Effect and molecular mechanism of mTOR inhibitor rapamycin on temozolomide-induced autophagic death of U251 glioma cells

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Abstract. Glioma is a malignant tumor of the glial tissue that is difficult to excise through surgery, with poor patient prognosis. The use of chemotherapeutic drugs alone to treat glioma following surgery results in a high probability of sequelae, such as tumor recurrence. The present study investigated the effects of a novel treatment combination on glioma cells and determined the molecular mechanisms underlying its action. The effect of temozolomide (TMZ) combined with rapamycin (RAPA) on the TMZ-induced autophagic death of U251 glioma cells was examined. The U251 cell line was treated with TMZ combined with RAPA, and the cell survival rate and half maximal inhibitory concentration (IC50) of TMZ/RAPA was detected using the Cell Counting Kit-8 (CCK-8) assay. Flow cytometry was used to detect changes in cell cycle distribution. The formation of acidic vesicular organelles (AVOs) in the cytoplasm was identified using fluorescence microscopy and quantitatively analyzed. Western blotting was performed to detect the expression levels of autophagy-associated proteins Beclin-1 and microtubule associated protein 1 light chain 3 alpha (MAP1LC3A)‑I and II. RAPA (1.25 nM) combined with 5 µM TMZ markedly inhibited U251 cell growth. RAPA reinforced TMZ-induced autophagic death, reducing the IC50 value of treatment when combined (TMZ alone, 22.5±3.23 µM vs. TMZ and RAPA, 10.35±2.81 µM). Compared with the control group, the proportion of cells in G1/M were markedly increased following treatment with TMZ combined with RAPA. Acridine orange staining demonstrated that TMZ combined with RAPA could markedly enhance the generation of intracellular AVOs compared with TMZ or RAPA alone. In addition, Beclin-1 and LC3-II protein expression was markedly increased compared with the control and single treatment groups (P<0.05). The results of the present study indicate that RAPA reinforces TMZ-induced autophagic death of U251 glioma cells, providing a novel therapeutic combination for the treatment of malignant glioma.

Introduction

Glioma has one of the highest incidence rates of all malignant tumor types and is the most difficult primary cerebral tumor to treat (1). Cerebral malignant glioma, also named high-grade glioma (HGG) (2), is the most common type of primary intracranial malignant tumor and originates in cranial nerve glial cells (3). The relapse rate of HGG is high and results in the generation of metastases; thus, the median survival time of patients with HGG is only 13 months (4). The typical biological characteristic of this type of malignant tumor is invasive growth, including that into the surrounding healthy brain tissue. Due to the location of such tumors within the brain, it is difficult to remove them using surgery alone, resulting in poor prognosis (5,6). Therefore, the conventional treatment for glioma involves a combination of surgery and chemotherapy. However, patients receiving chemotherapy alone following surgery frequently relapse. Further studies are required in order to develop novel and effective treatments for patients with glioma.

Temozolomide (TMZ) is a small lipophilic molecule that can efficiently pass through the blood-brain barrier (7). TMZ is an alkylating agent that can induce base mismatch and DNA abruption, thus inhibiting tumor cell growth and inducing cell death. Therefore, TMZ is regarded as an important chemotherapeutic drug for the treatment of glioma (8). However, Kalkan et al (9) identified that the expression of multidrug resistance-associated proteins, such as P-glycoprotein and O6-methylguanine-DNA methyl transferase (MGMT), was upregulated in malignant glioma cells following chemotherapy, which induced resistance to chemotherapeutic drugs, such as TMZ. In addition, this study identified an association between the expression of these proteins and the relapse rate of malignant glioma.

There may be other mechanisms of drug resistance in malignant glioma. Zhang et al (10) identified that the median survival time of malignant glioma only increased from 14.6 to 21.7 months following MGMT promoter methylation, which
resulted in the down regulation of MGMT. Previous studies have demonstrated that TMZ induces and molecularly regulates the autophagy of glioma cells (11). Based on these results, it has been speculated that TMZ resistance could be associated with autophagy. Therefore, in the present study, the combined effect of TMZ and a specific inhibitor of autophagy was investigated in order to elucidate the molecular mechanisms underlying autophagy in cerebral malignant glioma.

Rapamycin (RAPA) is a macrolide antibiotic, which was first widely used as an immunosuppressant, due to its significant effects on multiple autoimmune diseases. In addition, RAPA is used to prevent rejection following organ transplantation, with few side effects. In previous studies, RAPA demonstrated antitumor activity, in addition to specific autophagy inhibition (12-14). The present study aimed to investigate the combined biological effect of TMZ and RAPA on the proliferation, survival, apoptosis, cell cycle distribution and autophagy of cerebral glioma cells, and the underlying molecular mechanisms involved, in order to develop more effective treatments for patients with glioma.

Materials and methods

Reagents and apparatus. Dulbecco's modified Eagle's medium (DMEM) and fetal bovine serum (FBS) were purchased from Gibco (Thermo Fisher Scientific, Inc., Waltham, MA, USA). Trypsin (0.25%) was purchased from Hangzhou Greengreen (Hangzhou, China). Penicillin and streptomycin were purchased from Wuhan Boster Biological Technology, Ltd. (Wuhan, China). RAPA was purchased from the Beyotime Institute of Biotechnology (Haimen, China). TMZ and acridine orange (AO) were purchased from Sigma-Aldrich (Merck KGaA, Darmstadt, Germany).

Cell culture. The human glioma cell line U251, purchased from the Shanghai Institute of the Chinese Academy of Sciences (Shanghai, China), was cultured in DMEM containing 100 U/ml penicillin and 100 µg/ml streptomycin, supplemented with 10% FBS and incubated at 37°C with 5% CO₂, in a humidified atmosphere. Medium was changed every 1-2 days. When the cells covered 90% of the culture flask, they were digested with trypsin and passedaged.

Detection of U251 cell survival rate using the cell counting kit-8 (CCK-8) assay. Glioma cells in the logarithmic phase of growth were washed with PBS three times and then digested using 0.25% trypsin. DMEM containing 10% FBS was used to achieve a solution of ~1x10⁵ cells/ml. The resulting suspension was added to a 96-well plate (100 µl/well). The plate was incubated at 37°C with 5% CO₂ for 24 h. The cells were subsequently divided into the control groups (solvent and blank) and the experimental groups (TMZ alone, RAPA alone, and TMZ and RAPA). Different concentrations of TMZ (1-50 µM) and/or RAPA (0.625-20 nM) were added to the experimental groups.

Dimethyl sulfoxide (1 µg/ml) was added to the solvent control group and DMEM was added to the blank control group. There were six wells per treatment. Forty-eight hour following the addition of TMZ/RAPA, 10 µl CCK-8 reagent (Sigma-Aldrich; Merck KGaA) was added to each well prior to incubating the plate for a further 2-4 h. An enzyme-linked spectrophotometer was used to detect cell proliferation via measuring the absorbance at 450 nm. The following formula was used to calculate cell viability: (D-B/C-B) x100% (D, absorbance of drug well at 450 nm; B, absorbance of blank control well at 450 nm; C, absorbance of control well at 450 nm).

Detection of cell apoptosis using flow cytometry. A total of 48 h following treatment with drugs as described above, cells (∼5x10⁵ cells/ml) were collected by centrifugation at 200 x g for 5 min in room temperature. Cells were subsequently washed in ice-cold PBS and centrifuged again. Cells were resuspended in 100 µl 5X annexin binding buffer for flow cytometry (Thermo Fisher Scientific Inc.). An additional 5 µl APC annexin V and 5 µl propidium iodide (PI) (both 100 µg/ml) were added to the suspension and incubated at room temperature for 15 min. PI fluorescence excitation was performed using an argon ion laser at a wavelength of 488 nm. Annexin V staining was detected using flow cytometry, as aforementioned. Flow cytometry (Attune Nxt Flow Cytometer Software, version 2.1, Thermo Fisher Scientific Inc.) was used to detect the effect of the drugs on U251 cell apoptosis.

Detection of cell cycle progression using flow cytometry. A total of 48 h following treatment with drugs as described above, the cell culture medium was collected into a streaming dedicated pipe. Cells were washed once with 1 ml PBS and cleaning fluid was added to the pipe. The cells were collected into the pipe following digestion with trypsin. The remaining cells were washed with 1 ml PBS and collected (∼2x10⁵ cells/ml) through centrifugation at 200 x g for 5 min in room temperature. This step was repeated to wash the cells. Subsequently, 300 µl PBS was used to resuspend the cells and the cells were fixed with 700 µl of ice-cold 70% absolute ethanol, which was added dropwise. The cells were immobilized at -20°C for ≥24 h without light. The resultant mixture was centrifuged as described above to remove any stationary liquid. PBS (500 µl) was used to resuspend the cells prior to centrifugation as described above. PI (5 µl) was used for staining at 4°C for 30 min without light. Annexin V staining was detected using flow cytometry, as aforementioned.

Immunofluorescence detection. U251 cells were seeded into a 24-well plate with sterile cover slips in place. The culture medium was discarded when all cells had adhered. Cells were washed three times with PBS and paraformaldehyde (4%) was applied at room temperature for 20 min to immobilize cells. Cells growing on glass coverslips were permeabilized with Triton X-100 solution (0.3%) for 15 min. The cells were washed three times with PBS (5 min/wash). Normal goat serum (10%; Thermo Fisher Scientific Inc.) was used to block non-specific binding and the cells were subsequently incubated at 37°C for 60 min.

Cells were incubated with the primary antibody (1:100, CD81 Monoclonal Antibody (1D6), cat. no. MA1-80820, Thermo Fisher Scientific Inc.). Coverslips were then incubated at 4°C overnight. The cells were subsequently placed at room temperature for 30 min and rinsed three times with 0.01 M of PBS (5 min/wash). Secondary antibody [dilution: 10 µg/ml, donkey anti-Goat IgG (H+L) Cross-Adsorbed Secondary
Antibody, Alexa Fluor 350, catalog no. A-21081, Thermo Fisher Scientific Inc.] was added to the cells cultured on glass coverslips and incubated at 37°C for 30 min. Cells were counterstained with DAPI and incubated at 37°C for a further 10 min. Cells were washed three times with PBS (5 min/wash) and ProLong anti-fade solution (Thermo Fisher Scientific, Inc.) or glycerinum (Thermo Fisher Scientific Inc.) was used for sealing. Laser scanning confocal microscopy was used for observation and image acquisition.

Detection of acidic vesicular organelles (AVOs) in tumor cells using AO staining. The generation of AVOs is a specific process of autophagy. AO is a fluorochrome that can freely cross the cell membrane. AO accumulates in acidic cell components, generating red fluorescence. This red fluorescence appears when acid phosphatase activity increases during autolysosome formation. The present experiment was performed as described in a previous study, which used the FL1 pathway to detect green fluorescence with an emission maximum at 532 nm and the FL3 pathway to detect red fluorescence by using 6HFFRPWL Shower Lens, in order to analyze AVO generation in U251 cells (15). U251 cells in the logarithmic phase of growth (1x10^5 cells/ml), were inoculated into a 24-well plate (1 ml/well). Drugs were added the next day as described above and the cells were cultured for a further 72 h. AO (1 µg/ml) was used at 37°C to stain the cells for 15 min. The distribution of AVOs was subsequently observed under a fluorescence microscope. Flow cytometry (Attune NxT Flow Cytometer Software, version 2.1, Thermo Fisher Scientific Inc.) was used to quantitatively analyze the rate of AVO generation.

Detection of autophagy marker proteins, apoptosis-associated proteins and protein kinase B (Akt)/mammalian target of rapamycin (mTOR) signaling protein pathway expression using western blotting. After 48 h, cells treated with drugs as described above were collected and left on ice for 30 min, and then sonicated for 10 min for lysation. The cells were centrifuged at 2,400 x g, 12,000 rpm at 4˚C for 15 min. Protein concentration was detected using the protein assay kit (Quant-iT™ Protein Assay Kit, Thermo Fisher Scientific). Western blots were semi-quantified with Quantity One (version 4.6.2; Bio-Rad Laboratories, Inc., Hercules, CA, USA) using three different experimental results.

Statistical analysis. Results are presented as the mean ± standard deviation. SPSS software (version 13.0; SPSS, Inc., Chicago, IL, USA) was used for statistical analyses. An unpaired t-test was used to compare differences between groups. P<0.05 was considered to indicate a statistically significant difference.

Effect of TMZ and RAPA on the proliferation and survival of U251 glioma cells. Following treatment with TMZ, U251 cell growth was increased in a dose-dependent manner (Fig. 1A). Growth of the 5 µM TMZ-treated group was markedly inhibited by the fourth day (P<0.05, vs. the untreated cells). A total of 48 h following treatment with TMZ, the half maximal inhibitory rate (IC_{50}) of TMZ was calculated. The IC_{50} of TMZ was 22.5±3.23 M, as illustrated in Fig. 1B. Following treatment with mTOR inhibitor RAPA, the growth of U251 cells was inhibited in a dose-dependent manner (Fig. 1C, P<0.05, vs. the untreated cells). The IC_{50} of RAPA was 12.93±1.36 nM, as illustrated in Fig. 1D.

Effect of TMZ combined with RAPA on the survival of U251 glioma cells. The present study demonstrated that the inhibitory rates of 5 µM TMZ and 1.25 nM RAPA were 21.36 and 19.23%, respectively (Fig. 2). The inhibitory rate of 50 µM TMZ on U251 cell growth was 57.27%, but it showed strong cytotoxicity (data not shown). This is consistent with the results of a previous report (16). In the present study, 5 µM TMZ combined with 1.25 nM RAPA was used in order to reduce the cytotoxic effects of TMZ. The results obtained from the CCK-8 assay demonstrated that the inhibitory effect of TMZ combined with RAPA on cell growth was stronger compared with TMZ alone (P<0.05, in 72 h). The IC_{50} of TMZ and RAPA combined was notably lower compared with TMZ alone (10.35±2.81 µM vs. 22.5±3.23 µM, respectively; Fig. 2).

Effect of TMZ combined with RAPA on the apoptosis of U251 glioma cells. The high blue: low red apoptotic cells shows no significant difference, which indicates that the apoptosis-inhibiting effect of TMZ combined with RAPA was not notable compared with TMZ alone (Fig. 3).

Effect of TMZ combined with RAPA on the cell cycle distribution of U251 glioma cells. U251 cells were treated with 5 µM TMZ and/or 1.25 nM RAPA. Compared with the blank control group, the number of cells in G_{1} decreased and the number of cells in G_{2}/M increased in the groups treated with 5 µM TMZ or 1.25 nM RAPA alone. There was no change in 5 stage cell number (Fig. 4). Following treatment with 5 µM TMZ combined with 1.25 nM RAPA, G_{2}/M stage cell number was notably higher (61.07±2.37) compared with the groups treated with TMZ (31.07±1.39) or RAPA (27.07±1.82) alone (P<0.05, vs. the other groups; Fig. 4).
Effect of TMZ combined with RAPA on the autophagy of U251 glioma cells. In the present study, 5 µM TMZ combined with 1.25 nM RAPA was used to reduce the cytotoxic effects of TMZ. Following a 24 h treatment, the expression of autophagy-related protein LC3-II was notably increased compared with the control group (P<0.05; Fig. 5). There was no significant change in LC3-II expression. The Beclin-1 gene is a congener of yeast autophagy-related genes in mammals and an autophagy regulator gene. Expression of Beclin-1 protein was notably increased following treatment with TMZ combined with RAPA (P<0.05; Fig. 5 A and D).

Detection of U251 AVO generation by AO staining. AO staining is used to analyze autophagy mechanisms at a molecular level. The results from AO staining (Fig. 6) identified no AVO accumulation in the cytoplasm of the control group. The rate of AVO accumulation was 17.2% following treatment with 5 µM TMZ alone, which was significantly increased compared with the control group (0.065%; P<0.05). A similar result was seen following treatment with RAPA alone (AVO accumulation rate, 10.6%). This indicates that treatment with
TMZ or RAPA alone at low concentrations does not effectively induce AVO production in U251 cells. However, the rate of AVO accumulation was notably increased in the cytoplasm of the group treated with TMZ combined with RAPA (44.2%) compared with the groups treated with TMZ (17.2%) or RAPA (10.6%) alone (P<0.05; Fig. 6).
Effect of TMZ combined with RAPA on the autophagy of U251 cells through the mTOR signaling pathway. In order to investigate the molecular mechanisms underlying the induction of autophagy in U251 cells treated with TMZ combined with RAPA, the expression and activation of important components of the mTOR signaling pathway, such as p-mTOR, p-p70S6k and P-4EBP1, was measured following treatment with TMZ alone, RAPA alone or TMZ combined with RAPA. The results demonstrated that the expression of p-mTOR and p-p70S6k significantly decreased in the combination treatment group compared with the groups treated with TMZ or RAPA alone (P<0.05, Fig. 7). A similar trend was seen in the expression of mTOR (P<0.05, Fig. 7A and C). There was no notable change in the expression of p70S6k and P-4EBP1 in the combination treatment group compared with the single treatment groups (P>0.05, Fig. 7E and F). These results indicate that TMZ and RAPA-induced autophagy of U251 cells is associated with the mTOR signaling pathway.

Discussion

The three types of cell death include apoptosis, autophagy and necrosis. The majority of chemotherapeutic drugs induce the apoptosis of tumor cells. Autophagy is a type of programmed cell death (type II). In autophagy, an autophagosome is a monolayer or bilayer membrane that accumulates products to be degraded in the cytoplasm and delivers them to the lysosomes, thus forming autolysosomes. Autolysosomes carry out the digestion and degradation of these products using multiple enzymes, in order to achieve the metabolic needs of the cell and the renewal of specific organelles (17-19). Autophagy serves an important role in maintaining cell homeostasis. In certain tumors, inducing autophagy can significantly inhibit growth and proliferation (20,21). Using AO staining, the present study demonstrated that treatment with TMZ alone induces the generation of AVOs in U251 cells in a dose-dependent manner. The IC₅₀ of TMZ in U251 cells was 22.5±3.23 µM; however, this concentration is highly toxic and known to induce side effects in patients (20). Low concentrations of RAPA (≤2 nM) strengthened TMZ-induced autophagic cell death; thus, the dosage of TMZ was able to be decreased (20). The combined use of RAPA and TMZ increased the generation of AVOs in U251 glioma cells. As the primary indicator of autophagic cell death is the generation of AVOs, these results indicate that the combination of these two drugs may be used to synergistically inhibit glioma cell growth.

The induction of autophagy has been associated with a series of evolutionarily conserved gene products that were first discovered in yeast and named autophagy-related gene (Atg) proteins. In mammals, LC3-I is an autophagic vacuolar protein that is isogenous with Atg 8 (22). During cell autophagy, LC3-I is cleaved to produce LC3-II, which serves an important role in AVO generation and is regarded as a molecular marker of autophagic cell death (23,24). TMZ induces autophagy in other ways (25). Zou et al (26) identified that TMZ inhibited the growth of glioma cells and decreased the number of cells in S and G₂/M, leading to atypical cell apoptosis. The present study identified that the use of TMZ alone to treat U251 cells induces a slight increase in autophagy-related proteins and may induce the apoptosis of glioma cells to a certain extent, which is consistent with a previous study (15).

Figure 7. Effect of TMZ combined with RAPA on the expression of autophagy marker proteins, apoptosis-associated proteins and mTOR signaling pathway proteins in U251 cells. (A) Western blot analysis showing significantly lower levels of p-mTOR and p-p70S6k in the RAPA and TMZ+RAPA groups than in the other three groups. (B-F) The corresponding statistical analyses results for the western blotting are shown on the right sides of the histograms. The data were obtained from three independent experiments. *P<0.05 vs. the untreated cells and TMZ group; TMZ, temozolomide; RAPA, rapamycin; p-, phosphorylated; mTOR, mammalian target of rapamycin; P70S6k, ribosomal protein S6 kinase; 4EBP1, eukaryotic translation initiation factor 4E-binding protein 1.
mTOR is a central regulator of cell growth, proliferation, survival, migration, self-renewal and cell cycle progression (27-29). The downstream effectors of mTOR include eukaryotic translation initiation factor 4E-binding protein 1 and 70S6K. These two downstream effectors control expression of cyclin D1 and cyclin-dependent kinase in eukaryotes (30-32). Thoury et al (33) combined RAPA derivative RAD001 with AE788 (an epidermal growth factor receptor/vascular endothelial growth factor receptor double tyrosine kinase inhibitor) to treat D54MG glioma in a murine model. This combination was demonstrated to significantly inhibit tumor cell growth and proliferation, and the median survival time of the mice was significantly longer in the combined treatment group compared with the groups treated with AE788 or RAD001 alone. Wang et al (34) demonstrated that combining RAPA with an inhibitor of the mTOR upstream regulator molecules phosphoinositide 3-kinase and Akt significantly sensitized cells to RAPA-mediated autophagy and resulted in radiosensitization.

The present study demonstrated that autophagy-associated protein concentrations in U251 cells markedly increased following treatment with TMZ combined with RAPA. Furthermore, the number of G1/M stage cells significantly increased in the combined treatment group compared with the single treatment groups. AO staining identified that the amount of AVOs increased in cells treated with TMZ combined with RAPA, indicating that the combination promotes autophagic cell death. These results suggest that RAPA may be combined with other drugs or therapies to effectively inhibit the growth and proliferation of glioma cells.

The results of the current study demonstrated that RAPA combined with TMZ induces the expression of Beclin-1 and LC3-II in U251 cells. These proteins serve important roles in autophagy. Beclin-1 is upregulated by certain autophagy-inducing agents, such as the histone deacetylase inhibitor suberoylanilide hydroxamic acid-d5. The findings of the present study are consistent with those reported by Ye et al (35), in which small interfering RNA knockdown of Beclin-1 expression resulted in decreased RAPA-mediated autophagy and resulted in radiosensitization.

To conclude, the results of the present study indicate that a low concentration of RAPA significantly strengthens TMZ-induced autophagic death of U251 cells, thus providing a novel therapeutic approach for the treatment of patients with malignant glioma.

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