## Experimental study on Qizhu formula for modulating survivin, an inhibitor of apoptosis, in MGC-803 gastric cancer cells

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Abstract. The aim of this study was to investigate the mechanism of action of Qizhu formula, a Chinese medicinal empirical formula, in modulating the action of survivin, an inhibitor of apoptosis, in MGC-803 gastric cancer cells. Western blot analysis and reverse transcription-polymerase chain reaction (RT-PCR) were applied to detect the effect of varying concentrations of Qizhu formula in the modulation of the expression of survivin in MGC-803 human gastric adenocarcinoma cells. The western blot analysis results demonstrated that Qizhu formula exerted no significant effects on the protein expression of the  $\beta$ -actin housekeeping gene, whereas it exerted a significant inhibitory effect on the protein expression of the apoptosis-related survivin gene at concentrations of 250 µg/ml and, particularly, 500 µg/ml. RT-PCR was used to detect the effect of Qizhu formula on survivin mRNA in MGC-803 human gastric adenocarcinoma cells. The ratio of survivin/ $\beta$ -actin in the 0.1% dimethylsulfoxide and the 125, 250 and 500  $\mu$ g/ml groups of Qizhu formula was 0.4543, 0.4025, 0.2415 and 0.2235, respectively. Therefore, Qizhu formula exerted a distinct inhibitory effect on the mRNA expression of survivin in MGC-803 cells in a dose-dependent manner. In conclusion, Qizhu formula may modulate the apoptosis of MGC-803 human gastric adenocarcinoma cells, which is associated with the downregulation of survivin mRNA and protein expression.

## Introduction

Gastric cancer is the most common gastrointestinal cancer, with the highest morbidity and mortality rates in China,

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ranking second in the incidence of malignant tumors worldwide. Therefore, novel, more effective therapeutic options are required for the prevention and treatment of gastric cancer.

The occurrence and development of gastric cancer is a complicated, multistage and multifactorial process. An imbalance between cell apoptosis and proliferation is a key reason resulting in excessive proliferation of tumor cells. Survivin, which was screened and separated from the human genomic library by Ambrosini et al (1) by means of effector cell protease receptor-1 cDNA, is a novel member of the inhibitor of apoptosis (IAP) family and a bifunctional protein that may inhibit cell apoptosis and regulate cell division. Survivin is highly expressed in embryonic and developing fetal tissues, but it is almost absent from the terminally differentiated tissues of normal adults, whereas it is selectively expressed in tumors. Survivin expression was previously detected in colon, gastric, esophageal and non-small-cell lung cancers (2-5) and a previous study indicated that survivin is highly expressed in gastric cancer tissues, but is almost absent from the normal tissues surrounding gastric cancer (6). However, Yao et al (7) detected survivin expression in gastric adenocarcinoma using the immunohistochemical streptavidin-peroxidase method and the results demonstrated that the positive expression rate of survivin in the primary focus of gastric cancer, metastatic cancer cells of lymph nodes and basal germinal layer cells of normal glands were 49.2, 64.0, and 17.5%, respectively. Therefore, further investigation into the expression of survivin in tumor cells is required to elucidate the mechanisms underlying the occurrence and development of gastric cancer.

The majority of gastric cancers are middle- to advanced-stage at diagnosis and exhibit a low remission rate with chemotherapy. The anticancer role of traditional Chinese medicine (TCM) is under ongoing investigation in the medical field. In TCM, gastric cancer belongs to the classification of 'sick', 'dysphagia', 'stomachache' and 'accumulation' disease. Qizhu, an empirical formula for gastrointestinal tumors, has been subjected to numerous years of clinical research. Our earlier study (8) demonstrated that the Qizhu formula was able to inhibit the expression of telomerase and its related genes in MGC-803 gastric cancer cells.

In the present study, the effect of Qizhu on the downregulation of survivin protein and mRNA expression in MGC-803 human gastric adenocarcinoma cells was investigated by

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western blot analysis and reverse transcription-polymerase chain reaction (RT-PCR), with the aim to further demonstrate the induction of gastric cancer cell apoptosis by Qizhu and provide reliable experimental evidence regarding multitargeted treatment and application of TCM compounds in the treatment of gastric cancer.

## Materials and methods

*Tumor strain*. The MGC-803 human gastric cancer cells were purchased from Nanjing Kaiteng Technological Corporation (Nanjing, China) and preserved by subculturing.

Drugs. Qizhu formula crude extracts (referred to as Qizhu formula hereafter) were purchased from Nanjing Zhongshan Pharmaceutical Factory (Nanjing, China). Qizhu formula consists of Zhihuangqi (Radix Astragali Praeparata cum Melle), Yuzhu (Rhizoma Polygonati Odorati), Fabanxia (Rhizoma Pinelliae Praeparatum), Shengyiyiren (Semen Coicis, raw), Xianhecao (Herba Agrimoniae), Ezhu (Rhizoma Curcumae) and Baihuasheshecao (Herba Hedyotis Diffusae). All the compounds in the formula were identified by TCM Identification Teaching and Research Section of Nanjing University of Chinese Medicine and conformed to the national pharmacopoeia criterion. Steam distillation was applied to extract zedoary oil, with 11.25 g extracted from 500 g of crude Ezhu, yielding 11.25 ml, for a yield of 2.25%, namely 1 g=1 ml. The extracted residual Ezhu liquid was mixed with other crude drugs and the crude drugs were then extracted by poaching and alcohol precipitation. A total of 480.5 g crude extracts (dry extracts, 496.25 g + zedoary oil, 11.25 g) was collected from 4,500 g of crude drugs, for a yield of 10.68% (480.5/4,500).

## Reagents

Western blotting. Methanol, 200 ml 10X blotting buffer, 1,400 ml deionized water and Coomassie brilliant blue G<sub>250</sub> were collected for the western blotting procedure. First, 20 mg of Coomassie brilliant blue G250 were dissolved into 10 ml of 95% alcohol, followed by the addition of 20 ml H<sub>3</sub>PO<sub>4</sub> and deionized water to a final volume of 200 ml. The mixture was filtered using filter paper and stored at 4°C. Subsequently, 0.05 g bovine serum albumin (BSA) were dissolved in 100 ml of double-distilled water (0.5 mg/ml) and stored at 4°C. Finally, 0.01742 g phenylmethanesulfonyl fluoride (PMSF) were dissolved in 1 ml of isopropanol (100 mM), subpackaged and stored at -20°C. Acrylamide and ammonium persulfate were purchased from Longxi Chemical Company (Nanjing, China), tetramethylethylenediamine was obtained from American Amresco Co. (Solon, OH, USA) and horseradish peroxide solutions A and B were purchased from Beijing ZSGB-Biotechnology Co., Ltd. (Beijing, China).

*RT-PCR*. TRIzol (DP-405) was purchased from Beijing Tianwei Time Technology Co., Ltd. (Beijing, China), general RT-PCR kits (20 reactions) were obtained from Beijing Dingguo Biotechnology Co., Ltd. (Beijing, China) and synthetic primers of survivin and  $\beta$ -actin genes were obtained from Shanghai Sangon Biotech Service Co., Ltd. (Shanghai, China). The TRIzol kits were provided by Tianyao Time Company (Tianjin, China) and diethyl pyrocarbonate was obtained from American Amresco Co. Pure chloroform, isopropanol and ethanol were all analyzed by the Longxi Chemical Company and agarose was obtained from Oxoid Limited (Hampshire, UK).

Main experimental apparatus. A laminar flow hood was obtained from Suzhou Purification Equipment Factory (Suzhou, China); a Forman 3111 CO2 incubator was purchased from (Thermoscientific, West Palm Beach, FL, USA); a 550 enzyme-linked immunoassay detector was obtained from Bio-Rad Laboratories (Hercules, CA, USA); an LDZ5-2 medical centrifuge was purchased from Beijing Medical Centrifuge Factory (Beijing, China); a UV-2450 ultraviolet spectrophotometer was obtained from Shimadzu Co. (Milton Keynes, UK); a PHS-3C Precision pH meter was purchased from Shanghai Torpedo-Magnetic Instrument Plant (Shanghai, China); an FAZ104 electronic balance was obtained from Meigele Hardware Co., Ltd. (Guangzhou, China); a 101AS-3 stainless steel digital electrothermal air dry oven was purchased from Shanghai Shengxin Scientific Instrument Co., Ltd. (Shanghai, China); a YXQ-CS-30L double-layer stainless vertical sterilizer was obtained from the Medical Equipment Factory of Shanghai Boxun Industrial Co., Ltd. (Shanghai, China); a WD-940153 horizontal shaker was purchased from Beijing Liuvi Instrument Plant (Beijing, China); an XSZ-O2 inverted metallographic microscope was obtained from Chongqing Optical Instrument Plant (Chongqing, China); a PTC-100 PCR amplifier was purchased from MJ Reserach Inc. (Waltham, MA, USA); a SHO 3014 tabletop high-speed refrigerated centrifuge was obtained from Eppendorf (Hamburg, Germany); a Mini-PROTEAN vertical electrotransfer chamber was purchased from Bio-Rad Laboratories; and an FR-980 biological electrophoresis image analysis system was obtained from Shanghai Furi Technology Co., Ltd (Shanghai, China).

## Assessment of Qizhu formula effect on survivin protein expression in MGC-803 cells by western blot analysis

*Grouping*. A negative control group, a dimethylsulfoxide (DMSO) solvent control group and two medication groups with different concentrations of Qizhu formula (250 and 500  $\mu$ g/ml) were set up.

*Methods*. The MGC-803 cells were first cultured to a certain concentration. Four 10-ml culture flasks were then inoculated with  $2\times10^6$  cells and placed in a 5% CO<sub>2</sub> incubator for 24 h at 37°C. Subsequently, a negative control group, a 0.1% DMSO solvent control group and two medication groups with different concentrations of Qizhu formula (250 and 500  $\mu$ g/ml) were set up. Finally, 10 ml of the cultured cells were added to each of the four groups and placed in a 5% CO<sub>2</sub> incubator at 37°C for 24 h.

*Cell collection.* Cell scraping paper was used to collect 10 ml of cells in each group. The cells were centrifuged for 5 min at 352xg, followed by the addition of 1 ml PBS to transfer the centrifuged cells to Eppendorf tubes and the mixture was centrifuged at 1,570xg for 1 min. The supernatant was then removed, the cells were washed again with PBS, beakers were prepared and tap water was added and heated to 95°C. Furthermore, 500  $\mu$ l of lysis buffer were freshly prepared (497  $\mu$ l lysis buffer, 2.5  $\mu$ l dithiothreitol and 0.5  $\mu$ l PMSF).

Cell lysis. Lysis buffer (110  $\mu$ l) was added to each group to create a cell suspension and placed on ice for 30 min. A separation gel and a stacking gel were then prepared. Following lysis, the cells were centrifuged at 3,070xg for 10 min, the supernatant was absorbed and another 1.5-ml Eppendorf tube was prepared to measure the protein concentration and conduct protein quantification. Finally, equal amounts of protein were collected from each sample, 10  $\mu$ l lysis buffer and 20  $\mu$ l 2X sample buffer were added and the solution was incubated at 95°C for 5 min. The electrophoresis 10X running buffer was diluted to 1X, the samples were placed on ice for 5 min, centrifuged at 1,570xg for 1 min and the supernatant was absorbed to prevent trailing. The samples were placed in loading buffer and subjected to sodium dodecyl sulfate-polyacrylamide gel electrophoresis at 80 V for 2 h.

Electrotransfer membranes. In total, 500 ml 1X blotting buffer (100 ml methanol, 50 ml 10X blotting buffer and 350 ml water) were freshly prepared. The gel was cut into 5.5x5 cm pieces, placed in 1X blotting buffer and marked; two pieces of the PVDF mebrane were sheared into 6.5x6 cm pieces and activated in methanol over 5 min. Eight filter papers with two pieces on one side were then placed into 1X blotting buffer to siphon, the PVDF was removed from the methanol, washed once with distilled water and placed into 1X blotting buffer with the gel. In total, four sponges were prepared and soaked in 1X blotting buffer; subsequently, the sponges, two flat pieces of filter paper, the gel (peripheral absorption-desorption 1X blotting buffer) and the flat membranes (moistened with 1X blotting buffer) were placed on a wet-rotating device in turn to compress the filter papers and sponges, followed by placement in a 100 V vertical electrophoresis chamber for 1 h. The chamber was filled with 1X blotting buffer and an ice box was placed along its side.

Protein detection. The capsular membranes were placed into two small containers filled with blocking buffer at 4°C for 8 h to block the non-reactive sites on the membranes in order to inhibit the non-specific absorption of antibodies. The primary goat polyclonal  $\beta$ -actin (1:500) and rabbit polyclonal survivin (1:500) antibodies (Santa Cruz Biotechnology Inc., Santa Cruz, CA) were then diluted with blocking buffer, the opening was sealed with film and the table was agitated horizontally for 1 h to combine the antigens and antibodies. The membranes were washed three times with Tris-buffered saline (TBS) for 15 min each time, the secondary donkey antirabbit antibodies (1:5,000; Santa Cruz Biotechnology Inc.) with horseradish peroxidase were added and the mixture was diluted with blocking buffer. The opening was sealed with film and the table was agitated horizontally for 45 min. Finally, the samples were washed three times with TBS for 15 min per wash and developed in a dark room by X-ray radiography.

# Assessment of Qizhu formula effect on survivin protein expression in MGC-803 cells using RT-PCR

*Grouping*. A 0.1% DMSO group and three medication groups with different concentrations of Qizhu formula (125, 250 and  $500 \mu g/ml$ ) were set up.

Methods

*Cell collection*. First, the MGC-803 cells were collected at the logarithmic phase and added to four  $30\text{-cm}^2$  Petri dishes, with  $4.5 \times 10^6$  cells per dish. The cells were placed in a 5% CO<sub>2</sub>

incubator at 37°C for 24 h, followed by the addition of 6 ml 0.1% DMSO, 125 250, or 500  $\mu$ g/ml Qizhu formula and placement in a 5% CO<sub>2</sub> incubator at 37°C for 24 h.

Total MGC-803 cell RNA extraction. TRIzol one-step extraction was performed according to the manufacturer's instructions as follows: First, the supernatant was removed, 3 ml TRIzol (1 ml/10 cm<sup>2</sup>) were added to each Petri dish, the homogenate samples were repeatedly absorbed and blown 20 times with a 1-ml spearhead and the sample was left to rest for 5 min at 15°C. Second, the supernatant was obtained after a 10-min centrifugation at 10,000 x g at 4°C; the sample was then vigorously agitated for 15 sec following the addition of 0.6 ml trichloromethane (0.2 ml CHCl<sub>3</sub>/1 ml TRIzol) and was kept for 3 min at room temperature. Third, the sample was centrifuged for 15 min at 10,000 x g at 4°C, the upper layer of fluid was transferred to another dry Eppendorf tube, 1.5 ml of isopropanol (0.5 ml isopropanol/1 ml TRIzol) were added, the sample was kept for 10 min at room temperature after blending and centrifuged for 10 min at 10,000 x g at 4°C. Fourth, the supernatant was discarded, the sediment was removed, 3 ml 75% alcohol (1 ml 75% alcohol/1 ml TRIzol) were added and the solution was blended. The supernatant was then discarded by absorption with a spearhead following centrifugation for 5 min at <7,500 x g at 4°C and the sample was dried for 5-10 min at room temperature. Fifth, 20 µl water without RNA enzymes were added and the pellet was dissolved using the blow and beat of the spearhead. The optical density (OD) at 260 and 280 nm was measured using an ultraviolet spectrophotometer following addition of water without RNA enzymes to 2  $\mu$ l RNA to a final volume of 1 ml. Finally, the RNA concentration was calculated according to the formula  $[(OD_{260nm} - OD_{H_2O}) \times H_2O (ml)/template (\mu l)] \times 40 \ \mu g/\mu l$  and the RNA purity to  $OD_{260}/OD_{280}$ . The ranges of extracted RNA  $OD_{260}/OD_{280}$  of 1.78-2.0 were diluted to 1  $\mu g/\mu l$  and kept at -72°C.

RT-PCR was performed according to the instructions of a general RT-PCR kit. Based on the primer design of the gene sequences provided by the gene pools, the primer sequences were as follows through Blast comparison:  $\beta$ -actin gene primer sequence: upstream, 5'-ATCATGTTTGAGACCTTCAACA-3' and downstream, 5'-CATCTCTTGCTCGAAGTCCA-3' with a product length of 318 bp; survivin gene primer sequence: upstream, 5'-CACCGCATCTCTACATTCAAG-3' (97,118) and downstream, 5'-GAAGCAGCCACTGTTACCAG-3' (681,700) with an amplified fragment length of 612 bp.

*RT reaction*. Total RNA (5  $\mu$ g) was transferred to a 0.2-ml centrifuge tube, maintained at 65°C for 10 min and placed in an ice bath following centrifugation for several seconds. We then prepared a 20  $\mu$ l reactive system of table preparation in the centrifugal tube: 1  $\mu$ l solution A [random primer (RP)], 1  $\mu$ l solution B (dNTPs), 4  $\mu$ l solution D (5X RT buffer), 1  $\mu$ l solution E (enzymatic mixed solution), 5  $\mu$ l RNA and mRNA samples, followed by the addition of 8  $\mu$ l solution G to the final volume of 20  $\mu$ l. The sample was left to rest for 10 min at room temperature and the temperature was maintained for 30 min at 37°C, 5 min at 95°C and 4°C for 5 min, respectively. The samples were immediately used or stored at -20°C.

*PCR reaction*. The 50- $\mu$ l PCR sample was prepared as follows: 4  $\mu$ l RT reactive products, 4.5  $\mu$ l solution F (10X PCR buffer), 1  $\mu$ l solution B (dNTPs), 50 pmol upstream primers,



Figure 1. Protein expression of  $\beta$ -actin and survivin in MGC-803 human gastric adenocarcinoma cells. Lane 1, negative control; lane 2, dimethylsulf-oxide control; lane 3, 250  $\mu$ g/ml Qizhu formula; and lane 4, 500  $\mu$ g/ml Qizhu formula.



Figure 2. Functional electrophoretogram of the effect of Qizhu formula on survivin mRNA expression in MGC-803 human gastric adenocarcinoma cells. Lane M, marker; lane 1, 0.1% dimethylsulfoxide; lane 2, 125  $\mu$ g/ml Qizhu formula; lane 3, 250  $\mu$ g/ml Qizhu formula; and lane 4, 500  $\mu$ g/ml Qizhu formula.

50 pmol downstream primers, 1  $\mu$ l solution C (*Taq* enzyme) and solution G to a volume of 50  $\mu$ l. A PCR amplifier was then added following centrifugation for several seconds and the result was finally observed using agarose gel electrophoresis. The amplification parameters were as follows: denaturation for 2 min at 94°C (denaturation at 94°C for 45 sec; renaturation at 55°C for 45 sec; extension at 72°C for 45 sec), amplification for 30 cycles, extension for 5 min at 72°C and storage at 4°C.

Semi-qualitative agarose electrophoresis and analysis. First, ethidium bromide was added to  $10 \,\mu$ l of the PCR products and electrophoresis was conducted on a 2% agarose gel (1 h at 80 V). Images of the electrophoretogram were captured using an FR-980 biological electrophoresis image analysis system (Shanghai Furi Technology Co., Ltd., Shanghai, China) and the results were analyzed using SmartView analysis software (Everett, WA, USA). The survivin/ $\beta$ -actin ratio of each sample was calculated using  $\beta$ -actin as an internal standard. A smaller ratio indicated a lower level of survivin mRNA expression.

## Results

Experimental results of the Qizhu formula on survivin protein expression in MGC-803 cells with western blotting. A gray

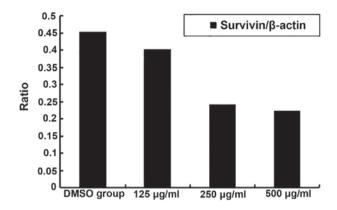


Figure 3. Efects of Qizhu formula on survivin mRNA expression in human MGC-803 gastric adenocarcinoma cells. DMSO, dimethylsulfoxide.

qualitative analysis of  $\beta$ -actin and survivin indicated that the Qizhu formula exerted no explicit effects on the protein expression of the housekeeping  $\beta$ -actin gene; however, it exerted a significant inhibitory effect on the protein expression of the apoptosis-related survivin gene at concentrations of 250  $\mu$ g/ml and, particularly, 500  $\mu$ g/ml (Fig. 1).

Experimental results of survivin mRNA expression in MGC-803 cells with RT-PCR. The effect of Qizhu formula on survivin mRNA expression in MGC-803 human gastric adenocarcinoma cells is shown in Fig. 2. The ratios of survivin/ $\beta$ -actin in the 0.1% DMSO group and in each of the Qizhu formula groups (125, 250 and 500 µg/ml) were 0.4543, 0.4025, 0.2415, and 0.2235, respectively (Fig. 3).

The Qizhu formula was shown to modulate the protein expression of survivin in MGC-803 human gastric adenocarcinoma cells, indicating that this formula acts by downregulating the inhibition of apoptosis of tumor cells. Additionally, Qizhu formula exerted a significant inhibitory effect on survivin mRNA expression in MGC-803 human gastric adenocarcinoma cells in a dose-dependent manner. These results further indicate that Qizhu formula may induce apoptosis of gastric cancer cells by downregulating IAP survivin mRNA expression.

## Discussion

Cell proliferation and apoptosis are essential in maintaining body homeostasis. Tumor cells accelerate cell cycles and downregulate apoptosis-related genes using complex molecular mechanisms, consequently contributing to the malignant proliferation characteristics of tumors. Apoptosis is a key event in the process of tumor growth and tumor cells frequently exhibit defective apoptotic signal transduction pathways.

Previous studies indicated that the high expression of survivin, the most potent IAP currently identified and the smallest of the IAP family, is able to inhibit apoptosis induced by various apoptosis-stimulating factors, such as caspases. Wen *et al* (9) demonstrated that survivin expression in gastric cancer tissue was negatively associated with caspase-3 expression, whereas the changes in the expression levels of caspase-3 in gastric cancer tissues were correlated with the occurrence and development of gastric cancer, indicating that survivin exerts its inhibitory effect on cell apoptosis by inhibiting caspase-3, resulting in tumorigenesis and cancer progression.

Our earlier study (8) also indicated that Qizhu formula not only decreases telomerase activity in MGC-803 human gastric adenocarcinoma cells and protein expression of the telomerase-related hTERT gene, but it can also induce caspase-3, an apoptosis-related protease in its core position.

The present study demonstrated the ability of Qizhu formula to modulate protein and mRNA expression of the survivin gene in MGC-803 cells. The gray qualitative analysis of  $\beta$ -actin and survivin revealed that this formula exerted no explicit effects on the protein expression of the  $\beta$ -actin housekeeping gene, whereas it exerted a significant inhibitory effect on the protein expression of the apoptosis-related survivin gene at concentrations of  $250 \,\mu g/ml$  and, particularly, 500  $\mu$ g/ml. RT-PCR was used to detect the effect of Qizhu formula on survivin mRNA expression in MGC-803 human gastric adenocarcinoma cells. The ratios of survivin/β-actin in the 0.1% DMSO group and in each of the Qizhu formula groups (125, 250 and 500 µg/ml) were 0.4543, 0.4025, 0.2415 and 0.2235, respectively, indicating that Qizhu formula exerted a significant inhibitory effect on MGC-803 survivin mRNA expression in a dose-dependent manner. These results demonstrated that the Chinese medicinal Qizhu formula decreases protein and mRNA expression of the IAP survivin gene, which may represent an additional target pathway via which Qizhu formula intervenes in the process of tumor apoptosis and exerts its antitumor effects.

It was previously demonstrated that the Qizhu formula inhibits the telomerase proliferation of MGC-803 gastric cancer cells and induces caspase-3, an apoptosis-related protease, in the core position and decreases the protein and mRNA expression of the IAP survivin gene (8). Therefore, it may be deduced that Qizhu formula intervenes in gastric cancer through multiple pathways, multiple targets and bidirectional regulation, which further provides objective evidence of the anticancer efficacy of this Chinese medicinal compound.

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