- 1 Ginsenoside Rg1 defenses PC-12 cells against hydrogen peroxide-caused damage
- via up-regulation of miR-216a-5p
- 4 Running title: Protective mechanism of ginsenoside Rg1
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19 Abstract

- 20 **Background:** Spinal cord injury (SCI) is a destructive trauma accompanying with
- 21 local injury, half of which cause chronic paralysis. Ginsenoside Rg1 exerts
- 22 anti-apoptosis and anti-autophagy properties. Therefore, our goal was to study the
- 23 protective mechanism of Rg1 in attenuating cell injury.
- Methods: MiR-216a-5p inhibitor was transfected into PC-12 cells, then cells were
- 25 pre-treated by Rg1 and treated with 300 μM hydrogen peroxide (H<sub>2</sub>O<sub>2</sub>) for 24 h.
- 26 CCK-8 and apoptosis experiments were done to test cell activity and apoptosis
- 27 respectively. Expression of miR-216a-5p and cell damage relative factors was tested
- via qRT-PCR and western blot experiments, respectively.
- 29 **Results:** H<sub>2</sub>O<sub>2</sub> induced cell activity suppression, apoptosis and autophagy well at the
- 30 concentration of 300 μM, leading cell injury. Rg1 could attenuate cell injury induced
- 31 by H<sub>2</sub>O<sub>2</sub> at the working concentration of 200 μM that it elevated cell activity,
- 32 attenuated apoptosis and autophagy and activated PI3K/AKT and AMPK signal
- pathways. Further, miR-216a-5p was up-regulated by Rg1. Rg1 played its role in
- relieving cell injury by positively regulating miR-216a-5p.
- 35 Conclusion: Our study demonstrated that Rg1 attenuated H<sub>2</sub>O<sub>2</sub>-caused cell injury
- through positively regulated miR-216a-5p.
- 37 **Key words:** ginsenoside Rg1, miR-216a-5p, cell injury

## Introduction

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41 Spinal cord injury (SCI), a common and destructive trauma (Hyun and Kim, 2010), is 42 mainly caused by external forces such as lateral bending, excessive stretching, 43 rotation, axial load and excessive bending, resulting in motor dysfunction, paralysis 44 and other symptoms (van den Berg et al., 2010). Because of the limited therapy 45 selection, the administration and care of SCI patients places a heavy burden on 46 patients and caregivers. Of particular note, more than 60% of damages occur at the 47 cervical level (Cripps et al., 2011), and the lifetime care costs are about at \$1.1-\$4.2 48 million per patient (Krueger et al., 2013). So, the precaution, curing and recovery of 49 SCI are a major topic in the medical area. SCI relates two different stages of tissue 50 injury, called primary and secondary hurt (Badner et al., 2019). Local tissue injury is 51 caused by SCI and is important in secondary hurt in SCI (Fu et al., 2018), leading to 52 apoptosis with loss of neurological roles. So, mechanism study of local injury after 53 SCI is highly significant for curing SCI. 54 Ginsenosides, considering as one of the main pharmacological active ingredients of 55 ginseng, is a steroid compound (Xiang et al., 2008). Ginsenoside contains the 56 Panaxatriol (Rg1, Rg2, Re and Rf) and Panaxadiol (Rb1, Rb2, Rc and Rd) classes 57 (Zhang et al., 2012a). Many beneficial effects of Rg1 have been proved in disorders 58 such as hypertension (Chen et al., 2012), hypoxia/reoxygenation (Zhang et al., 2012b), 59 Alzheimer's disease (Huang et al., 2012) etc. Importantly, it has been reported that 60 Rg1 exerts roles in inhibiting cell apoptosis, thereby exhibiting notable 61 cardioprotective effects against I/R damage through a variety of mechanisms (Lee and 62 Kim, 2014). Besides, Rg1 counteracts the aging of endothelial progenitor cells (Shi et 63 al., 2011) and human fibroblasts (Zhou et al., 2012) and exerts a notable influence in 64 suppressing cardiomyocytes and renal tubular cells' autophagy (Mao et al., 2016). The 65 influence of Rg1 in local injury after SCI still has been unknown yet. 66 MicroRNAs (miRNAs), short (22 nucleotides in length) non-coding RNAs, involve 67 in many biological processes (Jiang and Chen, 2012), such as differentiation of 68 ordinary tissues and are important in the pathogenesis of lots of human cancers (Taucher et al., 2016). MiR-216a-5p, known as an oncogene, involved in the 69 70 progression of many cancer subtypes (Chen et al., 2018). Chen et al. has proved that 71 miR-216a-5p elevates cell proliferation, activity and motility, and inhibits apoptosis 72 (Chen et al., 2018). This finding demonstrates that miR-216a-5p has a positive effect 73 on cell viability and anti-apoptosis. So it could be interesting to investigate if exerts

- 74 regulation relation of miR-216a-5p and Rg1 in cell injury after SCI. Based on the
- above questions and guesses, we probed mechanism of Rg1 against H<sub>2</sub>O<sub>2</sub>-caused cell
- 76 damage in PC-12 cells.

#### 77 Materials and Methods

- 78 *Cell*
- 79 PC-12 cells were bought form Kunming Institute of Zoology (Kunming, China) in
- this whole study. Seed cells at a denseness of  $1 \times 10^4$  cells/ml in Dulbecco's Modied
- 81 Eagle Medium (DMEM)/F-12 medium (Gibco, Carlsbad, CA, USA) adding with 10%
- 82 fetal bovine serum (FBS, Gibco), 100 μg/ml streptomycin and 100 U/ml penicillin
- 83 (Gibco). Cells were kept in a wet incubator carried 5% CO<sub>2</sub> and 95% air at 37°C.
- 84 Change fresh medium every day. Ginsenoside Rg1 (analysis level of 97% pureness)
- was bought from Sigma-Aldrich (St. Louis, MO, USA), solubled in ethanol and stored
- at -20°C. Pre-treatment of cells with Rg1 for 1 h, and then were treated with a series
- of consistences of hydrogen peroxide  $(H_2O_2)$  for 24 h.

#### 88 CCK-8 experiment

- 89 A Cell Counting Kit-8 (CCK-8, Dojindo Molecular Technologies, Gaithersburg, MD,
- 90 USA) was to test cell activity. Seed cells in 96-well plate with 5000 cells/well, and
- 91 then add CCK-8 solution, keep cells in a wet environment carried 95% air and 5%
- 92 CO<sub>2</sub> for 1 h at 37°C. Absorbance was tested at 450 nm via a Microplate Reader
- 93 (Bio-Rad, Hercules, CA, USA).

#### 94 Apoptosis experiment

- 95 Apoptosis analysis was done through propidium iodide (PI) and fluorescein
- 96 isothiocynate (FITC)-conjugated Annexin V staining (BD Pharmingen, San Diego,
- 97 CA, USA). Cells were cleaned in phosphatebuffered saline (PBS) for three times and
- 98 stained in PI/FITC-Annexin V with 50 μg/ml RNase A (Sigma-Aldrich). Keep cells in
- 99 dark processing at the room temperature for 1 h. Flow cytometry analysis was made
- through FACS can (Beckman Coulter, Fullerton, CA, USA). Data was analyzed via
- 101 FlowJo software (Tree Star Software, San Carlos, California, USA).

## 102 Transfection

- MiR-216a-5p inhibitor and its relative NC were compounded by Life Technologies
- 104 Corporation (Carlsbad, CA, USA) and transferred into cells. Transfection was done
- following the Lipofectamine 3000 reagent (Life Technologies Corporation). 48 h
- post-transfection was regarded as harvest moment in following assays.

## 107 *qRT-PCR*

- 108 Overall RNA was extracted through Trizol reagent (Life Technologies Corporation)
- and handled with DNaseI (Promega, Madison, WI, USA). Taqman MicroRNA
- 110 Reverse Transcription Kit and Taqman Universal Master Mix II with the TaqMan
- 111 MicroRNA Assay (Applied Biosystems, Foster City, CA, USA) were to test
- miR-216a-5p expression. U6 was taken as inside comparison.
- 113 Western Blot
- Overall protein was extracted through RIPA lysis buffer (Beyotime Biotechnology,
- Shanghai, China) with protease inhibitors (Roche, Basel, Switzerland), and then
- quantified through BCA<sup>TM</sup> Protein Assay Kit (Pierce, Appleton, WI, USA). A Bio-Rad
- 117 Bis-Tris Gel system was taken to build up a western blot system. Primary antibodies
- specific against Bax (ab32503, Abcam, Cambridge, MA, USA), pro-caspase-3
- 119 (ab183179), cleaved-caspase-3 (ab49822), pro-PARP (ab32064), cleaved-PARP
- 120 (ab4830), β-actin (ab8226), beclin-1 (ab62557), p62 (ab56416), LC3-I and LC3-II
- 121 (ab48394), t-PI3K (ab140307), p-PI3K (ab182651), t-AKT (ab179463), p-AKT
- 122 (ab38499), t-AMPK (ab131512) and p-AMPK (ab23875) were readied in 5%
- blocking buffer. Primary antibody was cultured with membrane at 4°C all the night,
- then washing and incubating with secondary antibody, marking by horseradish
- peroxidase for 1 h at room temperature. Then the Polyvinylidene Difluoride (PVDF)
- membrane taken along blots and antibodies were transferred into the Bio-Rad
- 127 ChemiDoc<sup>TM</sup> XRS system, adding 200 μl Immobilon Western Chemiluminescent
- 128 HRP Substrate (Millipore, MA, USA) to shroud film surface. At last, semaphores
- were seized and strength of strip was quantified via Image Lab<sup>TM</sup> Software (Bio-Rad).
- 130 Statistical analysis
- 131 All assays were duplicated for 3 times. Our consequences of multiplex assays are
- revealed as mean ± SD. Statistical analysis was done via Graphpad Prism 6.0
- 133 (GraphPad Software Inc., La Jolla, CA, USA). P-values were counted via a one-way
- analysis of variance (ANOVA). P-value of < 0.05 indicated statistical significant data.
- 135 Results
- 136 Rg1 extenuated  $H_2O_2$ -induced cell activity suppression and cell apoptosis
- PC-12 cells were treated in various H<sub>2</sub>O<sub>2</sub> consistences. From **Figure 1A**, we found
- that  $H_2O_2$  had notably inhibiting effect on cell viability when the concentration was
- 139 100 (P < 0.05), 200 (P < 0.05), 300 (P < 0.01), 400 (P < 0.01) and 500  $\mu$ M (P < 0.01)
- 140 0.001). We chose 300  $\mu$ M as the working concentration in the later assays because this
- was cell viability semi-lethal concentration. Besides, we tested effect of H<sub>2</sub>O<sub>2</sub> on cell

- apoptosis. We found that apoptosis was notably increased by  $H_2O_2(P < 0.001, Figure)$
- 143 1B). Similarly, apoptosis relative factors (Bax, cleaved-caspase-3 and
- cleaved-caspase-PARP) were obviously enhanced through H<sub>2</sub>O<sub>2</sub> (Figure 1C), and
- standards of these factors were notably raised (all P < 0.01, **Figure 1D**). We got that
- $H_2O_2$  caused cell activity suppression and apoptosis.
- For function of Rg1, following experimental results were clear. As shown in Figure
- 148 **1E**, there was no effect on cell viability by Rg1. We found that H<sub>2</sub>O<sub>2</sub> could notably
- reduce cell viability (P < 0.01), whereas Rg1 could notably attenuate this reduction at
- 200, 300 and 400 μM (all P < 0.05). We chose 200 μM as the working concentration
- in the following experiments because this is the concentration when cell viability was
- half restored. Besides, for cell apoptosis, we found that Rg1 attenuated apoptosis
- induced by  $H_2O_2$  (P < 0.01, Figure 1G). Similarly, Figure 1H revealed that
- expression of apoptosis relative factors was weakened by Rg1 compared with H<sub>2</sub>O<sub>2</sub>
- group. Levels of these factors were raised through  $H_2O_2$  (P < 0.01, P < 0.001 and P < 0.001
- 156 0.01), whereas Rg1 could decrease their levels (all P < 0.05, Figure 11). So we got
- that Rg1 attenuated cell activity suppression and apoptosis induced by H<sub>2</sub>O<sub>2</sub>.
- 158 Rg1 extenuated autophagy induced by  $H_2O_2$
- For autophagy, we tested three autophagy relative factors. Beclin-1 is autophagy gene
- and its overexpression can stimulate autophagy (Yue et al., 2003). Accumulation of
- p62 is a notable phenotype of autophagy-deficient tumor cells (Mathew et al., 2009).
- 162 LC3-II is a marker for mature autophagosomes. Autophagy could be analyzed by
- testing the conversion of the autophagosome marker LC3-I to LC3-II (Wu et al.,
- 2010). According to our results, **Figure 2A** showed the enhancement of beclin-1 and
- 165 LC3-II/LC3-I through H<sub>2</sub>O<sub>2</sub>, while Rg1 could weaken this enhancement. Expression
- of p62 was weakened by  $H_2O_2$ , while Rg1 could eliminate this mitigation (**Figure 2A**).
- 167 Besides, Figure 2B revealed the notable addition of beclin-1 and
- LC3-II/LC3-Ithrough  $H_2O_2$  (P < 0.01 and P < 0.001), whereas were opposite by
- adding of Rg1 (P < 0.05 and P < 0.01). p62 expression was notably weakened
- through  $H_2O_2$  (P < 0.05), whereas was increased by the adding of Rg1 (P < 0.05,
- 171 **Figure 2B**). So we got that Rg1 could attenuate H<sub>2</sub>O<sub>2</sub>-induced autophagy.
- 172 Rg1 positively regulated miR-216a-5p
- 173 From Figure 3, qRT-PCR assay indicated that miR-216a-5p was notably
- down-regulated after  $H_2O_2$  treatment (P < 0.05). But, it was specifically up-regulated
- by adding Rg1 (P < 0.01). So we got that Rg1 up-regulated miR-216a-5p.

- 176 Rg1 extenuated cell activity suppression, apoptosis and autophagy induced by  $H_2O_2$
- 177 through up-regulating miR-216a-5p
- 178 qRT-PCR revealed that miR-216a-5p expression was notably suppressed after
- miR-216a-5p inhibitor transfection (P < 0.01, Figure 4A). Cell viability was notably
- aggravated by Rg1 contrast with  $H_2O_2$  set (P < 0.05), whereas was notably alleviated
- when treated with Rg1 plus miR-216a-5p inhibitor (P < 0.05, **Figure 4B**). This result
- indicated that Rg1 attenuated H<sub>2</sub>O<sub>2</sub>-induced cell activity suppression by up-regulating
- miR-216a-5p. Besides, cell apoptosis was notably decreased by Rg1 contrast with
- $H_2O_2$  set (P < 0.01), whereas was notably increased when treated with Rg1 plus
- miR-216a-5p inhibitor (P < 0.01, Figure 4C). Figure 4D-E further indicated that
- levels of apoptosis relative factors were notably decreased through Rg1 contrast with
- H<sub>2</sub>O<sub>2</sub> set (P < 0.05, P < 0.01) and P < 0.01), whereas were raised in Rg1-treated cells
- with miR-216a-5p inhibitor (P < 0.05, P < 0.01 and P < 0.05). So we got that Rg1
- reduced H<sub>2</sub>O<sub>2</sub>-caused apoptosis by up-regulating miR-216a-5p. Additionally, **Figure**
- **4F-G** revealed the notable attenuation of beclin-1 and LC3-II/LC3-I (P < 0.01 and P
- 191 < 0.001) by Rg1 and level of p62 was increased (P < 0.05) by Rg1 constrast with
- H<sub>2</sub>O<sub>2</sub> set. However, beclin-1 and LC3-II/LC3-I were notably enhanced (both P < 0.05)
- and level of p62 was notably decreased when treated with Rg1 plus miR-216a-5p
- inhibitor (P < 0.05). So we got that Rg1 attenuated  $H_2O_2$ -induced autophagy by
- up-regulating miR-216a-5p.
- 196 Signal pathway
- 197 To further study the mechanism of Rg1, we focused on PI3K/AKT and AMPK signal
- pathways. Figure 5A-B indicated the notable addition levels of p-PI3K and p-AKT
- through Rg1 contrast with  $H_2O_2$  set (P < 0.01) and P < 0.05), whereas were notably
- alleviated in Rg1-treated cells with miR-216a-5p inhibitor (P < 0.01 and P < 0.05).
- Besides, **Figure 5C-D** revealed that level of p-AMPK was notably aggravated by Rg1
- compared with  $H_2O_2$  group (P < 0.01), whereas was notably alleviated in Rg1-treated
- cells with miR-216a-5p inhibitor (P < 0.01). These results indicated that Rg1 elevated
- 204 PI3K/AKT and AMPK pathways via positively modulating miR-216a-5p.
- 205 **Discussion**
- 206 SCI is a life-changing event. Recently, there is no effective treatment method to
- resume the functions of SCI patients. The complexity of SCI pathophysiology poses a
- 208 huge challenge for researchers and clinicians seeking to develop therapeutic
- interventions (Bareyre, 2019). Local injury is the main event of secondary injury in

210 SCI, eventually leading to apoptosis and ultimately loss of neurological function 211 (Genovese et al., 2009). Therefore, defensing cells against local injury or relieving 212 this injury can be an effective method to cure SCI. Rg1, an active component of 213 ginsenosides, has been proved to exert positive effect on anti-apoptosis (Zu et al., 214 2016). Cell damage has been reported to be induced by H<sub>2</sub>O<sub>2</sub> in cardiomyocytes 215 (Zeng et al., 2019) and neural cells (Chen et al., 2016). Therefore, our study also 216 found that H<sub>2</sub>O<sub>2</sub> was a mediator capable of inducing damage in PC-12 cells, including 217 activity suppression, promotion of autophagy and apoptosis. Our study firstly 218 researched the attenuated mechanism of Rg1 in H<sub>2</sub>O<sub>2</sub>-caused damage in PC-12 cells. 219 Rg1 was effective in defensing PC-12 cells against H<sub>2</sub>O<sub>2</sub>-caused damage. Rg1 could 220 extenuate H<sub>2</sub>O<sub>2</sub>-induced cell activity suppression, apoptosis and autophagy, and active 221 PI3K/AKT and AMPK pathways via positively regulating miR-216a-5p. Our results 222 indicated that Rg1 may be an effective treatment of curing SCI. Ginsenoside Rg1, the main bioactive ingredient in ginseng, has been proved to exert 223 224 low toxicity that there was no change on cell viability and proliferation (Li et al., 225 2017). Much evidence indicates that Rg1 exerts beneficial effects, like anti-aging 226 properties (Zhu et al., 2014). As we all know, Bax may control mitochondrial 227 permeability transition and promote releasing cytochrome c, ultimately triggering the 228 activation of caspases, leading to apoptosis (Li et al., 2017). Consistently, in our study, 229 treatment of Rg1 reduced the level of Bax, at the same time, cleaved-caspase-3 and 230 cleaved-caspase-PARP were weakened. These observations verified the anti-apoptosis 231 function of Rg1. Additionally, autophagy is an important cellular process where cytoplasmic components are digested by lysosomes to keep cell homeostasis and 232 energy production (Ravikumar et al., 2010). Rg1 has a notable pharmacological 233 234 influence in suppressing autophagy (Mao et al., 2016). Our study were consistent with 235 the report that Rg1 strongly inhibited autophagic factor (beclin-1 and LC3-II/LC3-I), 236 and counteracted attenuation of p62 induced by H<sub>2</sub>O<sub>2</sub>, leading to inhibiting 237 H<sub>2</sub>O<sub>2</sub>-induced autophagy in PC-12 cells. Consistently, our study indicated that Rg1 238 could counteract inhibition of cell activity induced by H<sub>2</sub>O<sub>2</sub> and attenuate cell 239 apoptosis and autophagy, suggesting the anti-oxidant and anti-autophagy functions of 240 Rg1 in SCI. 241 To further study the mechanism of Rg1, we turn our attention to miRNA. MiRNA is 242 important in cell growth, like proliferation and apoptosis (Ameres and Zamore, 2013).

MiR-216a-5p, acknowledged as an oncogenic gene, is involved in tumorigenesis and

244 development of human cancers (Liu et al., 2018). MiR-216a-5p significantly elevated 245 cell activity and reduced apoptosis in H<sub>2</sub>O<sub>2</sub>-caused 16 HBE cells of Asthma, 246 suggesting that miR-216a-5p could regulate H<sub>2</sub>O<sub>2</sub>-caused damage (Chaoyang et al., 247 2019). Besides, of interest, beclin-1 was the latent mark of miR-216-5p, which could 248 inhibit ox-LDL-induced autophagy in human umbilical vein endothelial cells 249 (HUVECs) through modulating levels of intracellular beclin-1 (Menghini et al., 2014). 250 These reports indicate that miR-216a-5p not only elevates activity, suppresses 251 apoptosis, but also inhibits autophagy, suggesting that miR-216a-5p may reduce 252 H<sub>2</sub>O<sub>2</sub>-caused cell damage. Also, functions of miR-216a-5p are similar to those of Rg1. 253 So it is worth to investigate if exerts a relation of Rg1 and miR-216a-5p. For the first 254 time, our study found the regulation relationship between Rg1 and miR-216a-5p. We 255 got that Rg1 could up-regulate miR-216a-5p to attenuate cell injury induced by H<sub>2</sub>O<sub>2</sub>. 256 This finding is a major discovery in SCI research. 257 Furthermore, the biological process is inseparable from the regulation of signal 258 pathways. It has been proved that AMPK/PI3K/AKT pathways are key coordinator 259 protecting cells from oxidative and inflammatory damage (Lv et al., 2017). PI3K was 260 reported to be related to many cellular functions, like proliferation and apoptosis 261 (Rasul et al., 2012). AKT is a key downstream effector of PI3K and exhibits 262 anti-apoptotic effects (Zheng et al., 2015). AMPK is present in metabolically related 263 organs. Cellular metabolism stimulation like cell stress can active it (Zheng et al., 264 2015). Moreover, Lin et al. found that the formation of autophagosomes was 265 accompanied by inhibition of the PI3K/AKT and AMPK signal pathways. This 266 finding indicated that there was negative regulation between these two pathways and 267 autophagy. Therefore, the above findings suggested that there might be positive 268 effects of these two pathways on H<sub>2</sub>O<sub>2</sub>-caused cell damage. It is worth noting that our 269 study verified this suppose. We firstly build up the relation among Rg1, miR-216a-5p 270 and AMPK/PI3K/AKT pathways that Rg1 activated PI3K/AKT and AMPK signal 271 pathways through positively modulating miR-216a-5p to reduce cell damage. This 272 regulation mechanism provides a theoretical basis for attenuating cell injury after SCI. 273 Conclusion 274 Our study firstly reported the underlying effects and mechanism of Rg1 in cell injury 275 of SCI. We demonstrated that Rg1 could up-regulate miR-216a-5p, attenuate cell 276 activity suppression, apoptosis and autophagy, and active PI3K/AKT and AMPK

signal pathways to decrease H<sub>2</sub>O<sub>2</sub>-caused damage in PC-12 cells. Because local cell

278 injury significantly aggravates SCI, we propose that Rg1, an effective biological 279 macromolecular, may supply a novel therapeutic approach for curing SCI. **Conflict of Interest statement** 280 The authors declare that there are no conflicts of interest. 281 282 Acknowledgments 283 None 284 **Fundings** 285 This research did not receive any specific grant from funding agencies in the public, 286 commercial, or not-for-profit sectors. 287 Availability of data and materials 288 The dataset(s) supporting the conclusions of this article is(are) included within the 289 article.

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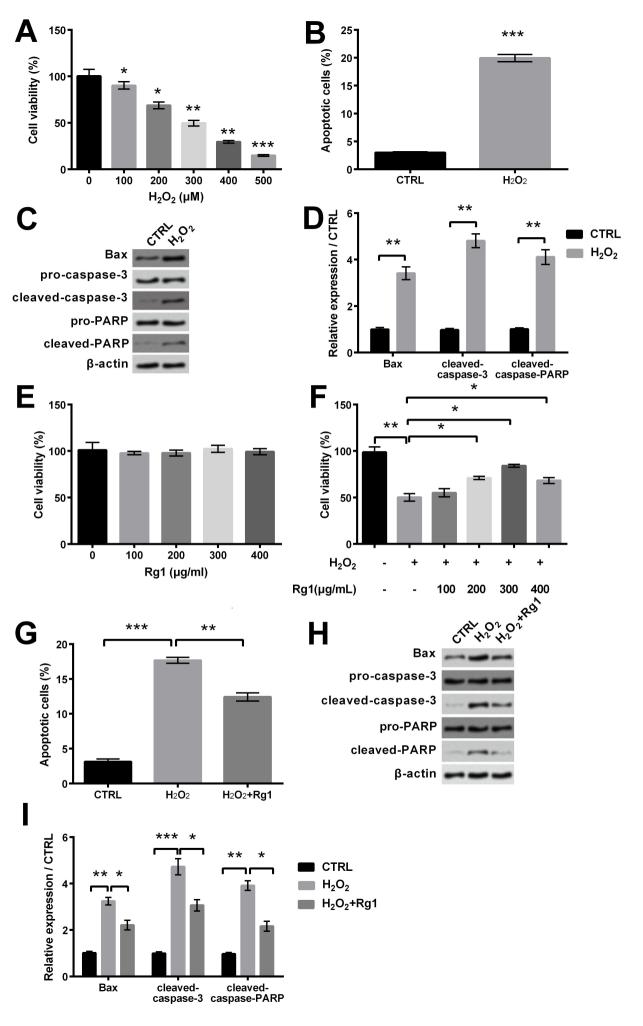
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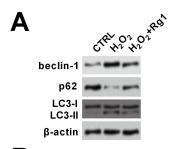
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# Figure legends

- Figure 1 Influence of Rg1 in cell activity and apoptosiscaused by H<sub>2</sub>O<sub>2</sub> in PC-12 cell.
- 393 (A) Cell viability was tested under diverse consistences of  $H_2O_2$  (0, 100, 200, 300,
- 400 and 500  $\mu$ M). 300  $\mu$ M was chose in the following experiments. (B) Cell apoptosis
- treated by H<sub>2</sub>O<sub>2</sub> was tested via flow cytometry. (C) Apoptosis relative elements
- expression was tested via western blot. (D) Level of apoptosis relative factors was
- tested via western blot quantitative. (E) Cell viability was tested via CCK-8 by Rg1.
- 398 (F) Rg1 attenuated H<sub>2</sub>O<sub>2</sub>-induced suppression of cell viability. (G) Rg1 attenuated
- 399 H<sub>2</sub>O<sub>2</sub>-induced cell apoptosis. (H) Expression of apoptosis relative factors was tested
- 400 via western blot. (I) Apoptosis relative elements standards were detected via western
- blot quantitative. \* P < 0.05, \*\* P < 0.01 and \*\*\* P < 0.001 contrast with control and
- the indicated set.
- Figure 2 Influence of Rg1 in autophagy caused by  $H_2O_2$ . (A) Standards of autophagy
- 404 relative factors were tested via western blot. (B) Standards of autophagy relative
- factors were tested via western blot quantitative. \* P < 0.05, \*\* P < 0.01 and \*\*\* P < 0.05
- 406 0.001 contrast with indicated set.
- Figure 3 MiR-216a-5p was up-regulated by Rg1. mRNA standard of miR-216a-5p
- 408 was tested via qRT-PCR. \* P < 0.05 and \*\* P < 0.01 contrast with indicated set.
- 409 Figure 4 Protective effects of Rg1 via up-regulating miR-216a-5p after transfected
- 410 with Rg1 plus miR-216a-5p and relative NC in PC-12 cells. (A) mRNA standard of
- 411 miR-216a-5p was tested via qRT-PCR after miR-216a-5p inhibitor transfection. (B)
- 412 Cell activity was tested via CCK-8. (C) Apoptosis was tested via flow cytometry. (D)
- 413 Expression of apoptosis relative factors was tested via western blot. (E) Level of
- 414 apoptosis relative factors was tested via western blot quantitative. (F) Standards of
- autophagy relative factors were tested via western blot. (G) Standards of autophagy
- 416 relative factors were tested via western blot quantitative. \* P < 0.05, \*\* P < 0.01 and
- 417 \*\*\* P < 0.001 contrast with indicated set.
- 418 **Figure 5** Rg1 elevated PI3K/AKT and AMPK signal pathways through positively
- 419 regulating miR-216a-5p after transfected with Rg1 plus miR-216a-5p and relative NC
- 420 in PC-12 cells. (A) Expression of PI3K/AKT pathway relative factors was tested via

western blot. (B) Standards of relative proteins were tested via western blot quantitative. (C) Expression of AMPK pathway relative factors was tested via western blot. (D) Level of AMPK pathway related proteins were detected via western blot quantitative. \* P < 0.05 and \*\* P < 0.01 contrast with indicated set.





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