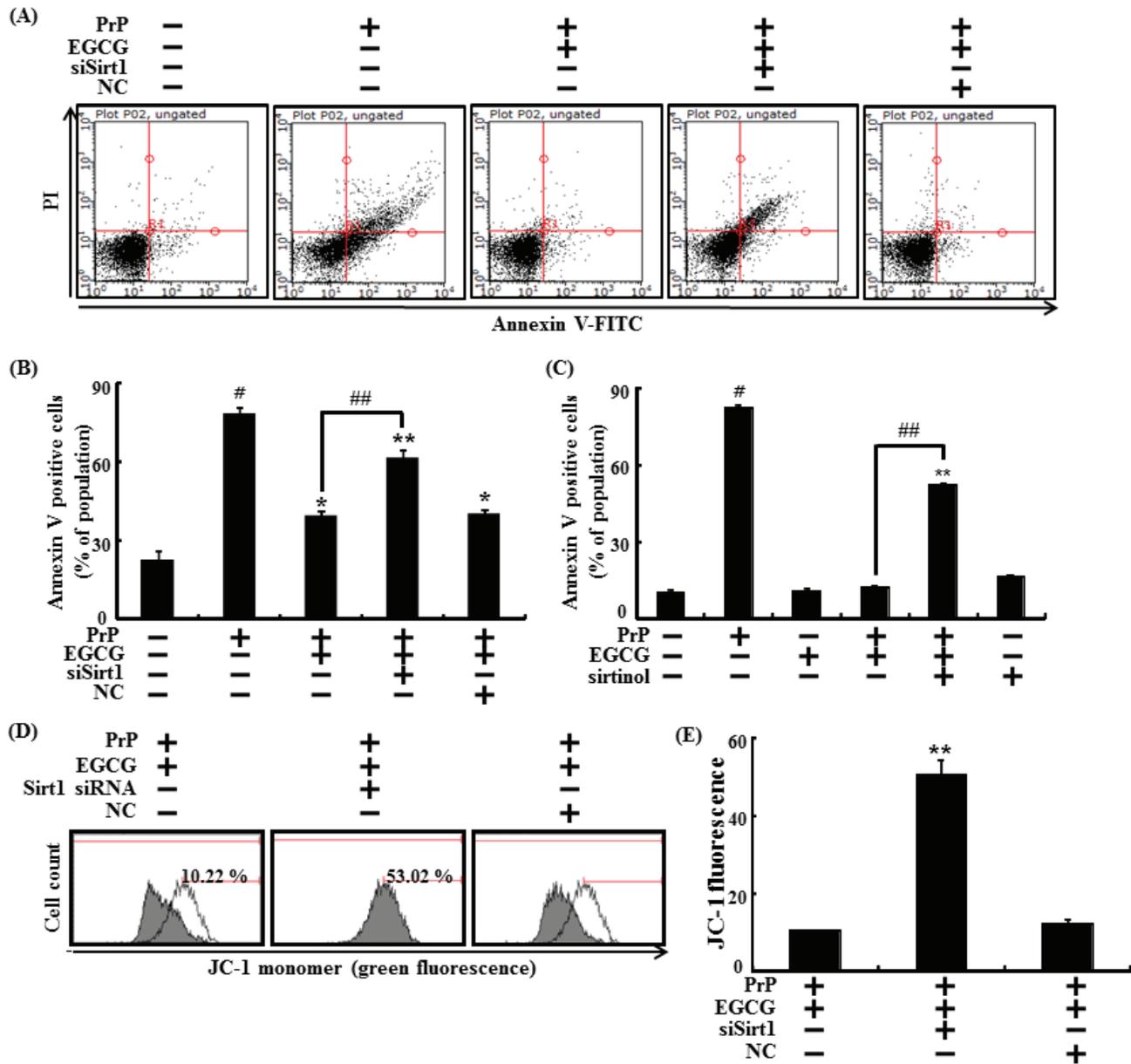


Figure 6: EGCG prevents neuronal cells from PrP (106-126)-induced cell death through the sirt1 pathway. Sirt1 small interfering RNA (siSirt1) or negative control siRNA (NC) transfected SH-SY5Y cells were incubated with 50 μ M PrP (106-126) for 36 hr in the presence of EGCG. Cell viability was measured by Annexin V assay **A, B**. SH-SY5Y cells were pretreated with sirtinol (10 μ M) and EGCG (10 μ M) for 1 h and then exposed to 50 μ M PrP (106-126) for 36 hr. Cell viability was measured by Annexin V assay **C**. The cells were measured for JC-1 mono form (green) by flow cytometry. M1 represents the population of JC-1 monomeric cells **D, E**. Bars indicates mean \pm standard error ($n = 4$). * $p < 0.05$, ** $p < 0.01$, # $p < 0.001$ significant differences between control and each treatment group, and ## $p < 0.01$; significantly different when compared with PrP (106-126)-treated group.

with sirt1 siRNA increased JC-1 monomer fluorescence intensity (Figure 6D and 6E).

We next investigated whether sirt1 knockdown inhibited sirt1 protein levels. SH-SY5Y cells transfected with sirt1 siRNA were treated with 10 μ M EGCG. As a result, EGCG increased sirt1 protein levels, which were blocked by sirt1 siRNA. Sirt1 knockdown contributed to p53 deacetylation, and p53 acetylation protein levels decreased in EGCG-treated cells (Figure 7A).

Sirt1 expression was measured by analyzing mRNA expression using RT-PCR to investigate whether sirt1 gene expression was blocked by sirt1 siRNA. Sirt1 mRNA level increased when SH-SY5Y cells were exposed to EGCG. However, sirt1 gene knockdown using sirt1 siRNA decreased sirt1 expression, compared to EGCG-treated cells (Figure 7B). In the same manner, transfecting sirt1 siRNA blocked increased sirt1 activity caused by EGCG treatment. Sirt1 activity did not change in the negative control (Figure 7C). Thus, we conclude that EGCG suppresses neuronal toxicity and mitochondrial damage by modulating both sirt1 activation and expression.

We also investigated whether sirt1 overexpression

by adenoviral transfection up-regulated protein expression. Ad-Sirt1, transfected at a multiplicity of infection (MOI) of 10-100, gradually increased sirt1 protein expression with protein levels at 100 MOI (Figure 8A). Consequently, sirt1 overexpression and EGCG increased sirt1 activities (Figure 8B). We then tested if sirt1 overexpression increased the neuroprotective effect. EGCG-treated or Ad-sirt1 transfected cells increased the protective effects compared to Ad-Lacz transfected cells (Figure 8C). These results suggest that EGCG up-regulates sirt1 protein levels and activates sirt1.

EGCG increase autophagy by activating sirt1

Next, we checked whether increased autophagy caused by EGCG was related to activating sirt1. We first measured autophagy protein levels when sirt1 was overexpressed using the sirt1 adenovirus to investigate the relationship between autophagy and sirt1. We found that overexpression of sirt1 up-regulated LC3-II and decreased p62 (Figure 9A). We then examined whether

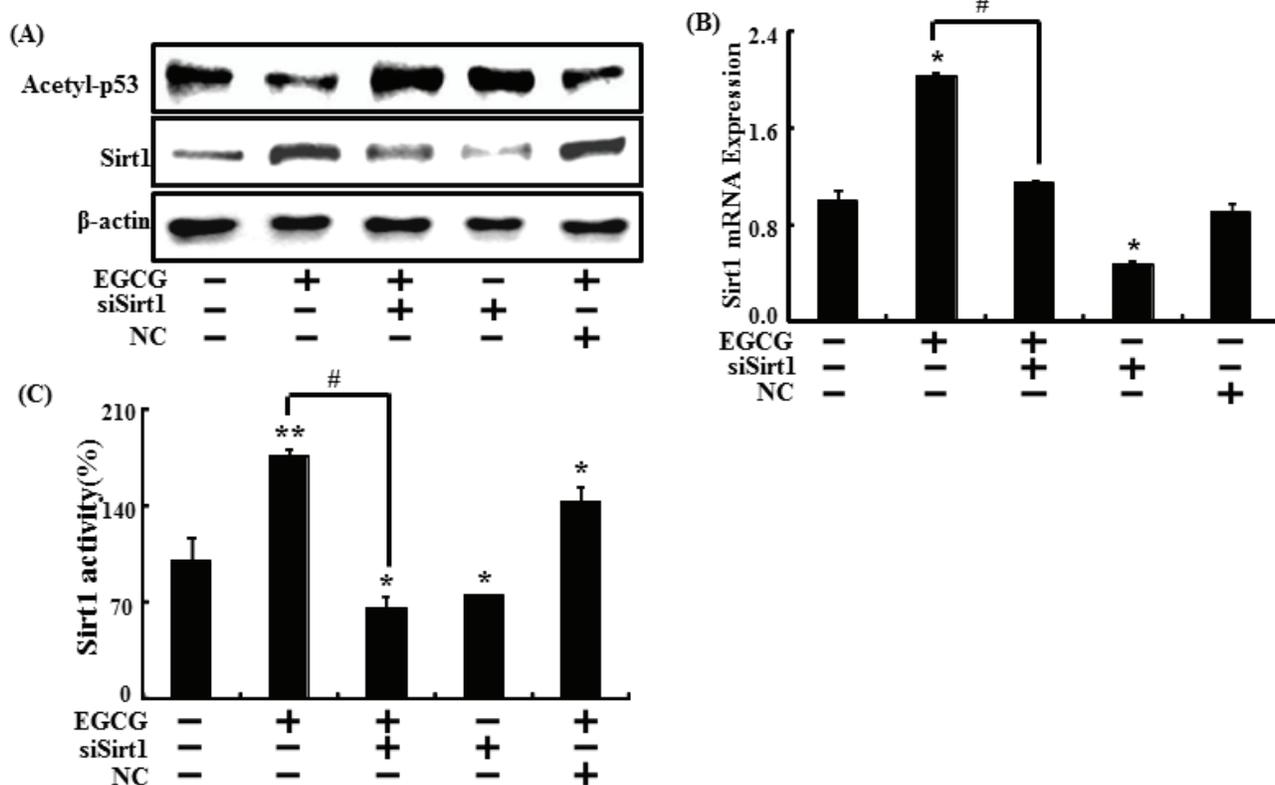


Figure 7: Inhibiting sirt1 decreased the sirt1 increase caused by EGCG. Sirt1 small interfering RNA (siSirt1) or negative control siRNA (NC) transfected SH-SY5Y cells were incubated with EGCG (10 μ M) for 30 hr. Western blot for sirt1 and acetyl-p53 proteins was conducted from SH-SY5Y cells. β -actin was used as the loading control **A**. Relative sirt1 mRNA expression levels were analyzed using quantitative real-time polymerase chain reaction. The indicated relative gene expression level shows expression levels that were normalized to β -actin expression as the standard **B**. siSirt1 or NC transfected SH-SY5Y cells were pre-incubated with EGCG (10 μ M) for 1 hr and then exposed to PrP (106-126) for 30 hr. Sirt1 deacetylase activities were analyzed in SH-SY5Y cell nuclei. **C**. Bars indicate mean \pm standard error ($n = 4$). * $p < 0.05$, ** $p < 0.01$, significant differences between control and each treatment group, and # $p < 0.01$; significantly different when compared with PrP (106-126)-treated group.

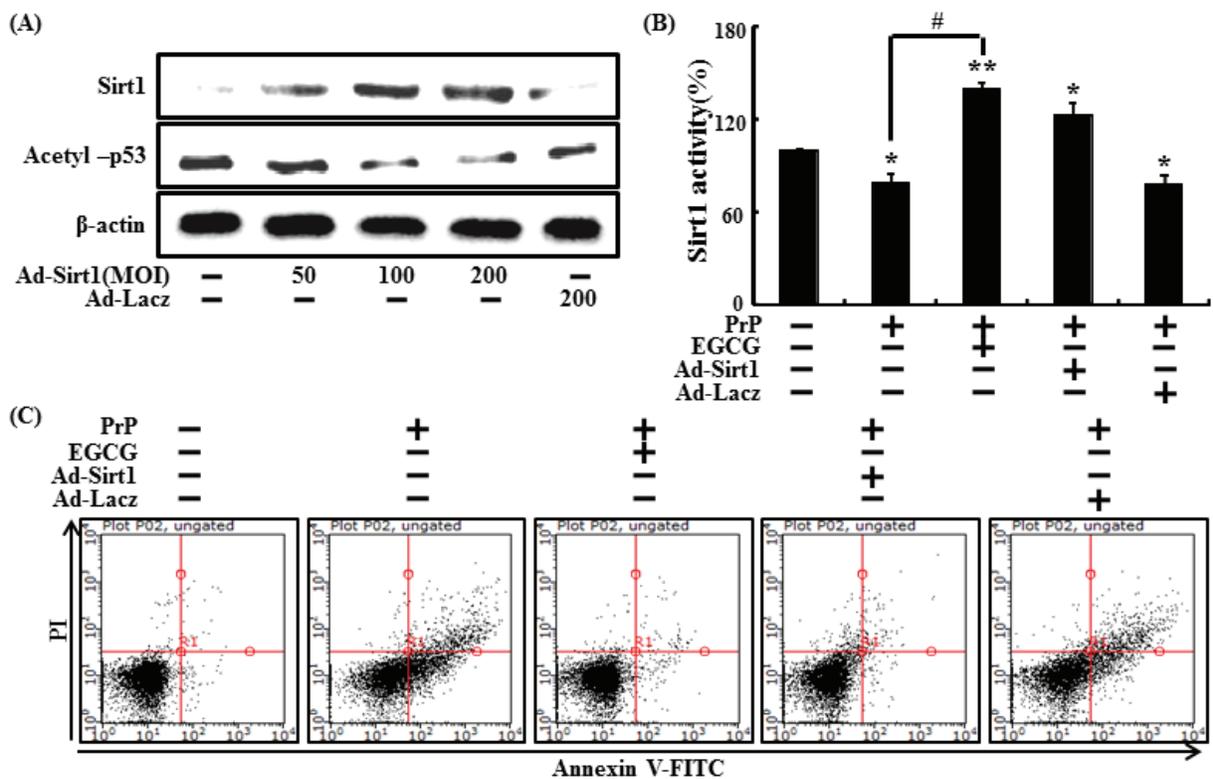


Figure 8: Overexpression of sirt1 and EGCG increased neuroprotective effects in SH-SY5Y cells. SH-SY5Y cells were transfected by overexpressing adenovirus (Ad-Sirt1) or lacZ-bearing adenovirus (Ad-lacZ). A Western blot for sirt1 and acetyl-p53 proteins was conducted from SH-SY5Y cells. Beta-actin was used as the loading control **A**. Ad-sirt1 or Ad-lacZ transfected cells were incubated with or without EGCG and exposed to PrP (106-126) for 36 hr. Sirt1 deacetylase activities were analyzed in SH-SY5Y cell nuclei **B**. Cell viability was measured by the Annexin V assay **C**. Bars indicate mean \pm standard error ($n = 4$). $*p < 0.05$, $**p < 0.01$, significant differences between control and each treatment group, and $\#p < 0.01$; significantly different when compared with PrP(106-126)-treated group.

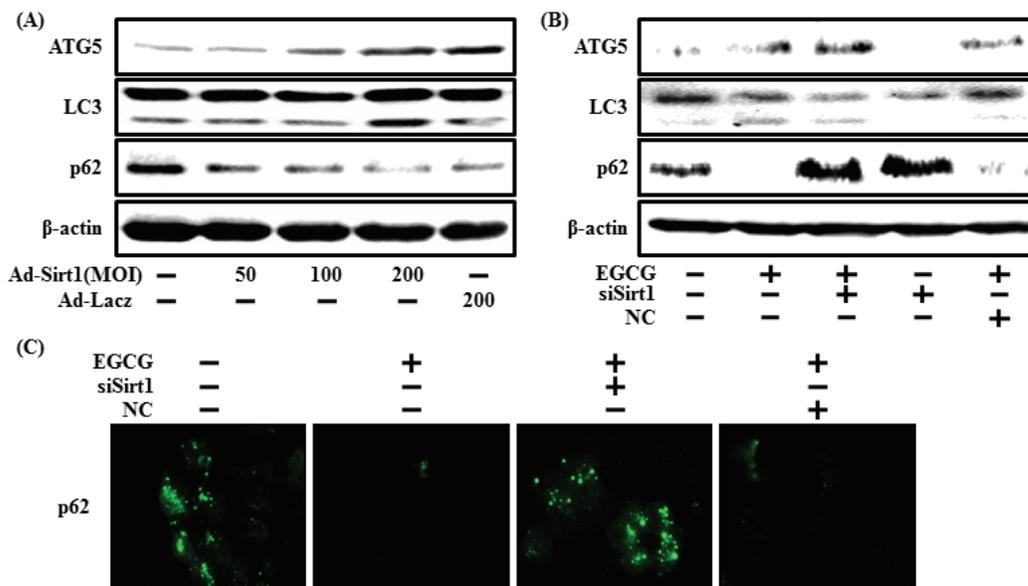


Figure 9: EGCG increases autophagy through the sirt1 pathway. The SH-SY5Y cells were transfected by overexpressing adenovirus (Ad-Sirt1) or lacZ-bearing adenovirus (Ad-lacZ). A Western blot of the LC3-II, ATG5 and p62 proteins was conducted in SH-SY5Y cells. Beta-actin was used as the loading control **A**. Sirt1 small interfering RNA (siSirt1) or negative control siRNA (NC) transfected SH-SY5Y cells were incubated with EGCG (10 μ M) for 30 hr. A Western blot for the LC3-II, ATG5, and p62 proteins was conducted with SH-SY5Y cells. Beta-actin was used as the loading control **B**. SH-SY5Y cells were treated as described in Figure 3A, and then ICC for p62 was analyzed **C**.

sirt1 knockdown by sirt1 siRNA affected EGCG-induced autophagy (Figure 9B). Inhibiting sirt1 decreased ATG5 and LC3-II but and increased p62 levels. Consistent with these results, ICC also showed increased p62 expression when sirt1 was knocked down. We conclude that inhibiting sirt1 decreases EGCG-induced autophagy and EGCG prevents prion-mediated neurotoxicity by inducing autophagy and activating sirt1.

DISCUSSION

This study demonstrated the possibility of attenuating prion protein-induced toxicity caused by EGCG, the major polyphenol in green tea, through autophagy activated by sirt1. We suggest that activating sirt1 with EGCG prevented mitochondrial damage related to prion-mediated toxicity. The biological effects of EGCG have already been demonstrated as it inhibits tumor growth of prostate, breast, liver, and lung cancers [4, 54, 55]. Studies have reported the potent effects of polyphenols against protein-misfolding diseases such as Alzheimer's, Parkinson's and Huntington's diseases.

In particular, EGCG leads to the formation of non-toxic amorphous aggregates through a disordered off-folding pathway [56] and modulates tau hyperphosphorylation [57]. The protective effects of EGCG have also been investigated in different experimental models of neurodegenerative diseases [58-61]. One study showed that EGCG plays an important role with PrPc, to protect cells from stress, and to degrade non-native PrPc conformations [62]. However, it was unknown if EGCG prevents prion-mediated neurotoxicity and the pathway of its protective actions is unknown.

The main goal of this study was to identify the mechanism of how EGCG acts in prion disease by examining the neuronal effects of EGCG against PrP (106-126) toxicity in neuroblastoma SH-SY5Y cells. The active polyphenol EGCG shows various pharmacological effects such as antioxidant, antimutagenic, antiproliferative, and anticarcinogenic properties, as well as neuroprotective activities [63]. EGCG also play roles as a neuroprotective compound at multiple levels and can chelate metal ions and scavenge free radical species directly [64]. EGCG may cross the blood-brain barrier (BBB) [65] by a specific uptake mechanism for glucuronides *in vivo*. The degree of BBB penetration of EGCG depends on lipophilicity. Brain entry may also be modulated by specific efflux transporters expressed in the BBB such as P-glycoproteins.

EGCG interacts with various protein kinase and lipid kinase signaling cascades in the brain such as phosphatidylinositol-3 kinase (PtdIns 3K)/Akt, protein kinase C, and mitogen-activated protein kinase (MAPK) pathways. These interactions increase the number of neurons and strengthen connections between neurons via neurotrophins such as brain derived neurotrophic factor that support and maintain cognitive function. The chemical

structures of the catechins, such as EGCG, contribute to their antioxidant properties. These compounds have a gallate moiety esterified site-specifically [42]. The potent free radical quenching activity of EGCG is affected by the presence of the gallate group [66]. Although we found protective effects of EGCG against apoptosis in neuroblastoma cells, many studies have revealed that EGCG induces non-apoptotic or apoptotic cell death in other cancer cells. Non-apoptotic cell death is stimulated by EGCG through autophagy and ROS [67, 68] and apoptosis occurs by activating caspases 3 and 9 [69]. Unlike other cancer cell lines, some studies have demonstrated that EGCG reduces apoptotic cell death in neuroblastoma cells [70, 71].

Autophagy is a complicated cellular process involved in cellular homeostasis and differentiation, as well as in tissue remodeling, aging, cancer, and other diseases [72]. During this process, targeted cytoplasmic constituents are isolated within autophagosomes, which are then fused with lysosomes for degradation or recycling [73]. We demonstrated that enhanced autophagy was caused by increased EGCG, due to increased expression of LC3-II and decreased expression of p62 in EGCG-treated SH-SY5Y cells, compared to control (Figure 3A). EGCG also prevented prion-mediated neuronal death (Figures 3 and 4). We further show the additional protective effect of EGCG as an autophagy inducer in SH-SY5Y cells. One study showed that EGCG may activate or inhibit various cellular signaling pathways such as NF- κ B, sirt1, MAPK, heat shock proteins, and numerous additional regulatory molecules. We confirmed that EGCG activates the sirt1 pathway by blocking sirt1 expression using sirt1 siRNA and by overexpressing sirt1 using sirt1 adenovirus (Figures 5,6,7,8).

Our results suggest that regulating sirt1 activation may be important for preventing neurodegenerative conditions including prion diseases. Overexpression of sirt1 also protects against neurodegenerative diseases including Alzheimer's and Huntington's disease as well as amyotrophic lateral sclerosis [31, 74, 75]. A previous study demonstrated that activating sirt1 protects against prion-mediated neurotoxicity [76]. In addition, resveratrol, which is a sirt1 activator, protects against MPTP-mediated cell degeneration and mitochondrial-oxidative damage by activating the sirt1/PGC-1 α pathways [77]. We showed here that prion-mediated neurotoxicity was protected by EGCG, as EGCG increased sirt1 protein expression and mRNA levels as well as sirt1 activities. Fewer neuroprotective effects were caused by sirt1 overexpression than by treatment with EGCG, indicating that although EGCG up-regulated sirt1 expression and activation, activating sirt1 has important effects beyond increased sirt1 expression.

We demonstrated that EGCG up-regulated autophagy via sirt1 as the main mechanism of neuroprotection against the prion peptide-induced

apoptotic pathway. This result suggests that EGCG may have pivotal role as a therapeutic target for prion diseases. Because PrP (106-126) is not equal to PrP^{Sc}, we could only confirm PrP^{Sc} toxicity. However, the 106-126 sequence of the prion protein is a useful model for *in vitro* study of prion-induced cell death, as it results in rapid depolarization of mitochondrial membranes. Some studies have shown that PrP (106-126) induces apoptotic-mediated cell death *in vivo* in model retinal neurons treated with an intravitreal injection of PrP fragments. [78]. As we only conducted our experiments with EGCG in cells, and not animal models, the neuroprotective effects of EGCG have yet to be demonstrated *in vivo*. Therefore, we shall further study whether EGCG has a neuroprotective effect that occurs through sirt1 and autophagy pathway in PrP (106-126) mouse models to identify possible anti-prion drugs and potential therapeutic role in prion diseases.

MATERIALS AND METHODS

Cell culture

The primary neurons were isolated from individual 1 day old pups. Briefly, tissues were collected in Hanks Buffered Saline Solution without Mg²⁺ and Ca²⁺ (HBSS; Invitrogen-GIBCO, Grand Island, NY, USA), and digested in 0.6% trypsin with 2.78mM glucose (30min, 37°C, 5% CO₂). Following trypsin neutralization with 1mg/ml soy trypsin inhibitor (Invitrogen, Carlsbad, CA, USA) containing 100U/ml DNase I (Invitrogen, Carlsbad, CA, USA), cells were mechanically dissociated and diluted in HBSS containing Mg²⁺ and Ca²⁺. Isolated neuron cells were diluted in DMEM containing 25mM glucose (abbreviated in text as 'glucose' medium; Thermo scientific) supplemented with 10% FBS, and plated in flasks pre-coated with 50µg/ml poly D-lysine. The human neuroblastoma cell line SH-SY5Y was obtained from the American Type Culture Collection (ATCC, Rockville, MD, USA). Cells were cultured in Minimum Essential Medium (MEM, Hyclone Laboratories, Logan, UT, USA) containing 10% fetal bovine serum (Invitrogen-GIBCO, Grand Island, NY, USA) and gentamycin (0.1 mg/mL) in a humidified incubator maintained at 37°C and 5% CO₂.

PrP (106-126) treatment

Synthetic PrP (106-126) with the sequence Lys-Thr-Asn-Met-Lys-His-Met-Ala-Gly-Ala-Ala-Ala-Ala-Gly-Ala-Val-Val-Gly-Gly-Leu-Gly and scrambled PrP (106-126) (sequence, Asn-Gly-Ala-Lys-Ala-Leu-Met-Gly-Gly-His-Gly-Ala-Thr-Lys-Val-Met-Val-Gly-Ala-Ala) was synthesized by Peptron, Inc. (Seoul, Korea). The peptide was dissolved in sterile dimethyl sulfoxide at 10 mM and stored at -80°C.

Annexin V assay

Apoptosis was assessed in detached cells using an Annexin V Assay kit (Santa Cruz Biotechnology, Santa Cruz, CA, USA), according to the manufacturer's protocol. Annexin V levels were determined by measuring fluorescence at 488 nm excitation and 525/30 emission using Guava EasyCyte HT System (Millipore, Bedford, MA, USA).

Lactate dehydrogenase assay

Cytotoxicity was assessed in supernatants using a lactate dehydrogenase (LDH) cytotoxicity detection kit (Takara Bio, Inc., Tokyo, Japan) according to the manufacturer's protocol. LDH activity was determined by measuring absorbance at 490 nm, using a microplate reader (Spectra Max M2, Molecular Devices, Sunnyvale, CA, USA).

Western blot analysis

After the SH-SY5Y cells were lysed in buffer (25 mM HEPES (4-(2-hydroxyethyl)-1-piperazineethanesulfonic acid), 100 mM NaCl, 1 mM EDTA, 5 mM MgCl₂, 0.1 mM DTT, and a protease inhibitor mixture, at pH 7.4), the proteins were electrophoretically resolved by 10-15% sodium dodecyl sulfate polyacrylamide gel electrophoresis and transferred to a nitrocellulose membrane. Immunoreactivity was detected through sequential incubations with horseradish peroxidase-conjugated secondary antibodies and enhanced chemiluminescence reagents. The antibodies used for immunoblotting were LC3 (Novus Biologicals, Littleton, CO, USA), Bax (Santa Cruz Biotechnology, Santa Cruz, CA, USA), cytochrome *c* (BD BioScience, San Jose, CA, USA), autophagy related 5 (ATG5) (Abcam, Cambridge, MA, USA), Sirt1 (Santa Cruz Biotechnology, CA, USA), acetyl-p53 (Abcam, Cambridge, MA, USA) and β-actin (Sigma Aldrich, St. Louis, MO, USA). Images were examined using a Fusion FX7 imaging system (VilberLourmat, Torcy Z.I. Sud, France).

Quantitative real-time polymerase chain reaction (qRT-PCR)

Total ribonucleic acid (RNA) was extracted from SH-SY5Y cells using the Easy-spin™ Total RNA Extraction kit (iNtRON Biotechnology, Seoul, Korea). The cDNA synthesis was carried out following the instructions of TaKaRa Prime Script™ 1st strand cDNA synthesis kit (TaKaRa Bio, Tokyo, Japan). For qRT-PCR, 1 µL of gene primers with SYBR Green (Bio-Rad Laboratories, Hercules, CA, USA) in 20 µL

of reaction volume was applied. The primers were: SIRT1 forward, 5'TGCTCGCCTTGCTGTAGACTTC3', reverse, 5'GGCTATGAATTTGTGACAGAGAGATGG3'; β -actin (as an internal control): forward, 5'GCAAGCAGGAGTATGACGAG3', reverse, 5'CAAATAAAGCCATGCCAATC3'. All reactions with iTaq SYBR Green Supermix (Bio-Rad Laboratories, Hercules, CA, USA) were performed on the CFX96 real-time PCR detection system (Bio-Rad Laboratories, Hercules, CA, USA).

Mitochondrial transmembrane potential assay

The change in mitochondrial transmembrane potential (MTP) was evaluated with a JC-1 cationic fluorescent indicator (Molecular Probes, Eugene, OR, USA) which aggregates in intact mitochondria (red fluorescence) indicating normal MTP and when it remains in monomeric form in the cytoplasm (green fluorescence) indicating low MTP. SH-SY5Y cells were incubated in MEM containing 10 μ M JC-1 at 37°C for 15 min, washed with PBS, and then transferred to a clear 96-well plate. The JC-1 aggregate fluorescent emissions were measured at 583 nm with an excitation wavelength of 526 nm, whereas the JC-1 monomer fluorescence intensity was measured with excitation and emission at 525 and 530 nm, respectively, using a Guava EasyCyte HT System (Millipore, Billerica, MA, USA) or a microplate reader (Spectra Max M2, Molecular Devices). SH-SY5Y cells were cultured on coverslips in a 24-well plate, incubated in MEM containing 10 μ M JC-1 at 37°C for 15 min, and then washed with PBS. Finally, the cells were mounted with Dako Fluorescent mounting medium and visualized via Nikon Eclipse 80i fluorescence microscopy (Nikon, Tokyo, Japan).

Construction of recombinant adenoviruses

The Sirt1 over-expressing adenovirus (Ad-Sirt1) was provided by Professor Byung-Hyun Park of Chonbuk National University (Jeonju, Jeonbuk, South Korea). The lacZ-bearing adenovirus (Ad-lacZ) was used as a control. Recombinant adenoviruses were amplified in human embryonic kidney (HEK)-293 cells and purified using the Vivapure AdenoPACK kit (Sartorius AG, Göttingen, Germany) according to manufacturer's instructions.

RNA interference

SH-SY5Y cells were transfected with ATG5 small interfering RNA (siRNA; oligoID HSS114104; Invitrogen, Carlsbad, CA, USA) or Sirt1 siRNA (oligoID VHS50608; Invitrogen) using Lipofectamine 2000 according to manufacturer's instructions. After a 48-hr culture,

knockdown efficiency was measured at the protein level by immunoblot analysis. Nonspecific siRNA (oligoID 12935-300; Invitrogen) was used as the negative control.

Sirt1 deacetylase activity assay

To measure cellular Sirt1 deacetylase activity, nuclear proteins were extracted from SH-SY5Y cells using a Nuclear/Cytosol Fractionation kit (BioVision, Milpitas, CA, USA). Sirt1 deacetylase activity was quantified following the protocols of the Sirt1 Fluorometric Assay kit (Sigma-Aldrich). Fluorescence intensities were measured with a microplate fluorometer (excitation wavelength = 360 nm, emission wavelength = 450 nm). The fluorescence intensities of Sirt1 deacetylase activity were normalized to the protein levels measured in the cell samples.

Statistical evaluation

All data are expressed as mean \pm standard deviation and were compared by Student's *t*-test, analysis of variance and Duncan's test using SAS statistical software ver. 9.1 (SAS Institute, Cary, NC, USA). Results were considered significant at **p* < 0.05, ***p* < 0.001 and #*p* < 0.01.

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CONFLICTS OF INTEREST

The authors declare no conflicts of interest.

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