Oleanolic Acid Inhibits the Proliferation and Invasion of U251 Glioma Cells and Promotes their Apoptosis through the IKK- β , MAPK3, and MAPK4 Signaling Pathway

Jinxiang Huang

Changzheng Hospital: Shanghai Changzheng Hospital

Shengnan Lin

Sanming First Hospital and First Hospital of Sanming Affiliated to Fujian Medical University

Feng Zhu

Sanming First Hospital and First Hospital of Sanming Affiliated to Fujian Medical University

Dengsheng Chen

Sanming First Hospital and First Hospital of Sanming Affiliated to Fujian Medical University

Fang Huang

Sanming First Hospital and First Hospital of Sanming Affiliated to Fujian Medical University

Junling xu (✉ xlning@fjmu.edu.cn)

Sanming First Hospital and First Hospital of Sanming Affiliated to Fujian Medical University

<https://orcid.org/0000-0003-2647-6130>

Research

Keywords: Glioma, Oleanolic acid, IKK- β , MAPK3, MAPK4

Posted Date: September 20th, 2021

DOI: <https://doi.org/10.21203/rs.3.rs-877888/v1>

License: This work is licensed under a Creative Commons Attribution 4.0 International License.

[Read Full License](#)

Abstract

Object: To investigate the effects of Oleanolic acid (OA) on proliferation, apoptosis, migration, and invasion of human glioma cell U251, as well as IKK- β and MAPK signaling pathways.

Methods: The binding of OA to IKK- β and MAPK signaling pathway essential proteins IKK- β , MAPK3, and MAPK4 was analyzed by molecular docking technique. U251 cells were treated with different concentrations of OA. The proliferation and apoptosis rates of U251 cells were detected by CCK-8 assay, MTT assay, cell cloning assay, and Annexin-V FITC/PI double staining assay. Transwell chamber assay was used to detect migration and invasion of U251 cells. Western blot was used to detect and analyze the expression levels of CYP17A1, IKK- β , PTGS2 and MAPK3/4 protein in U251 cells after OA treatment. Finally, the transcriptome sequencing method was used to detect the differentially expressed genes in the two groups, and the GO and KEGG enrichment analysis were performed.

Results: The results of molecular docking showed that OA could stably bind to IKK- β , MAPK3, and MAPK4 proteins. OA could not only effectively inhibit the proliferation and induce apoptosis of U251 cells ($P < 0.05$), but also significantly inhibit the invasion of U251 cells ($P < 0.005$). Western blot assay confirmed that OA could dramatically inhibit the protein expression levels of CYP17A1, IKK- β , PTGS2, MAPK3, and MAPK4 in U251 cells ($P < 0.01$). A total of 446 significantly differentially expressed genes were detected in transcriptome sequencing, of which 96 were up-regulated genes and 350 were down-regulated genes. These genes are mainly involved in processes such as inflammation, metabolism, immunity, and regulation of cell growth.

Conclusions: OA may inhibit the proliferation, migration, and invasion of glioma U251 cells by binding key molecules of the IKK- β signaling pathway and essential target proteins of MAPK3 and MAPK4 in the MAPK signaling pathway.

Background

Glioma is currently the most aggressive and fatal human intracranial primary malignant brain tumor ^[1]. Because of the specific and remote location of these tumors, removal of glioma is difficult, and thus glioma is associated with high morbidity and mortality rates ^[2]. The clinical treatment of glioma includes surgery, radiotherapy, and chemotherapy drugs, but the prognosis in patients remains unsatisfactory ^[3] and the 5-year survival rate of glioma is less than 10%. Therefore, identifying anti-tumor drugs with good curative effects, minor side effects, and high safety is critical. At present, targeted therapy has become a new research hotspot. Compared with traditional medicines, small molecule targeted drugs have increased efficacy in targeting tumors and show advantages such as specificity, accurate targeting, good tolerance, mild toxicity, and passing the blood-brain barrier ^[4].

Oleanolic acid (OA) is a type of natural pentacyclic triterpenoid compound that is present in many plants such as *Hedyotis officinalis*, *Ligustrum lucidum*, papaya, loquat leaves, olive, and others. OA has many

pharmacological effects, such as protecting the liver, lowering lipids, lowering glucose, and enhancing immunity. Studies have shown that OA can induce apoptosis in a variety of tumor cells, such as lung cancer, pancreatic cancer ^[5], cervical cancer ^[6], and ovarian cancer cells ^[7], and many other types of tumor cells that have anti-proliferation effect. Currently, OA is widely used in clinical medicine for the treatment of acute and chronic hepatitis and as an anti-cancer adjuvant drug ^[8].

In this study, we investigated the effects of OA on proliferation, migration, and invasion of glioma cells and the IKK- β , MAPK3, and MAPK4 proteins, with the aim of providing experimental basis for the clinical treatment of glioma patients with OA.

1. Materials And Methods

1.1 Materials

Oleanolic acid (Lot: AF9071301), its molecular formula is C₃₀H₄₈O₃, molecular weight is 456.70, purchased from Chengdu Alfa Biotechnology Co., Ltd. Purity of the compound used in the present study was shown to be higher than 98% by HPLC; The human glioma cell line U251 was purchased from the Cell Bank of Shanghai Academy of Sciences; DMEM was purchased from Hyclone, USA; fetal bovine serum was from GIBCO. The CCK-8 kit, PBS, 0.25% trypsin digestion solution, trypsin EDTA digestion solution, penicillin-streptomycin, and electrophoresis buffer were all purchased from Solarbio. The RIPA tissue cell rapid lysate and BCA protein quantification kits were purchased from Beyotime. HRP-labeled secondary antibody and the Annexin V-FITC cell apoptosis detection kit were purchased from Biyuntian Biotechnology Co., Ltd. Formaldehyde was purchased from Sinopharm Chemical Reagent Co., Ltd. Matrigel was purchased from Corning Company. IKK β and MAPK4 antibodies were purchased from Proteintech; MAPK3 antibody was purchased from Affinity; and GAPDH antibody was purchased from CST. The Type 311 CO₂ constant temperature incubator was purchased from Thermo Company. The AMR-100 microplate reader was purchased from Hangzhou Aosheng. The DMIL LED inverted fluorescence microscope was purchased from Leica. The Mini-PROTEAN 3 cell electrophoresis instrument was purchased from BIO-RAD. The Tanon-5200 imaging system was purchased from Tanon. The CytoFLEX flow cytometer was purchased from Beckman Coulter. The Transwell Chamber model 3422 was purchased from COSTAR Corporation.

1.2 Molecular docking

IKK- β (PDB ID: 4KIK), MAPK3 (PDB ID: 2QNJ) and MAPK4 (PDB ID: 5ES1) data were downloaded from the RSCB PDB database (<http://www.rcsb.org/pdb/home/home.do>). The 3D structures of the target proteins were saved as a PDB format file. The mol2 format file of OA was downloaded from the ZINC website (<http://zinc.docking.org/>). The PyMOL software was used to remove water, phosphate, and other irrelevant ions from the protein. The target protein receptor and ligand were separated, and the ligand molecules were extracted. The pre-processed target protein and the original ligand were saved as PDB format files. AutoDockTools 1.5.6 software was used to set the size of Grid Box and box, search for the

parameters of protein active site (active port), convert the active small molecules, pretreated target protein, and original ligand into pdbqt format file, run AutoDock Tools 1.5.6 software to perform Autogrid and Docking operations, and finally use Open Babel GUI and PyMOL software to analyze the docking results visually.

1.3 Cell culture

Human glioma U251 cells were cultured in DMEM high glucose medium containing 10% fetal bovine serum and 1% penicillin-streptomycin in a 5% CO₂ incubator at 37°C. Cells in logarithmic growth phase were used in experiments.

1.4 Cell viability assay

Cell viability of U251 cells was measured using a CCK-8 assay. Cells were plated in a 96-well plate with 3×10³ cells/well in 100 µL per well and cultured overnight. Cells were treated with OA (0–2.5×10²–50–100 µg/mL), with three wells for each treatment group, for 8 h. Next, 10 µL of CCK-8 solution was added to each well, and cells were incubated for 4 h. The OD value of each well was measured at 450 nm with an enzyme-linked immunoassay. The relative proliferation inhibition rate was calculated as (%) = [1- (OD value of each drug group-OD value of the blank group) / (OD value of the control group-OD value of the blank group)] × 100%.

1.5 MTT assay

Cells were plated in 96-well culture plates at 1–5×10⁴ cells/mL and cultured overnight. Cells were then treated with OA (0–10²–50–100–200–500–1000 µg/mL), with three wells for each treatment group, and cultured for an additional 24 h. Next, 20 µL of MTT solution (5 mg/mL) was added to each well, and cells were incubated for 4 h. The supernatant was then discarded. DMSO (150 µL) was added to each well and the plates were shaken for 10 min. Absorbance at 490 nm was measured with an enzyme plate analyzer. In addition, the OA group (100 µg/mL) was operated according to the above method, and the absorbance (OD value) at 490nm wavelength was measured after 0, 1, 2, 3, 4, 5, and 6 days of culture. The proliferation inhibition rate of each group was calculated.

1.6 Colony formation assay

The cell Colony formation assay were set up in control group and oleanolic acid group (100 µg/mL). Cells were inoculated in a 6 cm dish at 1×10³ cells per dish and then cultured for 2 to 3 weeks. The culture was discontinued when the cells formed visible clones. The supernatant was discarded and the plates were washed with PBS; cells were fixed with 4% paraformaldehyde for 15 min. The fixed solution was discarded, and cells were stained with crystal violet staining solution for 10–30 min. The staining solution was removed and plates were air-dried. The number of clones was calculated using a microscope, and the rate of clone formation was determined as = (Number of clones/Number of inoculated cells) × 100%.

1.7 Transwell assays

The Transwell assays were set up in control group and oleanolic acid group (100 µg/mL). Transfected cells were cultured in serum-free medium for 24 h before the experiment. Matrigel mixed with serum-free medium (1:2; 80 µL) was added to a Transwell chamber, and the chamber was incubated at 37°C for 30 min to solidify. Cells (300 µL, 3×10^5 cells/mL) were added to the top chamber of the Transwell chamber and 700 µL of 10% FBS-containing complete culture medium was added to the bottom chambers. Cells were cultured for 24 h. Next, 4% formaldehyde solution was added to each well and cells were fixed at room temperature for 10 min. The fixing solution was removed and the cells were washed by PBS, followed by staining with 1 ml of 0.5% crystal violet solution for 30 min. Samples were washed three times with PBS and air-dried. The cells were observed and photographed under a microscope (at 400×), and the number of invaded cells in each group was counted.

1.8 Flow cytometry assay

The Flow cytometry assay were set up in control group and oleanolic acid group (100 µg/mL). Cells were harvested (5×10^4 to 1×10^5 cells) and resuspended in PBS. Next, 195 µL Annexin V-FITC binding solution was added to cells, followed by 10 µL Annexin V-FITC and 5 µL propidium iodide staining solution. The samples were mixed and incubated in the dark at room temperature for 15 min and then placed in an ice bath. A sample without Annexin V-FITC and PI was used as a negative control. The apoptosis rate of each group was detected by flow cytometry.

1.9 Western blotting assay

Cells were lysed with RIPA rapid lysis buffer and the BCA protein quantitative method was used to determine protein concentration. After denaturation, protein samples were separated by 10% SDS-PAGE gel electrophoresis and transferred to a PVDF membrane. The membrane was blocked with 5% skimmed milk powder at room temperature for 1 h and then incubated with the following primary antibodies overnight at 4°C: IKKβ antibody (1:600), MAPK3 antibody (1:1000), MAPK4 antibody (1:400), and GAPDH antibody (1:2000). After three washes in TBST for 5 min, the membrane was incubated with HRP-labeled secondary antibody (1:1000) at 37°C for 1 h and then washed with TBST three times. The membrane was developed using the ECL chemiluminescence method and the protein bands were visualized using a chemiluminescence imaging system. The protein bands were analyzed by the Tanon-5200 image analysis system with GAPDH as the internal reference.

1.10 RNA-Sequencing of Oleanolic acid

The control group and oleanolic acid treatment group were set up, and the total RNA of human glioma cell U251 was extracted respectively. RNA purity was detected by one / onec, and life Invitrogen qubit was used. 3.0 fluorescence quantizer was used for accurate quantification, and Agilent 4200 tapestation system was used to evaluate RNA integrity (RIN value). After the sample is qualified, the library construction of the sample is carried out. Then the library concentration was accurately quantified by Kapa qPCR and the library fragment size was accurately detected by Agilent 4200 TapeStation. After the library is qualified, the different libraries are pooled according to the effective concentration and the target off-machine data volume for Illumina PE150 sequencing.

1.11 GO and KEGG Enrichment Analyses

Gene Ontology (GO) is a comprehensive database describing gene functions, which can be divided into three parts: molecular function, biological process and cellular component. Kyoto Encyclopedia of Genes and Genomes (KEGG) is a public database resource for high-level functions and biological systems (such as cells, organisms, and ecosystems). It integrates genome, chemistry, and system function information. The categories of KEGG are represented by pathways. The clusterProfiler hypergeometric distribution algorithm was applied to the selected differential genes to analyze the GO function of the differential gene enrichment and the KEGG pathway enrichment analysis, and the enrichment results were visualized.

1.12 Statistical analysis

GraphPad Prism 5.0 software was used for image processing. The SPSS 21.0 software was used for statistical analysis. Measurement data are expressed as $\bar{x} \pm s$, and the comparison of multiple groups was performed by the one-way analysis of variance (ANOVA) method. $P < 0.05$ indicated statistical significance.

2. Results

2.1 OA shows binding activity with IKK- β , MAPK3, and MAPK4

The interaction model of OA with IKK- β , MAPK3, and MAPK4 was examined through molecular docking. The docking results are shown in Table 1 and Fig. 1. Studies have shown that a lower binding energy required for the binding of the ligand and the receptor is associated with a more stable binding conformation and higher binding possibility. A binding energy less than 0 indicates that the ligand and receptor can spontaneously bind. We found that the binding energy of OA with IKK- β , MAPK3, and MAPK4 was -5 KJ/mol through hydrogen bonds with ASP-484, TYR-134, ILE-62, LYS-193, GLU-81, and ASP-187 in IKK- β , MAPK3, and MAPK4 kinases, indicating that OA has an excellent binding activity with the three target proteins.

Table 1
Docking results of OA with IKK- β , MAPK3, and MAPK4 proteins

Chemical compound	Target	PDB_ID	Binding energy(KJ/mol)
Oleanolic acid	IKK- β	4KIK	-6.1
	MAPK3	2QNJ	-7.11
	MAPK4	5ES1	-6.64

2.2 OA inhibits the proliferation of U251 cells

The CCK-8 assay results showed that OA inhibited U251 glioma cells proliferation. Concentrations of 50 $\mu\text{g/mL}$ induced 100% inhibition. (Fig. 2). Comparing the different treatment groups with the control group, the difference in inhibition rate was statistically significant ($P < 0.05$). These results suggested that OA inhibits the proliferation of U251 glioma cells, and it has a concentration-dependent effect at 2.5–50 $\mu\text{g/mL}$.

The MTT assay results showed that OA inhibited U251 glioma cells proliferation. Concentrations of 500 $\mu\text{g/mL}$ induced 91% inhibition. Significant inhibition of proliferation was seen at concentrations as low as 10 $\mu\text{g/mL}$. (Fig. 3A). Comparing the different treatment groups with the control group, the difference in inhibition rate was statistically significant ($P < 0.05$). The results indicate that oleanolic acid can significantly inhibit the proliferation of U251 cells, and its inhibition rate is positively correlated with the concentration of 10 $\mu\text{g/mL}$ ~ 500 $\mu\text{g/mL}$, and there is a phenomenon of inhibition and saturation afterwards. After treatment with 100 $\mu\text{g/mL}$ oleanolic acid for 3 days, the proliferation ability of U251 cells decreased significantly, and after the 4th day, it had almost no effect on the proliferation ability of U251 cells (Fig. 3B).

2.3 Cell clone assay to detect cell proliferation

We next performed cell clone assays to evaluate the effect of OA on the colony formation ability of U251 glioma cells. The number of cell clones (41 ± 2) in the OA group was significantly reduced compared with the number in the control group (124 ± 7) ($P < 0.005$) (Fig. 4). These results confirmed that OA treatment significantly decreased the colony formation ability of U251 glioma cells.

2.4 Cell invasion was detected by Transwell chamber assay

Transwell experiments showed that the number of invaded cells in the OA treatment group (34 ± 4) was significantly reduced compared with the control group (74 ± 5) ($P < 0.005$) (Fig. 5). These results indicated that OA substantially inhibits the invasion ability of human glioma U251 cells.

2.5 OA promotes apoptosis of human glioma U251 cells

We next evaluated apoptosis using Annexin V-FITC/PI double staining and flow cytometry. The results showed that the apoptotic rate of U251 cells after OA treatment was $25.27 \pm 1.11\%$, while the apoptotic rate of the control group was $4.39 \pm 0.22\%$ (Fig. 6). The apoptotic rate increased significantly after OA treatment, and the difference was statistically significant ($P < 0.01$). These results showed that OA promotes the apoptosis of human glioma U251 cells.

2.6 OA downregulates the expression of CYP17A1, IKK- β , PTGS2, MAPK3, and MAPK4 protein

We next used western blot analysis to evaluate the expressions of CYP17A1, IKK- β , PTGS2, MAPK3, and MAPK4 proteins in U251 glioma cells treated with OA. After OA treatment of U251 glioma cells, the

expression levels of CYP17A1, IKK- β , PTGS2, MAPK3, and MAPK4 protein were reduced compared with control cells, and the differences were statistically significant ($P < 0.05$) (Fig. 7). These results indicate that OA treatment downregulates the expressions of CYP17A1, IKK- β , PTGS2, MAPK3, and MAPK4 proteins in U251 glioma cells.

2.7 mRNA-seq test results

2.7.1 Significant difference gene analysis

In order to understand the potential mechanism of oleanolic acid against glioma, we performed RNA-seq analysis on human glioma cells U251 treated with oleanolic acid and cells without oleanolic acid treatment as controls. In RNA-seq analysis, a total of 446 significantly differentially expressed genes were detected. The Volcano Plot chart can be very intuitive to view the overall distribution of genes that are differentially expressed between two samples. According to the set threshold: $|\log_2(\text{FoldChange})| > 1$ and $\text{padj} < 0.05$, we screen the differential genes between samples to obtain FoldChange and padj corresponding to the up-regulated and down-regulated genes. Among them, there are 96 up-regulated genes and 350 down-regulated genes that are differentially expressed. The results are shown in Fig. 8A. These genes are mainly involved in processes such as inflammation, metabolism, immunity, and regulation of cell growth. The hierarchical clustering analysis of genes was carried out, and the hierarchical clustering results were displayed by heat maps. The clustering results are shown in Fig. 8B. The clustering heat map between samples is drawn according to the fpkm value of different genes.

2.7.2 GO and KEGG Enrichment Analyses

In order to study the functions of oleanolic acid significantly different genes, clusterProfiler hypergeometric distribution algorithm was used to analyze the GO function and KEGG pathway enrichment analysis of the different genes. The results of GO and KEGG enrichment of differential genes are visualized, as shown in Fig. 9. In the GO functional enrichment category, it is about cellular biological processes, defense response defense response, cellular response to interferon, basement membrane, endoplasmic reticulum, extracellular matrix, cytokine activity, cytokine There are many genes with significant differences in the activity of receptor binding, cytokine receptor binding, transcription factor activity, and transcription activator activity. The KEGG pathway is enriched, with a total of 236 signal pathways enriched, including TNF signaling pathway, NOD-like receptor signaling pathway, and AGE-RAGE signaling pathway in diabetic complications, etc. TNF- α can up-regulate the protein expression of ERK1/2 in glioma cells. MAPK extracellular signal-regulated kinase MEK-ERK inhibitor can significantly attenuate the invasion of U87 glioma cells mediated by TNF- α stimulation. Inhibition of ERK1/2^[9]. The combination of TBK-1, a member of the IKB kinase family, and IKK ϵ promotes autophagy and inflammation by activating important factors related to cancer development, including Akt and TNF-R-related factor 2TRAF2, and inhibits tumor suppressor factors. Its inhibitors promote the survival and growth of glioblastoma cells through anti-inflammatory, autophagy and apoptosis pathways^[10]. It can be seen from KEGG results that oleanolic acid may activate MAPK pathway through TNF pathway, thus inhibiting the proliferation of human neural U251 cells.

3. Discussion

Glioma is the most common malignant tumor among all intracranial malignancies^[11] and is a complex tumor lesion that is characterized by invasiveness and a high recurrence rate. Despite the availability of treatments, such as early surgical resection, radiation, or chemotherapy, the prognosis of patients with glioma is poor. In addition to the poor therapeutic effect of chemotherapy, early radiotherapy, and late adjuvant therapy on malignant glioma, these treatments also have toxic side effects. Therefore, identifying new therapeutic agents to improve the prognosis of patients with malignant glioma is critical^[12, 13]. To overcome the limitations of the existing treatment methods for malignant glioma, many new treatment methods such as molecular targeted therapy, immunotherapy, gene therapy, stem cell therapy, and nanotechnology have begun to appear in preclinical and clinical research. In recent years, traditional Chinese medicine has shown significant value in the treatment of human glioma. Bi^[14] found that Paris polyphylla saponin reduced the viability of U251 cells by inhibiting the expression of ARA1 and ARA3, leading to inhibited phosphorylation of Akt and p44/42 MAPK and induced apoptosis and cell cycle arrest. Liu^[15] showed that berberine inhibited the proliferation of glioma cells by interfering with wild-type and mutant p53.

OA is an effective anti-tumor drug that inhibits glioma cells with a variety of malignant phenotypes. Studies have shown that OA and its derivatives have no cytotoxicity to normal human cells^[16] and it inhibits tumor development in various malignant tumors through multiple methods and pathways. In vivo experiments showed that OA significantly reduced the size and quality of tumors, and in vitro experiments revealed that OA has a significant inhibitory effect on cell migration, cell viability and proliferation ability^[17, 18]. OA has a sensitizing effect on radiotherapy of C6 cell-bearing rats^[19]. The pathway by which OA inhibits cell processes varies in different tumor cells, and its anti-tumor effect is related to the inhibition of intracellular signaling pathways such as STAT3, VEGF, MAPK, ERK, JNK, Akt, mTOR, and NF- κ B pathways. In this study, molecular docking technology showed that the binding energies of OA and IKK- β , MAPK3, and MAPK4 proteins were all less than - 5 kcal /mol, indicating that OA had a good binding activity with the three proteins. We speculate that OA may inhibit glioma proliferation by downregulating IKK- β , MAPK3, and MAPK4 proteins.

To further study the effect of OA on glioma, we performed in vitro experiments. CCK-8 and MTT assays revealed that the proliferation inhibition rate of cells increased in response to OA in a dose-dependent manner. Colony formation assays showed that the number of U251 cells decreased significantly after OA treatment. Together these results indicated that OA inhibits the proliferation of U251 cells. Invasion is a critical biological feature of malignant tumors. It is the process by which tumor cells penetrate the blood vessel wall and enter the circulatory system. Transwell assays showed that the number of invaded U251 cells treated with OA was significantly reduced compared with the control group, suggesting that OA inhibits the invasion of U251 cells. Apoptosis is a process in which cells undergo programmed death through gene regulation. Many tumor suppressor genes inhibit tumor development through inducing apoptosis, and apoptosis induction is one of the main mechanisms of current tumor treatments. Flow

cytometry showed that the apoptosis rate of cells treated with OA was significantly increased, indicating OA has a promoting effect on cell apoptosis. Together these results show that OA inhibits the proliferation and invasion of glioma U251 cells and promotes glioma cell apoptosis, thereby exhibiting anti-cancer effects in glioma.

The IKK and MAPK pathways play an essential role in the proliferation, migration, and invasion of glioma cells. The overexpression of IKK- β is closely related to the occurrence of a variety of malignant tumors, and inhibition of IKK- β in tumor cells results in specific anti-tumor effects [20–22]. Recent studies found a close relationship between malignant tumors and chronic inflammation. IKK- β activation helps promote the occurrence of cancer and inflammatory lesions, and inflammation can inhibit cell apoptosis through the IKK- β pathway, thus enabling the development malignant tumors. The specific inhibition of IKK- β provides a new strategy and approach for the prevention and targeted treatment of inflammation-associated tumors. Activation of the IKK/NF- κ B pathway induces tumor cells to proliferate and migrate and inhibits apoptosis [23]. Wang et al. showed that TGF β 1 regulates TNF- κ B activity induced by TNF- α in glioma cells through the PP1A/IKK- β pathway and promotes the progression and development of gliomas [24].

The MAPK pathway participates in the regulation of cancer cells through roles in proliferation, differentiation, and apoptosis, and other critical biological responses [25, 26]. Recent studies showed that the MAPK signaling pathway has a series of active metastases in glioma and other malignant tumor tissues, and MAPK directly mediates the anti-tumor effects of a variety of drugs. One study showed that miR-326 inhibits the malignant biological behavior of glioma cells through the MAPK signaling pathway [27]. MAPK3 and MAPK4 are members of the MAPK pathway. Yang et al. showed that miR-127-3p inhibits the proliferation of U251 glioma cells by down-regulating the target gene MAPK4 [28]. Some researchers also reported that a new MAPK3 and MAPK4 double-molecule inhibitor PCC0208017 reduced glioma cell migration, glioma cell invasion, and angiogenesis and may be a promising lead compound for the treatment of glioma [29]. Studies have confirmed that CYP17A1 gene silencing significantly inhibits the invasiveness of glioma cells, and CYP17A1 gene silencing promotes the apoptosis of glioma cells [30]. Recent studies have shown that PTGS2 can not only promote the proliferation rate of glioma cells, but also improve radiation tolerance [31]. The studies described above showed that the CYP17A1/IKK- β /PTGS2/MAPK3/MAPK4 signaling pathway plays an essential role in the development of glioma. In this study, our western blot analysis showed that the protein expression levels of CYP17A1, IKK- β , PTGS2, MAPK3, and MAPK4 were suppressed in glioma cells treated with OA. We speculate that OA may inhibit IKK- β , MAPK3, and MAPK4 protein expression by directly binding to the proteins. In this study, in vitro experiments observed that OA has inhibitory effects on the proliferation, cloning and invasion of glioma cells, and can promote their apoptosis. Furthermore, through transcriptome sequencing, it was found that the significantly differentially expressed genes are mainly involved in processes such as inflammation, metabolism, immunity, and regulation of cell growth. In the GO function enrichment analysis, information about cellular biological processes, defense response, cellular response to type interferon, basement membrane, endoplasmic reticulum, extracellular matrix, cytokine activity, cytokine receptor binding,

transcription factor activity and transcription activator activity, etc. There are many genes with significant differences in activity. It can be seen from KEGG results that oleanolic acid may activate MAPK pathway through TNF pathway, thus inhibiting the proliferation of human neural U251 cells.

Conclusions

In summary, we found that OA inhibits the proliferation and invasion of U251 glioma cells and promotes cell apoptosis. We further showed that OA inhibited the IKK- β /MAPK3/MAPK4 signaling pathway in glioma cells. These findings suggest that OA may have a good application prospect in the clinical treatment of glioma and provides ideas for the development and study of tumor drugs. The treatment of human glioma with oleanolic acid still needs to be further studied in vivo. Further in-depth research can lay a theoretical foundation for finding new molecular targeted therapeutic drugs at the gene level.

Abbreviations

OA: Oleanolic acid; U251: The human glioma cell line U251; CYP17A1: Cytochrome P450 Family 17 Subfamily A Member 1; IKK- β : inhibitor kappa B kinase β ; PTGS2: prostaglandin-endoperoxide synthase 2; MAPK3: mitogen-activated protein kinase 3; MAPK4: mitogen-activated protein kinase 4; GAPDH: glyceraldehyde-3phosphate dehydrogenase; RNA-seq: RNA Sequencing; GO: Gene Ontology; KEGG: Kyoto Encyclopedia of Genes and Genomes.

Declarations

Acknowledgments

None.

Authors' contributions

Luning Xu designed the research. Jinxiang Huang, Shengnan Lin, Feng Zhu, Dengsheng Chen, and Fang Huang performed experiments. Luning Xu, Jinxiang Huang, and Shengnan Lin performed data analyses. Luning Xu and Jinxiang Huang wrote the paper. All authors have read and approved the final manuscript.

Funding

This work was supported by Natural Science Foundation of Fujian Province (2018J01212, China).

Availability of data and materials

All data generated or analysed during this study are included in this published article.

Ethics approval and consent to participate

Not applicable.

Consent for publication

Not applicable.

Competing interests

The authors declare that there is no competing interests.

Author details

¹ Department of Neurosurgery, Shanghai Institute of Neurosurgery, Shanghai Changzheng Hospital, Second Military Medical University, Shanghai, China,

² Department of Pharmacy, Sanming First Hospital, Affiliated Hospital of Fujian Medical University, Sanming City, Fujian Province, China,

³ Department of Neurosurgery, Sanming First Hospital, Affiliated Hospital of Fujian Medical University, Sanming City, Fujian Province, China.

References

1. Goudarzi Kaveh M, Espinoza Jaime A, Guo Min B, Jiri, Nistér Monica, Lindström Mikael S, et al. Reduced Expression of PROX1 Transitions Glioblastoma Cells into a Mesenchymal Gene Expression Subtype. *Cancer research*,2018,78(20):5901–5916.
2. Diao T, Zhang Y, Lu W, Min Q. Inhibitory effect of total flavonoids from hawthorn leaves on glioma U87 cells. *Chinese Journal of Pharmacology*. 2019;35(10):1448–52.
3. Christopher M, Jackson J, Choi M, Lim. Mechanisms of immunotherapy resistance: lessons from glioblastoma. *Nature Immunology*,2019,20(14):1100–1109.
4. Andrew DJ, Pearson R, Herold R, Rousseau C, Copland B, Bradley-Garelik D, Binner, et al. Implementation of mechanism of action biology-driven early drug development for children with cancer. *Eur J Cancer*. 2016;62:124–31.
5. Xiaojing Shen H, Zhao L, Zhao. Weiwei Jiang. Research progress of oleanolic acid. *Guangzhou Chemical Industry*,2019,47(24).
6. Xiaofei Jiang M, Shi M, Sui Y, Yuan S, Zhang Q, Xia, et al. Oleanolic acid inhibits cervical cancer Hela cell proliferation through modulation of the ACSL4 ferroptosis signaling pathway. *Biochem Biophys*

Res Commun. 2021;545:124e131.

7. Du G, Zhang Y, Xiaohui Hu. Effects of Olenolic acid on cell proliferation and metastasis of human ovarian cancer SKOV3 cells. *China Pharmacy*,2020,31(10):1190–1197.
8. Tang S, Gao D, Zhao T, Zhou J, Zhao X. An evaluation of the anti-tumor efficacy of oleanolic acid-loaded PEGylated liposomes. *Nanotechnology*. 2013;24(23):235102.
9. Ramaswamy P, Goswami K, Nanjaiah ND, et al. TNF- α mediated MEK-ERK signaling in invasion with putative network involving NF- κ B and STAT-6: a new perspective in glioma. *Cell Biol Int*. 2019;43(11):1257–66.
10. Scuderi SA, Lanza M, Casili G, et al. TBK1 inhibitor exerts anti-proliferative effect on glioblastoma multiforme cells. *Oncol Res*. 2021 Mar 19.
11. Quinn T, Ostrom H, Gittleman J, Xu C, Kromer Y, Wolinsky C, Kruchko, et al. CBTRUS Statistical Report: Primary Brain and Other Central Nervous System Tumors Diagnosed in the United States in 2009–2013. *Neuro-oncology*,2016,18(suppl_5): v1-v75.
12. Ping Liu J, Yu X, Tian J, Chang Y, Zhang R, Zhang, et al. The effect of downregulation of Stathmin gene on biological behaviors of U373 and U87-MG glioblastoma cells. *Biol Res*. 2018;51(11):492–507.
13. Katharina Seystahl W, Wick M, Weller. Therapeutic options in recurrent glioblastoma—An update. *Critical Reviews in Oncology / Hematology*,2016,99: 389–408.
14. Linlin Bi Y, Liu Q, Yang X, Zhou H, Li Y, Liu, et al. Paris saponin H inhibits the proliferation of glioma cells through the A1 and A3 adenosine receptor–mediated pathway. *International Journal of Molecular Medicine*,2021,47(4): 1.
15. Ziqiang Liu Y, Chen H, Gao W, Xu C, Zhang J, Lai, et al. Berberine Inhibits Cell Proliferation by Interfering with Wild-Type and Mutant P53 in Human Glioma Cells. *OncoTargets and therapy*,2020,13: 12151–12162.
16. Guo G, Yao W, Zhang Q. Yongli Bo. Oleanolic Acid Suppresses Migration and Invasion of Malignant Glioma Cells by Inactivating MAPK/ERK Signaling Pathway. *PLOS ONE*,2013,8(8): e72079.
17. Edathara PM, Chintalapally S, Makani VKrishnaK, Pant C, Yerramsetty S, Manohar D, Rao, et al. Inhibitory Role of Oleanolic Acid and Esculetin in Hela Cells Involve Multiple Signaling Pathways. *Gene*,2020,771: 145370e145370.
18. Xiaofei Jiang M, Shi M, Sui Y, Yuan S, Zhang Q, Xia, et al. Oleanolic acid inhibits cervical cancer Hela cell proliferation through modulation of the ACSL4 ferroptosis signaling pathway. *Biochem Biophys Res Commun*. 2021;545:124e131.
19. Zhang Yu, Xu H, Wang H, Yu W, Zhao X, Xue Y. Fluorine-18-deoxyglucose positron emission tomography/computed tomography with Ki67 and GLUT-1 immunohistochemistry for evaluation of the radiosensitization effect of oleanolic acid on C6 rat gliomas. *Nuclear medicine communications*,2015,36(1):21e27.
20. Zhu Y, Zhou Y, Zhou X, Guo Y, Huang D, Zhang J, et al. S100A4 suppresses can-cer stem cell proliferation via interaction with the IKK/NF-kappaB signaling pathway. *BMC Cancer*. 2018;18(1):763.

21. Nikee Awasthee V, Rai S, Chava P, Nallasamy, Ajaikumar B, Kunnumakkara A, Bishayee, et al. Targeting IkappaB kinases for cancer therapy. *Seminars in Cancer Biology*,2019,56:12e24.
22. Jack A, Prescott, Simon J, Cook. Targeting IKK β in Cancer: Challenges and Opportunities for the Therapeutic Utilisation of IKK β Inhibitors. *Cells*,2018,7(9): e115.
23. Amanda L, Rinkenbaugh PC, Cogswell B, Calamini DE, Dunn AI, Persson, William A, Weiss, et al. IKK/NF- κ B signaling contributes to glioblastoma stem cell maintenance. *Oncotarget*, 2016, 7(43).
24. Huizhi Wang Z, Chen S, Wang X, Gao M, Qian W, Qiu, et al. TGF β 1-induced beta-site APP-cleaving enzyme 2 upregulation promotes tumorigenesis through the NF- κ B signalling pathway in human gliomas. *Molecular Oncology*, 2020.
25. Khaled A, Selim H, Abdelrasoul M, Aboelmagd, Ahmed M, Tawila. The role of the MAPK signaling, topoisomerase and dietary bioac-tives in controlling cancer incidence. *Diseases*. 2017;5(2):13.
26. Nan Tang J, Chang H-C, Lu Z, Zhuang H-L, Cheng J-X. Shi, et al. Rhein induces apoptosis and autophagy in human and rat glioma cells and mediates cell differentiation by ERK inhibition. *Microbial Pathogene-sis*. 2017;113:168–75.
27. Jingxu Zhou T, Xu Y, Qin YR, Wang H, Zhang X, et al. MicroRNA-326 Functions as a Tumor Suppressor in Glioma by Targeting the Nin One Binding Protein (NOB1). *PLoS ONE*. 2013;8(7):e68469.
28. Yan yang. Han meiwen, Wang yingrui. mechanism of miR-127-3p inhibiting glioma proliferation by downregulating MAPK4. *stroke nervous diseases*. 2018;25(04):435–9 + 470.
29. Li F, Liu Z, Sun H, Li C, Wang W, Ye L, et al. PCC0208017, a novel small-molecule inhibitor of MARK3/MARK4, suppresses glioma progression invitro and in vivo. *Acta Pharmaceutica Sinica B*. 2020;10(2):289–300.
30. Zhou J, Xu T, Yan Y, et al. MicroRNA-326 Functions as a Tumor Suppressor in Glioma by Targeting the Nin One Binding Protein (NOB1). *PLoS ONE*. 2013;8(7):e68469.
31. Niu WX, Zhou CX, Cheng CD, et al. Effects of lentivirus-mediated CYP17A1 gene silencing on the biological activity of glioma. *Neurosci Lett*. 2019;692:210–5.

Figures

Figure 1

Molecular docking diagram of OA and IKK- β , MAPK3, and MAPK4

Figure 2

CCK-8 assay to detect the inhibitory effect of OA on the proliferation of U251 glioma cells. Data represent mean $\pm$ SD of three different experiments. *** $P < 0.05$ compared with control group.

Figure 3

MTT assay to detect the inhibitory effect of OA on the proliferation of U251 glioma cells. Data represent mean $\pm$ SD of three different experiments. * $P < 0.05$ vs control group, *** $P < 0.05$ vs control group.

Figure 4

Cell cloning to detect the proliferation inhibitory effect of OA on human glioma U251 cell. Data represent mean $\pm$ SD of four different experiments. *** $P < 0.005$ compared with control group.

Figure 5

Transwell chamber method to detect the effect of OA on the invasion of human glioma U251 cells. Data represent mean $\pm$ SD of six different experiments. $*** P < 0.005$ compared with control group.

Figure 6

Annexin V-FITC/PI double staining method to detect the promoting effect of OA on the apoptosis of human glioma U251 cells. Data represent mean $\pm$ SD of three different experiments. $** P < 0.01$ compared with control group.

Figure 7

Western blot detection of OA on the expression of CYP17A1, IKK-β, PTGS2, MAPK3, and MAPK4 proteins in glioma U251 cells. Data represent mean ± SD of three different experiments. ** P < 0.01 compared with control group.

Figure 8

Volcano map and clustering heatmap of differentially expressed genes between oleanolic acid and the control group (Figure A represents the down-regulated genes in blue and red represents the up-regulated genes; in Figure B, U251_KBa, b, and c are controls Group genes, U251_4a, b, and c are the oleanolic acid experimental group genes, green indicates low gene expression, red indicates high gene expression, and the connecting line represents the clustering result)