

## Gastrodin has *in vitro* anticancer effects in human glioma U118 cells.

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### Abstract

Gastrodin is a biologically active substance extracted from *Gastrodia elata Bl.* which is a traditional Chinese medicine. The aim of this study was to determine the *in vitro* anticancer effects in human glioma U118 cells. Two concentrations of gastrodin, (0.5, 1.0 and 2.0 mg/mL) both had the strong growth inhibitory effects in U118 cells determined by MTT assay. Using flow cytometry assay, high concentration of gastrodin (2.0 mg/mL) treated U118 cells contained the most apoptosis cells (43.2%) and the low concentration of gastrodin (1.0 and 0.5 mg/mL) treated U118 cells also had the more apoptosis cells (23.3% and 9.8%) than control cells (2.1%). By qPCR assay, the experiments results showed that gastrodin could raise caspase-3, caspase-8, caspase-9, Bax, p53, p21, I $\kappa$ B- $\alpha$ , Fas, FasL, TIMP-1, TIMP-2 and reduce Bcl-2, Bcl-xL, NF- $\kappa$ B, EGF, EGFR, VEGF, Fit-1, MMP-2, MMP-9 mRNA expressions. From these results, gastrodin could be used as a medicine for cancer treatment.

**Keywords:** Gastrodin, Cancer, U118 cells, mRNA expression.

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### Introduction

Cancer is the second most serious disease that threatens human health. Therefore, it is a big dream of scientists to conquer cancer. Natural product is one of the most important way from which new drug and lead compound may possibly be discovered and from which actually many kinds of anticancer drugs come directly or indirectly. Paclitaxel is now the most outstanding natural anticancer drug among those discovered which is used widely in clinical practice in treatment of breast cancer, ovarian cancer and some of head and neck cancer and lung cancer [1,2].

Cancer inhibitors exist in various plants naturally, and have very good effects on human cancer prevention. These cancer inhibitors that occur naturally are safer as having low toxicity, as well as can reduce the pain of patients during the treatment [3]. But the activity of many cancer inhibitors existing in natural plants is lower than that of synthetic drugs, and these natural cancer inhibitors can substantially improve the treatment effect of cancer, so finding out a useful cancer inhibitor becomes the most important thing to improve and enhance the anti-cancer effects resources that occurs naturally.

Gastrodin is extracted from dried root of *Gastrodia elata Blume* in China using a medicine, gastrodin has a good sedative and hypnotic effects, and can relieve the symptoms of neurasthenia, insomnia and headache in traditional Chinese medicine. Gastrodin is used for blood pressure, antiplatelet effects, action on nerve cells and also showed the effects of

anti-free radical, protects cell membrane, increases immune function, and treats acute lung injury [4-8].

Studies reported written by experts and scholars as well as clinical application showed that gastrodin has good curative effects on hepatic ascitic cancer and colonic adenocarcinoma [9,10]. Gastrodin had anticancer effects in H22 hepatic ascitic tumor cells through NF- $\kappa$ B signaling activation in CD4+ T cells [4]. Gastrodin also had anticancer effects in colonic adenocarcinoma Caco-2 cells using passive paracellular transport pathway. In this study, we study the anticancer effects of gastrodin in human glioma U118 cells, to induce the apoptosis of cancer cells, meanwhile observe the mechanism of these anticancer effects.

### Materials and Methods

#### Cell line

Human glioma U118 cell line was purchased from ATCC (American Type Culture Collection, Manassas, VA, USA). The RPMI-1640 medium (Gibco Co., Birmingham, MI, USA) containing 10% foetal bovine serum (Gibco Co., Birmingham, MI, USA) and 1% penicillin-streptomycin (Gibco Co.) was used for Human glioma U118 cells culturing. The cancer cells were cultured in the RPMI-1640 medium at 37°C in a humidified atmosphere containing 5% CO<sub>2</sub> (Forma, model 311 S/N29035; Waltham, MA, USA). The medium was changed once per 2-3 d.

### MTT assay

Culture solution was added to adjust the concentration of cancer cells in logarithmic growth phase to  $2 \times 10^4$ /dish, which were added to the 96-well culture plate with 50  $\mu$ L per well, and placed in incubator with 5% CO<sub>2</sub> at 37°C for 24 h. Gastrodin were added into the 96-well plate with 50  $\mu$ L per well, to adjust the concentration of cancer cells to 1 or 2 mg/mL. The 50  $\mu$ L culture solution was added to the blank control group, which was cultured in CO<sub>2</sub> incubator for 48 h. Followed by, the blank control group which was added with MTT solution after the supernatant was removed and then incubated for 4 h. The 100  $\mu$ L DMSO was added to the blank control group after the supernatant was removed and shocked for 30 min, the enzyme standard instrument were used to detect at 492 nm [11].

### Flow cytometry assay

Single cell suspension was centrifuged to remove stationary liquid and washed by 3 mL PBS twice, and then centrifuged for 5 min; added with 1 ml PI staining solution and incubated in refrigerator at 4°C for 30 min without explosion to sunshine; and then filtered by 500-well copper mesh; flow cytometry detection and argon ion laser with 15 mA excitation light source and 488 nm wavelength were used for testing, and 630 nm band-pass filter to receive the light. The  $1 \times 10^4$  cells were collected by FSC/SSC scattered point diagram method, with gating technology used to exclude adhesive cells and cell debris, to analyze the percentage of apoptotic cells in PI fluorescence histogram [11].

### qPCR assay

RNAzol reagent was used to extract the total RNA from cancer cells, and DNase RNase-free was adopted to digest total RNA at 37°C for 15 min, and then RNeasy kit to purify RNA to adjust its concentration to 1  $\mu$ g/ $\mu$ L. RNA (2  $\mu$ g) was used as the template to synthesize cDNA by reacting with reverse transcriptase at 37°C for 120 min, at 99°C for 4 min, and at 4°C for 3 min respectively. After that, reverse transcription-polymerase chain reaction method was adopted to amplify the DNA expressions (Table 1), to measure the transcription level of mRNA, and GAPDH was used as the housekeeping genes of internal control group [12].

**Table 1.** Sequences of primers were used in this study.

Gene Name	Sequence
Caspase-3	Forward: 5'-CAA ACT TTT TCA GAG GGG ATC G-3'
	Reverse: 5'-GCA TAC TGT TTC AGC ATG GCA-3'
Caspase-8	Forward: 5'-CCC CAC CCT CAC TTT GCT-3'
	Reverse: 5'-GGA GGA CCA GGC TCA CTT A-3'
Caspase-9	Forward: 5'-GGC CCT TCC TCG CTT CAT CTC-3'
	Reverse: 5'-GGT CCT TGG GCC TTC CTG GTA T-3'

Bax	Forward: 5'-AAG CTG AGC GAG TGT CTC CGG CG-3'
	Reverse: 5'-CAG ATG CCG GTT CAG GTA CTC AGT C-3'
Bcl-2	Forward: 5'-CTC GTC GCT ACC GTC GTG ACT TGG-3'
	Reverse: 5'-CAG ATG CCG GTT CAG GTA CTC AGT C-3'
Bcl-xL	Forward: 5'-CCC AGA AAG GAT ACA GCT GG-3'
	Reverse: 5'-GCG ATC CGA CTC ACC AAT AC-3'
p53	Forward: 5'-GCT CTG ACT GTA CCA CCA TCC-3'
	Reverse: 5'-CTC TCG GAA CAT CTC GAA GCG-3'
p21	Forward: 5'-CTC AGA GGA GGC GCC ATG3'
	Reverse: 5'-GGG CGG ATT AGG GCT TCC3'
NF- $\kappa$ B	Forward: 5'-CAC TTA TGG ACA ACT ATG AGG TCT CTG G-3'
	Reverse: 5'-CTG TCT TGT GGA CAA CGC AGT GGA ATT TTA GG-3'
I $\kappa$ B- $\alpha$	Forward: 5'-GCT GAA GAA GGA GCG GCT ACT3'
	Reverse: 5'-TCG TAC TCC TCG TCT TTC ATG GA3'
Fas	Forward: 5'-GAA ATG AAA TCC AAA GCT-3'
	Reverse: 5'-TAA TTT AGA GGC AAA GTG GC-3'
FasL	Forward: 5'-GGA TTG GGC CTG GGG ATG TTT CA-3'
	Reverse: 5'-TTG TGG CTC AGG GGC AGG TTG TTG-3'
TIMP-1	Forward: 5'-GTC AGT GAG AAG CAA GTC GA-3'
	Reverse: 5'-ATG TTC TTC TCT GTG ACC CA-3'
TIMP-2	Forward: 5'-TGG GGA CAC CAG AAG TCA AC-3'
	Reverse: 5'-TTT TCA GAG CCT TGG AGG AG-3'
MMP-2	Reverse: 5'-CTT CTT CAA GGA CCG GTT CA-3'
	Forward: 5'-GCT GGC TGA GTA CCA GTA3'
MMP-9	Reverse: 5'-TGG GCT ACG TGA CCT ATG AC3'
	Forward: 5'-GCC CAG CCC ACC TCC ACT CC-3'
EGF	Reverse: 5'-GCC AAG CTC AGA AGG CTA C-3'
	Forward: 5'-CAG GCC AGC CTC GTC TCA T3'
EGFR	Reverse: 5'-TCG GTG CTG TGC GAT TTA3'
	Forward: 5'-TTT CTG GCA GTT GCT CCT C-3'
VEGF	Reverse: 5'-GCA CCC ATG GCA GAA GGA GGA G-3'
	Forward: 5'-GTG CTG ACG CTA ACT GAC C-3'
Fit-1	Reverse: 5'-CAA GTG GCCAGA GGC ATG GAG TT3'
	Forward: 5'-GAT GTA GTC TTTACC ATC CTG TTG-3'
GAPDH	Reverse: 5'-CGG AGT CAA CGG ATT TGG TC-3'
	Forward: 5'-AGC CTT CTC CAT GGT CGT GA-3'

### Statistical analysis

The *in vitro* experiments were presented as mean  $\pm$  standard deviation (SD). Differences between the mean values for

individual groups were assessed with one-way analysis of variance (ANOVA) with Duncan's multiple range test using SAS version 9.1 (SAS Institute Inc., Cary, NC, USA).

## Results

### Growth inhibitory effects of gastrodin in U118 cells

By the MTT assay, the untreated U118 cells showed the OD540 value at 0.479 (Table 2), after 0.5, 1.0 and 2.0 mg/mL gastrodin treatment, the OD540 values were reduced at 0.317, 0.197 and 0.063 respectively. The 0.5, 1.0 and 2.0 mg/mL gastrodin showed the inhibitory effects at 33.8%, 58.9% and 86.8%, respectively.

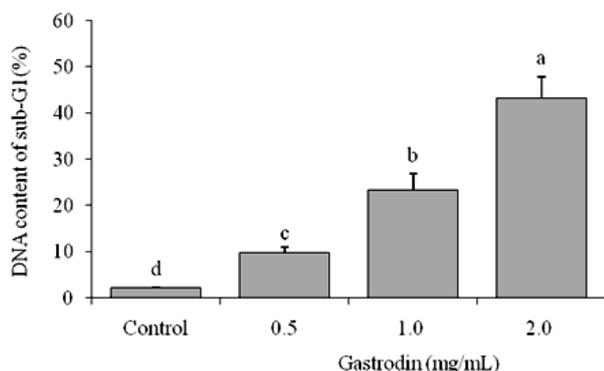
**Table 2.** Growth inhibitory effects of human glioma U118 cells treated by gastrodin by MTT assay.

Treatment	OD492 value	Inhibitory rate (%)
Control	0.479 ± 0.006 <sup>a</sup>	/
Gastrodin (mg/kg)	0.5	0.317 ± 0.011 <sup>b</sup>
	1	0.197 ± 0.009 <sup>c</sup>
	2	0.063 ± 0.005 <sup>d</sup>

<sup>a,b,c,d</sup>Mean values with different letters in the same column are significantly different (P<0.05) according to Duncan's multiple-range test.

### DNA content of sub-G1 U118 cells

The flow cytometry showed that control cells has only 2.1 ± 0.2% DNA content of sub-G1 of U118 cells (apoptotic cells), and as per the results in the other groups the cells have more apoptosis because of gastrodin treatment (Figure 1). The 2.0, 1.0 and 0.5 mg/mL gastrodin treated U118 cells had 43.2 ± 4.7%, 23.3 ± 3.6% and 9.8 ± 1.2% apoptosis cells, respectively.



**Figure 1.** Gastrodin induced apoptotic cells (sub-G1 DNA content) in OS-732 human osteosarcoma cells. <sup>a,b,c,d</sup>Mean values with different letters over the bars are significantly different (P<0.05) according to Duncan's multiple-range test.

### mRNA expressions of caspases

By qPCR experiment, control cells showed the weakest caspase-3, caspase-8 and caspase-9 also the mRNA expressions were weakest (Table 3). Gastrodin treated cells

showed remarkably stronger caspase-3, caspase-8 and caspase-9 expressions than control cells, and 2.0 mg/mL gastrodin treatment had the strongest caspase-3 (4.87 fold of control), caspase-8 (5.32 fold of control) and caspase-9 (4.33 fold of control) expressions.

**Table 3.** Quantitative analysis of mRNA expressions of caspase-3, caspase-8 and caspase-9 in U118 cells treated by gastrodin.

Group	Caspase-3	Caspase-8	Caspase-9
Control	1.00 ± 0.21 <sup>d</sup>	1.00 ± 0.17 <sup>d</sup>	1.00 ± 0.11 <sup>d</sup>
Gastrodin (mg/kg)	0.5	2.31 ± 0.29 <sup>c</sup>	1.68 ± 0.23 <sup>c</sup>
	1	2.92 ± 0.21 <sup>b</sup>	3.06 ± 0.15 <sup>b</sup>
	2	4.87 ± 0.48 <sup>a</sup>	5.32 ± 0.52 <sup>a</sup>

<sup>a,b,c,d</sup>Mean values with different letters in the same column are significantly different (P<0.05) according to Duncan's multiple-range test.

### mRNA expressions of Bax, Bcl-2 and Bcl-xL

Gastrodin treated U118 cells had the high Bax mRNA expressions and low Bcl-2, Bcl-xL expressions (Table 4). Whereas the high concentration of 2.0 mg/mL treatment were showing the highest Bax (3.36 fold of control) expressions than other groups cells, but this concentration treatment showed the lowest Bcl-2 (0.22 fold of control), Bcl-xL (0.12 fold of control) expressions.

**Table 4.** Quantitative analysis of mRNA expressions of Bax, Bcl-2 and Bcl-xL in U118 cells treated by gastrodin.

Group	Bax	Bcl-2	Bcl-xL
Control	1.00 ± 0.08 <sup>d</sup>	1.00 ± 0.06 <sup>a</sup>	1.00 ± 0.05 <sup>a</sup>
Gastrodin (mg/kg)	0.5	1.86 ± 0.22 <sup>c</sup>	0.77 ± 0.04 <sup>b</sup>
	1	2.31 ± 0.19 <sup>b</sup>	0.53 ± 0.05 <sup>c</sup>
	2	3.36 ± 0.33 <sup>a</sup>	0.22 ± 0.04 <sup>d</sup>

<sup>a,b,c,d</sup>Mean values with different letters in the same column are significantly different (P<0.05) according to Duncan's multiple-range test.

### mRNA expressions of Fas and FasL

The control cells had the lowest Fas and FasL mRNA expressions (Table 5), after gastrodin treatment, these expressions were elevated, raising concentration treated cells had the remarkably higher Fas (6.39 fold of control at 2.0 mg/kg gastrodin) and FasL (2.82 fold of control at 2.0 mg/kg gastrodin) expressions with respect to low concentration treated cells.

**Table 5.** Quantitative analysis of mRNA expressions of Fas and FasL in U118 cells treated by gastrodin.

Group	Fas	FasL
Control	1.00 ± 0.14 <sup>d</sup>	1.00 ± 0.09 <sup>d</sup>
Gastrodin	0.5	3.05 ± 0.38 <sup>c</sup>
	2.0	1.45 ± 0.12 <sup>c</sup>

(mg/kg)	1	4.38 ± 0.42 <sup>b</sup>	2.32 ± 0.21 <sup>b</sup>
	2	6.39 ± 0.50 <sup>a</sup>	2.82 ± 0.36 <sup>a</sup>

<sup>a,b,c,d</sup>Mean values with different letters in the same column are significantly different (P<0.05) according to Duncan's multiple-range test.

### mRNA expressions of p53 and p21

The p53 and p21 mRNA expressions of control cells were weakest (Table 6), gastrodin could increase these expressions, and high concentration of gastrodin had the stronger capability to increase p53 (5.31 fold of control at 2.0 mg/kg gastrodin) and p21 (4.88 fold of control at 2.0 mg/kg gastrodin) activities.

**Table 6.** Quantitative analysis of mRNA expressions of p53 and p21 in U118 cells treated by gastrodin.

Group		p53	p21
Control		1.00 ± 0.15 <sup>d</sup>	1.00 ± 0.14 <sup>d</sup>
Gastrodin (mg/kg)	0.5	2.31 ± 0.30 <sup>c</sup>	2.11 ± 0.18 <sup>c</sup>
	1	3.50 ± 0.45 <sup>b</sup>	3.09 ± 0.37 <sup>b</sup>
	2	5.13 ± 0.44 <sup>a</sup>	4.88 ± 0.46 <sup>a</sup>

<sup>a,b,c,d</sup>Mean values with different letters in the same column are significantly different (P<0.05) according to Duncan's multiple-range test.

### mRNA expressions of NF-κB and IκB-α

Gastrodin (2.0 mg/mL) group cells had the lowest NF-κB (0.19 fold of control) mRNA expression and the highest IκB-α (3.81 fold of control) expression (Table 7). Control cells had the highest NF-κB expressions and the lowest IκB-α expression.

**Table 7.** Quantitative analysis of mRNA expressions of NF-κB and IκB-α in U118 cells treated by gastrodin.

Group		NF-κB	IκB-α
Control		1.00 ± 0.05 <sup>a</sup>	1.00 ± 0.06 <sup>d</sup>
Gastrodin (mg/kg)	0.5	0.69 ± 0.06 <sup>b</sup>	1.49 ± 0.11 <sup>c</sup>
	1	0.41 ± 0.04 <sup>c</sup>	2.41 ± 0.25 <sup>b</sup>
	2	0.19 ± 0.03 <sup>d</sup>	3.81 ± 0.33 <sup>a</sup>

<sup>a,b,c,d</sup>Mean values with different letters in the same column are significantly different (P<0.05) according to Duncan's multiple-range test.

### mRNA expressions of TIMP-1, TIMP-2, MMP-2 and MMP-9

The TIMP-1, TIMP-2 mRNA expressions in control cells were lowest than other groups cells, but MMP-2, MMP-9 expressions were highest than other groups cells (Table 8). Gastrodin could raise TIMP-1, TIMP-2 expressions and reduce MMP-2, MMP-9 expressions as compared to the control cells, and after raising its concentration showed higher TIMP-1 (4.20 fold of control at 2.0 mg/kg gastrodin), TIMP-2 (3.80 fold of control at 2.0 mg/kg gastrodin) expressions and lower MMP-2 (0.14 fold of control at 2.0 mg/kg gastrodin), MMP-9 (0.22

fold of control at 2.0 mg/kg gastrodin) expressions than low concentration treated cells.

**Table 8.** Quantitative analysis of mRNA expressions of TIMP-1, TIMP-2, MMP-2 and MMP-9 in U118 cells treated by gastrodin.

Group		TIMP-1	TIMP-2	MMP-2	MMP-9
Control		1.00 ± 0.08 <sup>d</sup>	1.00 ± 0.09 <sup>d</sup>	1.00 ± 0.06 <sup>a</sup>	1.00 ± 0.09 <sup>a</sup>
Gastrodin (mg/kg)	0.5	2.03 ± 0.28 <sup>c</sup>	1.75 ± 0.19 <sup>c</sup>	0.70 ± 0.08 <sup>b</sup>	0.68 ± 0.06 <sup>b</sup>
	1	2.68 ± 0.24 <sup>b</sup>	2.59 ± 0.31 <sup>b</sup>	0.39 ± 0.05 <sup>c</sup>	0.41 ± 0.05 <sup>c</sup>
	2	4.20 ± 0.41 <sup>a</sup>	3.80 ± 0.39 <sup>a</sup>	0.14 ± 0.04 <sup>d</sup>	0.22 ± 0.04 <sup>d</sup>

<sup>a,b,c,d</sup>Mean values with different letters in the same column are significantly different (P<0.05) according to Duncan's multiple-range test.

### mRNA expressions of EGF, EGFR, VEGF and Fit-1

Gastrodin treatment reduce EGF, EGFR, VEGF, Fit-1 mRNA expressions as compared to the control cells (Table 9), and high concentration gastrodin showed further reduction in expression of EGF (0.34 fold of control at 2.0 mg/kg gastrodin), EGFR (0.28 fold of control at 2.0 mg/kg gastrodin), VEGF (0.11 fold of control at 2.0 mg/kg gastrodin), Fit-1 (0.29 fold of control at 2.0 mg/kg gastrodin).

**Table 9.** Quantitative analysis of mRNA expressions of EGF, EGFR, VEGF and Fit-1 in U118 cells treated by gastrodin.

Group		EGF	EGFR	VEGF	Fit-1
Control		1.00 ± 0.11 <sup>a</sup>	1.00 ± 0.12 <sup>a</sup>	1.00 ± 0.08 <sup>a</sup>	1.00 ± 0.10 <sup>a</sup>
Gastrodin (mg/kg)	0.5	0.83 ± 0.07 <sup>b</sup>	0.76 ± 0.08 <sup>b</sup>	0.64 ± 0.07 <sup>b</sup>	0.73 ± 0.09 <sup>b</sup>
	1	0.62 ± 0.06 <sup>c</sup>	0.52 ± 0.05 <sup>c</sup>	0.35 ± 0.07 <sup>c</sup>	0.45 ± 0.06 <sup>c</sup>
	2	0.34 ± 0.06 <sup>d</sup>	0.28 ± 0.06 <sup>d</sup>	0.11 ± 0.05 <sup>d</sup>	0.29 ± 0.07 <sup>d</sup>

<sup>a,b,c,d</sup>Mean values with different letters in the same column are significantly different (P<0.05) according to Duncan's multiple-range test.

## Discussion

Apoptosis of cancer cell plays an important role in the occurrence and development of cancer, Wong found that a lot of receptor-mediated cell signal transduction and many different genes are involved in the activation of cancer cells apoptosis, and regulation of cancer cell apoptosis respectively [13]. As an upstream protein involved in exogenous apoptosis, caspase-8 shears and activates downstream apoptosis-inducing proteins such as caspase-3, caspase-6 and caspase-7, causing cell apoptosis [14]. Apaf-1 can bond to the original structural domain of the precursor of caspase-9 through the complementary domain of caspase, leading to the self-activation of caspase-9, which further activates downstream caspase-3, caspase-6 and caspase-7, and ultimately inducing endogenous apoptosis of cells [15]. Caspase-3 involves both exogenous and endogenous apoptosis, and many apoptotic factors work on downstream effector caspase-3 ultimately to induce cell apoptosis [16].

The inhibition of apoptosis has a vital significance to the incidence and development of cancer. Proteins in Bcl-2 family play the important roles in regulating the apoptosis of cancer cells. Bcl-2 family is made up of apoptosis inhibitory factor (*Bcl-2* and *Bcl-xL*) and apoptosis-promoting factor (Bax); their ratio determines whether the cell is able to accept the apoptotic signal [17]. To a certain extent, apoptosis or apoptosis inhibition are regulated by the above two genes. The disturbance of apoptosis regulation is crucial in the development of tumor, and Bcl-2 family plays a major role in this process [18]. As the main members of Bcl-2 family, Bcl-2, Bax and Bcl-xL mainly regulate the apoptosis of cells by affecting mitochondrial pathway. When cells get death signals, the Bax which is bonded to Bcl-2 or Bcl-xL will be displaced, results increase in the permeability of the mitochondrial membrane and leading to the release of a series of substances, thus eventually causing the death of cells [19].

Fas, FasL and caspase-3 are the important proteins mediating the apoptosis of cells. At present, it has been found that FasL can be induced by certain stress responses, such as ultraviolet and DNA damage, and the interactions between FasL and Fas can induce programmed death of cells, which may be an important mechanism of the body to clear cells having mutation [20]. FasL can express on the surface of tumor cells, and tumor-specific antigen can induce tumor infiltrating T lymphocytes (TIL) to express Fas in large quantity, it enhances the sensibility of T cells to apoptosis. Tumor cells induce the apoptosis of T lymphocyte which cause the high expression of Fas by FasL, resulting in immunosuppression. Fas-mediated apoptosis is also related to many other factors, such as *p53* gene mutation or the lack of co-stimulatory factor [21].

*p53*, the major protein regulating Bcl-2 family, regulates different proteins of Bcl-2 family in various ways, affecting the biological behaviours of pancreatic cancer. *p53* can up-regulate Bax and down-regulate Bcl-2 or Bcl-xL, affecting the apoptosis of cancer cells, and changing the permeability of mitochondria, thus affecting the function of downstream pro-apoptotic genes [22]. As the clumping factor of CDK, low concentrations of tumor suppressor gene *p21* positively regulates the function of CDK, facilitating the development of cells and promoting the transition from G1 stage to S stage, but highly expressed *p21* protein and cyclin bind to CDK competitively to inhibit the activity of CDK, causing the cell development stagnating in G1 stage, thus inhibiting cell proliferation or inducing cell apoptosis [23]. *p73* and *p53* protein have homology in target gene binding, but their functions have great differences. As *p73* can arrest cell cycle and induce cell apoptosis, it can inhibit tumor to certain extent [24].

NF- $\kappa$ B system is composed of NF- $\kappa$ B family and its inhibitor I $\kappa$ B- $\alpha$ . NF- $\kappa$ B is an extremely important transcriptional activator, and I $\kappa$ B- $\alpha$  is the inhibitory protein of NF- $\kappa$ B [25]. NF- $\kappa$ B is important to inflammation process, and also serves as regulatory protein in the development of cancer. It plays an important role in information transmission in relation to tumor growth, closely related to the incidence and development of

tumor [26]. Studies have found that NF- $\kappa$ B highly expresses in many types of tumors, and activated NF- $\kappa$ B promotes the expression of a variety of genes which involve the development of cancer [27,28]. Wu et al. found that Hp infection activated NF- $\kappa$ B and the expression of COX-2 play important roles in the incidence and development of cancer [29].

Malignant tumors are characterized by local invasion and distant metastasis, which are the main reasons that malignant tumor threaten patients' health and life. MMPs play an important role in the invasion and metastasis of tumor, it not only mediates tumor cells' degradation of extracellular matrix including the basement membrane, but also controls the process of angiogenesis, that affects the function of cell adhesion molecules and regulates the growth of tumor cells [30]. Study has shown that the expression of MMP-2 and MMP-9 is closely related to cancer angiogenesis; tumor cells which can secrete MMP-2 and MMP-9 have high invasion and metastases ability, drugs can also be used to inhibit the growth of tumor cells through lowering the activity of MMP-2 and MMP-9 [31]. In addition, ECM play a key role in local invasion and distant metastasis of cancer cells, the degradation of ECM is complex, as it involved a lot of factors, and MMPs and inhibitors play important functions, MMPs can degrade ECM, while TIMPs can inhibit the degradation of ECM through lowering the activity of MMPs, to protect normal cells [32]. The formation of intravascular cavity depends on the balance of MMPs and TIMPs, introducing exogenous inhibitors may break the balance of MMPs and TIMPs, inhibiting the process of angiogenesis, as well as the invasion and metastasis of tumor cells. Therefore TIMPs can inhibit tumor invasion and metastasis, and has a remarkable use in the research of tumor treatment [33].

EGF is a kind of growth factor which can affect many reactions by combining with EGFR. Study has shown that EGF and other growth factors can promote the proliferation of human cells. EGFR is a member of ErbB receptor family located on the surface of cells, involved in cells proliferation, growth, migration and infiltration [34]. Study has found that EGFR can be adjusted by EGF-mediated cancer cell proliferation through sialylation [35].

VEGF can promote the growth of tumor and angiogenesis, and provides a foundation for tumor metastasis, affecting the prognosis of patients with tumor. VEGF is the strongest vascular endothelial cell growth factor which can directly work on blood vessels, and specifically promote the division, proliferation and migration of endothelial tumor cells, playing an important role in the formation of tumor blood vessel, and it is also one of the key factors of promoting angiogenesis [36]. Fit-1 is the receptor of VEGF; it can bind to VEGF in high affinity. Fit-1 receptor deficient mice are mainly characterized by vascular endothelial cell damage; the expression of Fit-1 is mainly related to the early-stage angiogenesis and wound healing of mouse embryos [37].

## Conclusion

In this study, MTT assay, flow cytometry and qPCR assays were used to determine the anticancer effects of gastrodin in human glioma U118 cells. Gastrodin had a good anticancer effect on U118 cells, and from these results, we can conclude that gastrodin could be used as a cancer cell inhibitor for glioma treatment, and this inhibitor might be used in clinical application to save human life in future.

## Conflict of Interest

There is no conflict of interest.

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