Dual PI3K/mTOR Inhibitor, XL765, suppresses glioblastoma growth by inducing ER stress-dependent apoptosis

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Hang Zhao
Guangyong Chen
Huaxin Liang

Department of Neurosurgery, China-Japan Union Hospital of Jilin University, Changchun, Jilin 130033, People's Republic of China

Background: Deregulated phosphoinositide 3-kinase (PI3K)/mTOR signaling commonly exists in glioblastoma (GBM), making this axis an attractive target for therapeutic manipulation. A recent dual inhibitor of PI3K/mTOR pathway, XL765, exhibited an attractive suppression effect on GBM tumor growth. However, the exact functional mechanisms of tumor suppression mediated by XL765 have not yet been fully characterized.

Purpose: In this study, we took efforts to assess the effects of PI3K/mTOR blockade by XL765 on GBM growth *in vitro* and *in vivo*.

Methods: We analyzed the cytotoxicity of XL765 in three different GBM cell lines, A172, U87MG, and T98G, by using Hoechst 33258 (Invitrogen), Annexin V/propidium iodide (PI), as well as Cell Counting Kit -8 (CCK-8) assay. We also used A172 xenograft model to study the effect of XL765 *in vivo*.

Results: We found that XL765 inhibits GBM viability with a wide range of potencies. Importantly, XL765 suppressed GBM cell growth by inducing endoplasmic reticulum (ER) stress dependent apoptosis. The activation of CHOP/DR5 pathway by XL765 induced ER stress is responsible for the induction of apoptosis. Moreover, the inhibition of mTOR signal by XL765 is the major source of ER stress, rather than inhibition of PI3K. At last, we demonstrated that combination of XL765 with GBM chemotherapeutic drug, temozolomide (TMZ), can achieved better therapy effect *in vitro* and *in vivo*.

Conclusion: Overall, our data show that targeting PI3K/mTOR by XL765 is a promising therapeutic strategy to relieve tumor burden in GBM patients.

Keywords: XL765, glioblastoma, apoptosis, ER stress

Introduction

Glioma, as the most frequent primary brain tumor, accounts for 81% of malignant tumors in central nervous system, which has been increasing as a severe threat to human life.¹ Currently, the standard therapy strategy is surgery, radiotherapy, and adjuvant temozolomide (TMZ) chemotherapy.² However, the glioblastoma (GBM) tumors usually possess mixed cytological subtypes in phenotype, multiple mutations and gene amplifications in genotype, which make any conventional or pathway-specific therapy less effective.³⁻⁵ With the development of molecular biology, it has been identified several molecular pathway abnormal expression, which are related to the pathogenesis and prognosis of GBM, including *TP53* mutations, *TERT* promoter region mutations, *EGFR* amplification, *PTEN* deletions, and 1p/19q combined deletions.⁶

Correspondence: Huaxin Liang
Department of Neurosurgery, China-Japan Union Hospital of Jilin University, No. 126, Xiantai Street, Changchun, Jilin 130033, People's Republic of China
Email huaxinliang2010@jlu.edu.cn

As its potential function in cancer, the phosphoinositide-3-kinase (PI3K)/mTOR pathway signaling axis is receiving more and more attention in recent years.⁷ Database analysis by the Cancer Genome Atlas shows that the PI3K/mTOR pathway is activated in around 88% in GBM patients.^{6,8} This critical pathway acts as upstream effector of multiple signal pathways, and in turn stimulates numerous downstream effectors, which are involved in cellular survival, growth, protein synthesis, motility, and other functions of pro-tumorigenic effects.⁷ The PI3K pathway is manipulated by different signaling cascades, including loss function of the phosphatase and tensin homologue (PTEN) protein, amplification and/or mutation of the EGFR, which occurs in 40% and 50% of GBM cases, respectively.⁸ These perspectives led to the development of novel drugs targeting single (eg, mTORC1 inhibitors) or multiple (eg, dual PI3K/mTOR inhibitors) components of this pathway.⁹ Currently, several drugs targeting this signaling pathway are in Phase I and II clinical trials, either in combination with other chemotherapeutic agents such as TMZ, or as single agents.^{10–12} Although first-generation PI3K pathway inhibitors have effects on the most downstream node of mTOR, an enhanced PI3K signal by feedback loops was observed, which has led to the development of second-generation drugs.^{13–15}

XL765 (SAR245409) is a potent class I dual inhibitor of PI3Ks and mTOR. In cellular assays, treatment with XL765 suppresses phosphorylation of PI3K and mTOR downstream effectors in multiple tumor cell lines, such as AKT and ribosomal protein S6 (S6RP).¹⁶ In multiple tumor cell lines, XL765 exhibits wide range of inhibitory potencies.¹⁴ It was also reported that XL765 treatment in GBM tumor in mice model is associated with enhanced survival,¹⁷ but without clear mechanisms. The goal of the current study was to reveal the underlying mechanism of XL765 suppression of GBM tumor growth, which will provide more insights about PI3K/mTOR axis as a potential target for anti-GBM therapy.

Materials and methods

Cell lines and reagents

A172, U-87MG, and T98G GBM cell lines were obtained from ATCC (Manassas, VA, USA). The cells were cultured with DMEM consisting of Ham's F12 medium (1:1) (Invitrogen) was mixed with 10% FBS (HyClone, Logan, UT, USA), 100 units/mL penicillin, and 100 µg/mL streptomycin. All of the cells were cultured at 37°C and 5% CO₂.

Chemicals, including TMZ, XL765, Z-VAD-FMK (z-VAD), salubrinol were purchased from Sigma-Aldrich (Shanghai, People's Republic of China).

Cell survival and viability assay

Cell survival was determined by using Cell Counting Kit-8 (CCK-8) (Sigma-Aldrich). A172, U-87MG, and T98G cells were seeded at 2×10^3 – 4×10^3 cells/well in 96-well plates. On the following day, cells were treated with different concentrations of drugs. After 24 and 48 hrs, viable cells were quantified by using a CCK-8 assay according to the manufacturer's protocol.

The cell viability was determined by crystal violet staining. Briefly, 1×10^5 A172, U-87MG, and T98G cells were seed in 12-well plates, and treated with different concentrations of drugs for 24 hrs. The attached cells were washed with PBS and stained with a 0.05% crystal violet solution (containing 3.7% paraformaldehyde prepared in distilled water) at room temperature.

Apoptosis analysis

The apoptosis of GBM cells was analyzed by nuclear staining with Hoechst 33,258 (Invitrogen), or Annexin V/propidium iodide (Invitrogen) followed by flow cytometry as described previously.¹⁸ For the nuclear Hoechst 33,258 staining, the treated GBM cells were stained with Hoechst 33,258 (3.7% formaldehyde, 0.5% Nonidet P-40, and 10 µg/mL Hoechst 33,258 (Invitrogen)), and the condensed chromatin and micro-nucleation was counted under microscopic visualization.

Intracellular Ca²⁺ detection

The levels of intracellular Ca²⁺ were determined using Fluo-3 AM (S1056, Beyotime, Shanghai, People's Republic of China) staining followed by flow cytometry analysis. Human GBM cell A172 and U87 were treated with 10 µM XL765 for 24 hrs. After treatment, cells were stained with Fluoro-3AM for 30 mins and collected for flow cytometry analysis.

Transfection of small-interfering RNA (siRNA) and plasmid

Transfection of siRNA and plasmid was conducted using Lipofectamine[®] 2000 (Invitrogen) as described by the manufacturer. Active Akt plasmid (myr Akt delta PH, #10,841) and S6K expressing plasmid (pDONR223-RPS6KB1, #23,686) were purchased from Addgene (Cambridge, MA, USA). siRNA duplexes were synthesized by Genepharma (Shanghai, People's Republic of China) and included: DR5 (AAGACCCUUGUGCUCGUU GUC), CHOP (GCACAGCUAGCUGAAGAGA).

Western blot analysis

RIPA lysis buffer supplemented with protease inhibitor cocktail (Complete Mini; Roche, Basel, Switzerland) was used for Whole cell lysates preparation. After sonicated for 5 mins, the lysates were centrifuged at $12,000\times g$ for 20 mins at 4°C , and the supernatants were collected. Western blot was performed as previously described.¹⁶ The following primary antibodies were used: Tubulin (Sigma), cleaved caspase-3, p-AKT, total AKT, p-S6K, total S6K, PERK (Cell Signaling, Danvers, MA, USA), CHOP (EMD Millipore, Burlington, MA, USA), DR5 (Santa Cruz, Shanghai, People's Republic of China).

Quantitative reverse transcription-polymerase chain reaction (qRT-PCR)

RNeasy Mini Kit (Qiagen, Hilden, Germany) was used to extract total RNA from the cultured cells in accordance with the manufacturer's protocol. One microgram of total RNA was used to generate cDNA using Superscript II reverse transcriptase (Invitrogen, Carlsbad, CA, USA). qRT-PCR was performed on the StepOnePlus Real-Time PCR System (Applied Biosystems, Foster City, CA, USA) using SYBR Green qPCR Master Mix (ThermoFisher Scientific, Shanghai, People's Republic of China). Values were normalized to the expression of *ACTIN* mRNA. Primers used in this study are: DR5: 5'-AAGACCCTTGTGCTCGTTGT-3' and 5'-AGG TGGACACAATCCCTCTG-3'; ACTIN: 5'-AGAGCTACG AGCTGCCTGAC-3' and 5'-AGCACTGTGTTGGCGTAC AG-3'.

Xenograft experiment

The animal studies were approved by the Institutional Animal Ethics Committee of Jilin University and experiments were performed in accordance with the Animal Ethics guidelines of Jilin University. For in vivo tumorigenicity assays, severely immuno-compromised female BALB/C nude mice were used. Briefly, 1×10^6 of A172 cells in 0.1 mL ice-cold Hanks Balanced Salt Solution were implanted subcutaneously into the right flank of female nude mice (N=5). Seven days after cell injection, the mice were received intraperitoneal (IP) injection of TMZ (25 mg/kg) every day, and/or oral gavage of XL765 (100 mg/kg) every 2 days. PBS injection was used as a negative control (control). The drugs were administrated for total of 2 weeks. The tumor volume was measured each other day, and determined using the formula: $4/3\pi (\sqrt{\text{major axis}/2\times \text{minor axis}/2})$.

Statistical analysis

Data are presented as means $\pm$ standard error of the mean. Two-tailed Student's *t* test was used for statistical calculation. Differences were considered statistically significant at $P<0.05$.

Results

XL765 suppressed GBM cells viability

To investigate the effect of XL765 on the GBM cell lines, different human GBM cell lines A172, U87, and T98G were treated with different concentrations of XL765 (0, 1, 2.5, 5, 10, and 20 μM) for 24 and 48 hrs. We found that 5, 10, and 20 μM XL765 could inhibit the growth of human GBM cell lines in a time- and dose-dependent manner (Figure 1A). The IC₅₀ values of these GBM cells were shown in Figure (1B). A cell viability assay by plate crystal violet staining was performed to further confirm the effect of XL765 on GBM cell viability, where the survival of GBM cells was suppressed by XL765 treatment in a dose-dependent manner (Figure 1C). The effects of XL765 on the PI3K/mTOR signaling pathway were then examined by Western immunoblot of AKT and S6K. XL765 was found to induce a marked dose-dependent decrease in the phosphorylation of AKT and S6K (Figure 1D). Induction of apoptosis is the major mode of cytotoxicity for many anti-cancer drugs.¹⁹ A flow cytometry assay and Hoechst 33,258 nuclei staining were performed to investigate whether XL765 could induce GBM cell apoptosis in vitro. The percentage of apoptotic A172 cells was higher after XL765 treatment compared to the control groups (Figure 1E). The apoptosis rates increased after higher XL765 treatment concentrations (Figure 1E). Furthermore, Hoechst 33,258 staining also revealed that XL765 can induce nuclei fragmentation in multiple GBM cells, including A172, U87, and T98G cells (Figure 1F). Inhibition of apoptosis by pan-caspases inhibitor, z-VAD, compromised the nuclei fragmentation induced by XL765 in these 3 cell lines (Figure 1G). Thus, our data demonstrated that XL765 suppressed GBM cell viability by inducing apoptosis in a time- and dose-dependent manner.

XL765 induces endoplasmic reticulum (ER) stress-related apoptosis in GBM cells

It was reported that ER stress was negatively regulated by AKT/TSC/mTOR pathway,²⁰ and is a major cellular stress to induce apoptosis.²¹ To investigate this phenomenon, calcium disorder, an important marker of the ER

Figure 1 XL765 suppressed GBM cell viability by inducing apoptosis. **(A)** The CCK8 assay analysis for the survival of A172, U87, and T98G cells treated with different doses of XL765 for 24 and 48 hrs. **(B)** The 24 and 48 hrs IC₅₀ values of XL765 in A172, U87, and T98G cells. **(C)** The cell viability of A172, U87, and T98G cells treated with different concentrations of XL765 for 24 hrs. **(D)** The expression of p-AKT, total AKT, p-S6K, total S6K in A172 cells treated with different concentrations of XL765 for 24 hrs. **(E)** The Annexin-V/PI staining of A172 cells treated with different concentrations of XL765 for 24 hrs. Left, the representative flow pictures, right, the bar graph for Annexin V positive cells. **(F)** The Hoechst-333,255 staining for apoptosis of A172, U87, and T98G cells treated with different concentrations of XL765 for 24 hrs. **(G)** The Hoechst-333,255 staining for apoptosis in A172, U87, and T98G cells treated with 10 μM XL765 with or without 10 μM z-VAD combination for 24 hrs. N=3 for each experiment.

stress, was analyzed by Fluo-3 AM cytoplasmic calcium staining followed by flow cytometry assay. We found that a significantly elevated cytoplasmic calcium level was detected in XL765 treatment GBM cell groups (Figure 2A). To further clarify this, the expression levels of ER stress marker PERK, p-Eif2α, and CHOP were assessed. We found that XL765 treatment could significantly increase the expression of PERK, p-Eif2α, and

CHOP in a dose-dependent manner (Figure 2B). Inhibition of ER stress by Eif2α inhibitor, salubrinal, suppressed the phosphorylation of Eif2α and induction of CHOP by XL765 (Figure 2C). Suppression of ER stress by salubrinal also compromised the XL765-induced apoptosis in A172 cells, as showed by cleaved caspase-3 expression and fragmented nuclei staining (Figure 2C and D). Moreover, knockdown of CHOP,

Figure 2 XL765-induced ER stress-dependent apoptosis in GBM cells. **(A)** A172 and U87MG cells were treated with 10 μ M XL765 for 24 hrs, then cells were stained with Fluo-3AM for 30 mins, cytosolic calcium was monitored by flow cytometry. Left, the representative flow pictures, right, the bar graph for cytosolic calcium positive cells. **(B)** The expression of indicated proteins in A172 cells treated with different concentrations of XL765 for 24 hrs. **(C)** A172 cells were treated with 10 μ M XL765 with or without pretreatment of 10 μ M salubrinal for 24 hrs. The expression of indicated proteins was analyzed by Western blot. **(D)** The Hoechst-333,255 staining for apoptosis of A172 treated as in **(C)**. **(E)** A172 cells transfected with control or CHOP siRNA were treated with 10 μ M XL765 for 24 hrs. The expression of indicated proteins was analyzed by Western blot. **(F)** The Hoechst-333,255 staining for apoptosis of A172 treated as in **(E)** N=3 for each experiment. **, $P < 0.01$.

Abbreviations: C, control; X, XL765; S, salubrinal; XS, XL765+ salubrinal.

the ER stress sensor that has a protective effect in ER stress-related apoptosis, which significantly suppressed XL765-induced GBM apoptosis (Figure 2E and F). These findings demonstrated that XL765 could induce ER stress-related apoptosis in human GBM cell lines.

DR5 induction mediated the GBM apoptosis induced by XL765

Death receptor 5 (DR5) plays an critical role in ER stress-induced apoptosis,¹⁸ we therefore investigated its regulation in response to XL765 treatment. We found that XL765 treatment induced DR5 expression in a dose-dependent manner (Figure 3A). Accordingly, the mRNA level of DR5 was also induced

by XL765 in A172 cells with a dose-dependent manner (Figure 3B). Suppression of ER stress by salubrinal, depletion of CHOP by siRNA suppressed the induction of DR5 in protein and mRNA level, further confirmed that DR5 is the downstream effector of ER stress induced by XL765 (Figure 3C and D). To further evaluate the role of DR5 in XL765-induced apoptosis, we applied the siRNA to depletion the expression of DR5. Our results showed that DR5 knockdown by siRNA compromised the cleavage of caspase-3 (Figure 3E), and abrogated the apoptosis induced by XL765 in A172 cells (Figure 3F). Collectively, our results suggested that DR5 is the downstream effector of XL765-induced ER stress to promote GBM cell apoptosis.

Figure 3 DR5 mediated the apoptosis induced by XL765. **(A)** The protein expression of DR5 in A172 cells treated with different concentrations of XL765 for 24 hrs. **(B)** The mRNA expression of DR5 in A172 cells treated with different concentrations of XL765 for 24 hrs. **(C)** A172 cells were treated with 10 μM XL765 with or without pretreatment of 10 μM salubrinal (Sal) for 24 hrs. The mRNA expression of DR5 was analyzed by RT-PCR. **(D)** A172 cells transfected with control or CHOP siRNA were treated with 10 μM XL765 for 24 hrs. The mRNA expression of DR5 was analyzed by RT-PCR. **(E)** A172 cells transfected with control or DR5 siRNA were treated with 10 μM XL765 for 24 hrs. The expression of indicated proteins was analyzed by Western blot. **(F)** The Hoechst-333,255 staining for apoptosis of A172 treated as in **(E)** N=3 for each experiment. **, $P<0.01$.

Inhibition of mTOR by XL765 contributes to ER stress-associated apoptosis

We next addressed the mechanism of ER stress raised by XL765 treatment in GBM cells. As we found XL765 treatment inhibited S6K and AKT activation, we further investigated whether inhibition of PI3K or mTOR can induce ER stress in GBM. Transfection of constitutively active AKT in A172 cells suppressed XL765 inhibition of phosphorylation of AKT (Figure 4A). However, restore of AKT activity has mild effect on suppressing the XL765-induced ER stress, as indicated by expression CHOP (Figure 4A). Accordingly, enhanced AKT activity did not significantly suppress XL765-induced apoptosis,

caspase-3 activation, and DR5 induction in mRNA level (Figure 4A–C), suggesting that inhibition of PI3K by XL765 did not contribute to ER stress-induced apoptosis. It was reported that PI3K is the upstream effector of mTOR,²² however, we found restore AKT activity only slightly reduced the inhibition of mTOR by XL765, also suggesting that XL765 inhibits PI3K-independent mTOR signaling.¹⁶ Inhibition of mTOR is also an important source of ER stress,²³ we therefore tested whether XL765-induced ER stress by mTOR inhibition. Transfection of constitutively active S6K in A172 cells suppressed XL765 inhibition of phosphorylation of S6K, but did not affect the inhibition of phosphorylation of

Figure 4 XL765-induced ER stress in GBM by suppressing mTOR activity. (A–C) A172 cells transfected with control or active AKT plasmids were treated with 10 μ M XL765 for 24 hrs. (A) The expression of indicated proteins was analyzed by Western blot. (B) The apoptosis of A172 cells. (C) The mRNA expression of DR5. D–F. A172 cells transfected with control or active S6K plasmids were treated with 10 μ M XL765 for 24 hrs. (D) The expression of indicated proteins was analyzed by Western blot. (E) The apoptosis of A172 cells. (F) The mRNA expression of DR5. N=3 for each experiment. N, $P>0.05$; *, $P<0.05$; **, $P<0.01$.

AKT (Figure 4D). As predicted, enhanced S6K expression in A172 cells suppressed XL765-induced CHOP expression, cleavage of caspase-3, cell apoptosis, as well as DR5 expression in mRNA level (Figure 4D–F). Therefore, these data collectively indicated that XL765-induced ER stress-related apoptosis in GBM by suppression of mTOR activity.

XL765 enhancing the killing effect of TMZ in vitro and in vivo

We found that XL765 treatment suppressed cell viability by apoptosis in different GBM cell lines (Figure 1A–C), which has different responses to TMZ treatment, a major chemotherapy drug in treating GBM. We therefore tested whether XL765 can benefit the TMZ-based chemotherapy of GBM. We found that TMZ treatment induced higher apoptosis and caspase-3 activation in TMZ sensitive cells, U87 and A172 cells (Figure 5A and B). In contrast, TMZ treatment did not obviously induce apoptosis in TMZ resistant cells, T89G cells (Figure 5C). However, the combination of XL765 and TMZ can maximize the killing effect of TMZ in different GBM cells (Figure 5A–C),

suggesting that XL765 combination can overcome the TMZ resistance, and achieve better therapy results. To further confirm the therapy effect of XL765 in GBM, we tested the effect of combination treatment of TMZ and XL765 on the growth of A172 xenograft in vivo. We found that XL765 single treatment suppressed the tumor growth, which was comparable with the effect of TMZ single treatment (Figure 5D and E). And combination of TMZ with XL765 treatment dramatically had more suppression effects on tumor growth, compared with the TMZ or XL765 single treatment (Figure 5D and E). Therefore, our data verified that XL765 has antitumor activity in vivo, and combination with TMZ treatment could enhance its killing effect in vitro and in vivo.

Discussion

The PI3K/mTOR pathway, as one of the most important signaling pathways in human GBM, has multiple amplification, mutation, or deletion in its key pathway regulatory components.⁸ Multiple evidence have shown that aberrant PI3K/mTOR pathway signaling has a critical role in many aggressive malignancy tumors,²⁴ supporting the development of drugs targeting this axis. Studies here demonstrated

Figure 5 XL765 enhanced the killing effect of TMZ in vitro and in vivo. (A–C) A172 (A) U87MG (B) and T98G (C) cells treated with 10 μ M XL765 and/or 15 μ M TMZ for 24 hrs. Upper, the apoptosis was analyzed by Hoechst-33,258 staining, lower, the expression of cleaved caspase-3. (D) Tumor growth of A172 cells xenografted nude mice (n=5) received TMZ (25 mg/kg) and/or XL765 (100 mg/kg) treatment. *, $P < 0.05$; **, $P < 0.01$. (E) The representative tumors in each treatment group.

that dual PI3K/mTOR blockade by XL765 markedly suppressed the growth of human GBM in vitro and in vivo. In this study, we found that inhibition of PI3K/mTOR by XL765-induced apoptosis in GBM cells, which was dependent on the ER stress-mediated DR5 activation. The ER stress induced by XL765 was more related to mTOR inhibition rather than PI3K inhibition. Furthermore, our studies investigated the effect of XL765 by using oral administration of clinically relevant dosing regimens (100 mg/kg every 2 days),^{14,17} and combination treatment with the standard care of TMZ in GBM xenograft mice model. Our in vivo data suggested that combination of XL765 with chemotherapy has a better anti-cancer effect, which provided a novel therapy combination strategy for GBM patients.

It has been reported that XL765 exhibits a wide range of anti-tumor activity in multiple cancers.¹⁴ There is a trend that enhanced sensitivity to XL765 in the cells exhibiting mutations in *PI3K* pathway, which is consistent with previous reports about PI3K inhibitors GDC-0941 and CH5132799.^{25,26} However, it is still unclear whether *PI3K* mutations or *PTEN* deficiency can be the general predictive marker for clinical responsiveness to PI3K pathway inhibitors.²⁷ Large-scale molecular profiling of

tumors in the ongoing XL765 clinical studies is being performed to further address this question. It was reported that oral administration of XL765 resulted in substantial tumor growth suppression in multiple xenograft tumor models, suggesting that XL765 may have broad utility in tumors.¹⁶ In contrast, *PTEN*-absent cell lines had multiple sensitivities to XL765, since some are very sensitive but some are intractable.¹⁶ In our study, the three human GBM cells, A172, U87-MG, and T98G, are *PTEN* deficient or mutant, and showed high sensitivity to XL765-induced apoptosis, which suggested that XL765 might have favorable therapy outcomes in *PTEN* mutated tumors.⁶

It has been reported that XL765 could promote apoptosis through genetic and pharmacologic autophagy blockade in malignant peripheral nerve sheath tumors.²² Meanwhile, Radoul et al described the phenomenon that XL765 treatment either alone or in combination with TMZ significantly reduced the tumor burden in xenografted mice,¹⁷ but without further mechanism discovered. Here in this study, we first discovered that XL765 could block GBM cell survival by inducing apoptosis in a time- and dose-dependent manner. Further research on these XL765-treated GBM cell lines detected an increasing expression of ER stress markers after XL765 treatment, such

as PERK, EIF2 α , and CHOP. PERK is pivotal for cell adaptation to ER stress, promoting either survival or apoptosis.²⁸ The activation of PERK/EIF2 α axis leads to the expression of CHOP, a pro-apoptotic protein in the downstream of PERK/EIF2 α axis.²⁹ Our results showed that pharmacological inhibition of ER stress by EIF2 α inhibitor resulted into abandon of cell apoptosis, suggesting the activation apoptotic pathway by of PERK/EIF2 α pathway.

As to the induction of ER stress by XL765, we found that the ER stress is more dependent on the mTOR inhibition. mTOR and unfolded protein response have been shown to act coordinately in a number of biological processes.³⁰ mTOR inhibitors were able to suppress the tumor growth by trigger ER stress-dependent apoptosis.³¹ Up to date, several pre-clinical studies using rapamycin or its derivatives have yielded favorable outcomes. However, accumulating data from other solid malignancies suggest that the clinical effects of mTOR inhibitors only have transient tumor stabilization, and there is tumor re-growth during and/or after treatment discontinuation.^{32,33} XL765, as a novel developed small molecule, inhibits both class I PI3Ks and mTOR. Although XL765 seems to be more potent against PI3K, it also inhibits mTOR-dependent phosphorylation events and PI3K-independent in tumor cells.¹⁶ Different from previous described mTOR inhibitor, XL765 not only suppressed the GBM cell growth, but also demonstrated obvious killing effect on GBM cells. However, the clinical usage of XL765 still needs to be further investigated.

XL765 is currently in Phase I and II clinical studies, and used as a single agent or in combination with other chemotherapy agents in various cancer patients.^{34,35} Consistent with previous observation,³⁶ XL765 has demonstrated significant efficacy in a GBM xenograft model, both as a single treatment or combination with TMZ. Taken together, these data support the ongoing clinical function of XL765 for the treatment of GBM cancer.

Disclosure

The authors declare no conflicts of interest in this work.

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