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***Basic Study***

**Guggulsterone induces apoptosis in human hepatocellular carcinoma cells through** **intrinsic mitochondrial pathway**

Shi JJ *et al*. Guggulsterone induces apoptosis in HepG2 cells

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**Abstract**

**AIM:** To investigate the effects of guggulsterone on cell proliferation and apoptosis of human hepatoma HepG2 cells and relevant mechanisms *in vitro*.

**METHODS:** Human hepatocellular carcinoma HepG2 cells and normal human liver L-02 cells were treated with different concentrations of guggulsterone (5-100 μmol/L) for 24-72 h. Cell proliferation was tested by MTT assay. Cell cycle and apoptosis were investigated using flow cytometry (FACS). Bcl-2, Bax mRNA and protein expression were detected by real-time PCR and western bloting. TGF-β1, TNF-α, VEGF contents were determined by ELISA kit.

**RESULTS:** Guggulsterone significantly inhibited HepG2 cells proliferation in a dose-dependent and time-dependent manner. FACS showed that guggulsterone arrested HepG2 cell cycle at G0/G1 phase. Guggulsterone induced apoptosis was also observed in HepG2 cells, with 24.91% ± 2.41% and 53.03% ± 2.28% apoptotic percentage in response to the treatment of 50 μmol/L and 75 μmol/L guggulsterone respectively. Bax mRNA and protein expression were significantly increased and Bcl-2 mRNA and protein expression were decreased. ELISA analysis showed concentration of TGF-β1 and VEGF were significantly decreased and TNF-α concentration was increased.

**CONCLUSIONS:** Guggulsterone exerts its anticancer effect by inhibiting cell proliferation and inducing apoptosis in HepG2 cells and guggulsterone induces apoptosis by activation of the intrinsic mitochondrial pathway.

**Key words:** Guggulsterone; Hepatocellular carcinoma cells; Apoptosis; Cell cycle; Mitochondrial pathway

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**Core tip:** Hepatocellular carcinoma (HCC) is the fifth most prevalent cancer and the second leading cause of tumor mortality worldwide. Guggulsterone (GS) is a phytosterol extracted from the gum resin of guggul plants that shows pro-apoptotic effect has been found to play a pivotal role in its anti-carcinogenic mechanisms. In this study, we investigated the anticancer effects of GS-induced apoptosis in human HCC cells and the underlying molecular mechanisms. Our results demonstrated that GS induced HepG2 cells apoptosis through regulating Bcl-2 and Bax expression levels.

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**INTRODUCTION**

Hepatocellular carcinoma (HCC) is the fifth most prevalent cancer and the second leading cause of tumor mortality worldwide[1]. The number of new cases of HCC are increasing year by year and about 748000 new cases are being diagnosed annually, accounting for 9.2% of all new global tumor cases[2,3]. HCC has a low resectability rate, high recurrence rate, insidious onset, rapid progression, grave prognosis and high mortality[1], therefore development of anticancer drugs with explicit efficacy on HCC has now become a challenge worldwide.

Many phytochemicals derived from edible plants have shown cancer therapeutic potential[4,5]. Guggulsterone (GS) is a phytosterol extracted from the gum resin of guggul plants that shows strong biological properties of antioxidant, anti-inflammatory, hypolipidemic and hypocholestremic[6-9], and has been used for in treatment for several malignant diseases[10-13]. Pro-apoptotic effect of GS has been found to play a pivotal role in its anti-carcinogenic mechanisms[10,13].

Apoptosis is known as programmed cell death, accompanying specific morphological and biochemical changes such as cell shrinkage, nuclear and DNA fragmentation, apoptotic body formation[14,15]. These changes are associated with a distinct set of signaling pathways including the intrinsic mitochondrial pathway and the extrinsic death receptor pathway[16]. A common step is the caspases activation in both pathways[17]. The mitochondrial pathway is activated by a variety of stimulus such as heat shock, DNA damage and reactive oxygen to increase permeability of mitochondrial outer membrane and release cytochrome C into the cytoplasm. When the presence of ATP, the cytochrome C is combined with apoptosis protease-activating factor (Apaf-1) that is combined with caspase recruitment domain and caspase-9 precurs or to activate caspase-9, leading to activation of caspase-3 and caspase-7[14]. Bcl-2 family, which plays an important role in the mitochondrial pathway, contains anti-apoptotic proteins including Bcl-2, Bcl-XL, Mcl-1 and pro-apoptotic proteins such as Bax, Bad and Bek, in which Bcl-2 and Bax are more important proteins [16].

Previous studies have shown that GS can induce human colon cancer HT-29 cells apoptosis through decreasing anti-apoptotic proteins expression including Bcl-2, cIAP-1, cIAP-2 and increasing the Bid protein expression[10]. Moreover, GS has been found to activate JNK signaling pathway in human prostate cancer cells, which subsequently upregulates the Bcl-2 family members including Bax and Bak, leading to the apoptosis of cancer cells[18,19]. Several studies have indicated that GS enhanced the sensitivity of HCC cell line Hep 3B and Hep G2 to TRAIL-induced apoptosis *via* ROS-dependent ER stress induction[12,20,21]. However, it has not been determined whether GS has anti-HCC effect in other signaling pathways, such as the intrinsic mitochondrial pathway.

In this study, we investigated the anticancer effects of GS-induced apoptosis in human HCC cells and the underlying molecular mechanisms. Our results demonstrated that GS induced HepG2 cells apoptosis through regulating Bcl-2 and Bax expression levels.

**MATERIALS AND METHODS**

***Reagents and antibodies***

Z-guggulsterone (Z-GS) was purchased from ENZO (USA) and was dissolved in Dimethyl sulfoxide (DMSO) (Sigma-Aldrich, St Louis, MO) as a 16 mmol/L stock solution and stored at -20℃. Various concentrations of Z-GS (0-100 μmol/L) were diluted in serum free RPMI1640 medium (HyClone, Utah, United States) with 0.5% (v/v) DMSO was used as a vehicle control. 0.25% (w/v) trypsase were obtained from Hyclone (Utah, United States). The rabbit monoclonal antibodies against Bcl-2, Bax and β-actin were purchased from Santa Cruz Biotechnology (CA, United States). Horseradish peroxidase conjugated goat anti-rabbit and goat anti-mouse were purchased from ABGENT Biotechnology (SD, United States).

***Cell lines and cell culture***

Human HCC cell line HepG2 and the normal human hepatic cell line L-02 were obtained from the Experimental Center of Xi’an Jiaotong University. Cells were cultured in RPMI1640 medium supplemented with 10% (v/v) fetal bovine serum (FBS, HyClone, Utah, United States), 100 unites/ml penicillin (Sigma-Aldrich, St Louis, United States) and 100 μg/ml streptomycin (Sigma-Aldrich, St Louis, United States) in a humidified atmosphere of 95% (v/v) air and 5% (v/v) CO2 at 37 ℃. Culture medium was changed every other day. When cells covered 80%-90% of the bottom of culture flasks, cell were washed twice with phosphate buffered saline (PBS, 137 mmol/L NaCl, 2.7 mmol/L KCl, 4.3 mmol/L Na2HPO4, 1.4 mmol/L KH2PO4, pH 7.4) and then were digested by 0.25% (w/v) trypsase. Cells were harvested using RPMI1640 medium followed by centrifugation at 1000rpm for 10 min. Cells were re-suspended in RPMI1640 medium that were plated in appropriate plates at appropriate density and serum-starved for 24 h using serum free RPMI1640. Then the cells were treated with RPMI1640 medium containing various concentrations of Z-GS. After 24, 48 or 72 h of culture, cells were harvested as usual.

***MTT assay***

Cell viability was tested using 3-(4,5-dimethylthiazol-2 -yl)-2,5-diphenyltetr azolium bromide (MTT, Sigma-Aldrich, St Louis, MO) assay as described previously[22]. Briefly, cells were plated in 96-well plates at a density of 6 × 103cells/well followed by starvation for 24 h using serum free RPMI1640 culture medium. The culture medium were then replaced with RPMI1640 medium containing various concentrations of Z-GS (0-100 μmol/L). After 24, 48, and 72 h of culture, 20 μl of MTT solution (5 mg/ml) was added to each well and cells were continuously cultured for 4 h. Culture medium was then removed and 150 μl of DMSO was added to each well. After shaking the culture plates for 5 min, the solution was collected and the optical density (OD) was measured using a spectrophotometer (ND-1000, Thermo Fisher, United States) at wavelength of 570nm. The cell viability rate (%) = (OD treated/ODcontrol) × 100%.

***Cell cycle analysis***

The logarithmic phase HepG2 cells and L-02 cells were plated in 6-well plates at a density of 6 × 105cells/well and incubated a humidified atmosphere of 95% (v/v) air and 5% (v/v) CO2 at 37 ℃ for 24 h. Cells were then treated with 50 μmol/L and 75 μmol/L Z-GS in RPMI1640 medium for 24 h. After washing with cold PBS twice, cells were fixed in 70% (v/v) ice-cold ethanol overnight at -20 ℃. Cells were treated with Tris-HCl buffer (10 mmol/L Tris-HCl, pH 7.5) containing 1% (w/v) RNase A (Sigma-Aldrich) for 15 min followed by incubation with popidium iodide (PI, Sigma-Aldrich) for 15 min. Cell cycles were then analyzed using flow cytometer (CALIBUR, BD, United States), and the outputs were processed using ModFit LT2.0 software (Verity Software House, United States).

***Apoptosis assay***

Cell apoptosis was determined using the Annexin V-FITCand PI double staining (KaiJi, NanJing, China) as previously described[22]. Briefly, HepG2 cells and L-02 cells were plated in 6-well plates at 6 × 105 cells/well and treated with various concentrations of Z-GS (0 μmol/L, 50 μmol/L and 75 μmol/L) for 24 h. Cells were then harvested and stained with Annexin V-FITC and PI in the cold binding buffer (50 mmol/L HEPES, 700 mmol/L NaCl, 12.5 mmol/L CaCl2, pH 7.4) for 15 min at room temperature in the dark. After washing, cell apoptosis was analyzed using flow cytometer (CALIBUR, BD, United States).

***Real-time PCR analysis***

Total RNA was extracted using Trizol reagent (Invitrogen, United States) as described previously[23]. 1 μg of RNA was then used to syntheses complementary DNA (cDNA) with AMV Reverse Transcriptase (RT) kit (TAKARA, Japan) according to the manufacturer’s protocols. The relative expression of Bcl-2 and Bax were analyzed by quantitative real-time polymerase chain reaction (PCR) with SYBRPremix Ex TaqⅡkit (TAKARA, Japan) with GAPDH as an internal control. The reaction system contained 12.5 μl of 2 × SYBR Ex TaqⅡ, 2 μl RNA reaction, 1 μl each of forward and reverse primers and 8.5 μl sterile distilled water in a final volume of 25 μl. The reaction was performed at 95 ℃ for one cycle for 30 s, 95 ℃ for 5 s and 60 ℃ for 60 s for 40 cycles, 72 ℃ for 60 s. Table 1 shows all the primer sequences. PCR products were run on 2% (w/v) agarose gel (Sigma-Aldrich, United States) and stained with ethidium bromide (Sigma-Aldrich, United States). RT-PCR outputs were detected using Bio-Rad iQ5 software (CA, United States).

***Western blotting analysis***

Bcl-2 and Bax protein expression was investigated using Western blotting analysis as described previously[23]. Briefly, cells were harvested and re-suspended in RIPA lysis buffer [20 mmol/L Tris, 150mM NaCl, 1% (v/v) Triton X-100, 1% (w/v) digestive phosphatase inhibitors, 1% (w/v) protease inhibitors, 1% (w/v) phenylmethyl sulfonylfluoride (PMSF), pH 7.5] (Sigma-Aldrich). Protein concentration was determined using BCA assay (Kangweishiji, BeiJing, China) according to the manufacture’s protocols. Equal amounts of protein (30 μg/lane) were separated in a 10% (w/v) sodium dodecyl sulfate-polyacrylamide (SDS-PAGE) gel (Sigma-Aldrich), and were then electrotransferred onto polyvinylidenedifluoride (PVDF) membranes (Sigma-Aldrich). After blocking with 5% (w/v) bovine serum albumin (BSA, Sigma-Aldrich) in Tris-buffered saline (TBS, 0.1 mol/L, pH 7.4), membranes were incubated with primary antibodies (1:1000 dilution) overnight at 4 ℃. After washing, membranes were incubated with horseradish peroxidase-conjugated secondary antibodies (1:5000 dilution) for 1 hour at room temperature with agitation. All membranes were detected using the ECL Westren Blotting Kit Reagent (Thermo Fisher, United States).

***Cytokine detection***

Cell supernatants were collected from Z-GS treated and untreated control cells and filtered through a 0.22 μm filter (Millipore, United States). The levels of TGF-β1, TNF-α and VEGF in the culture medium were determined using enzyme-linked immunosorbent assay kit (ELISA, eBioscience, United States) accoriding to the manufacture’s protocols.

***Statistical analysis***

Data are presented as mean ± SD, and tested for normality andequal variance. Student's *t* test or One-way analysis of variance (ANOVA) plus Bonferroni's post-test was carried out using SPSS 15.0 software (SPSS Inc, United States). Differences were considered significant at *P* values of less than 0.05.

**Results**

***Guggulsterone*** ***inhibits the*** ***viabilityof******human HCC cells***

Compared with the control group, the viability of HepG2 and L-02 cells were significantly decreased in a dose- and time-dependent manner in response to 5, 10, 25, 50, 75 and 100 μmol/L of GS treatment for 24 h (Figure 1A). Interestingly, 25, 50, 75 and 100 μmol/L of GS significantly reduced the HepG2 cell viability when compared with that in the L-02 cells after 24 h treatment (Figure 1B), suggesting a less sensitivity to GS in L-02 cells relative to the HepG2 cells. Similary results were revealed after 48 and 72 h treatment (data not shown). The half maximal inhibitory concentration (IC50) of GS treatment in HepG2 cells for 24 h was 75 μmol/L. Therefore, 50 μmol/L and 75 μmol/L GS were used for all subsequent experiments in HepG2 cells.

***Effect of*** ***Guggulsterone on cell cycle*** ***in human HepG2 and L-02 cells***

After treatment by 75 μmol/L of GS for 24 h, there was an increase in G0/G1 fraction (62.88% ± 2.67% *vs* 44.02% ± 1.07%) but decrease in G2/M fraction of HepG2 cells (13.33% ± 1.84% *vs* 28.33% ± 1.25%) in comparison with the untreated control cell (*p* < 0.05, Figure 2A). However, 50 μmol/L GS treatment did not induced a significant difference in cell cycle fractions in HepG2 cells when compared with the control group (*p* > 0.05, Figure 2A). Defferently, neither 75 μmol/L nor 50 μmol/L of GS treatment induced significant alteration in L-02 cell cycles (*p* > 0.05, Figure 2B).

***Guggulsterone induced apoptosis in human HepG2 cells***

After treatment by 50 μmol/L and 75 μmol/L GS for 24 h, the percentage of apoptotic cells in HepG2 cells was 24.91% ± 2.41% and 53.03% ± 2.28% respectively, significantly higher than that in the untreated control cells (5.18% ± 1.74, *P* < 0.01) (Figure3A). However there was no significant defferences in L-02 cell apoptosis in response to 50 μmol/L and 75 μmol/L GS treatment (*p* > 0.05, Figure 3B).

***Guggulsterone induced apoptosis through activation of intrinsic mitochondrial signaling pathway***

There was significantly increase in Bax mRNA and protein levels in HepG2 cells after treated by 50 μmol/L and 75 μmol/L of GS for 24 and 48 h respectively (Figure 4A). Differently, Bcl-2 mRNA and protein contents in HepG2 cells were significantly reduced in response to 50 μmol/L and 75 μmol/L of GS treatment (Figure 4B). However, neither 50 μmol/L nor 75 μmol/L induced significant alterations in Bax (Figure 4C) or Bcl-2 expression (Figure 4D) in L-02 cells (*p* > 0.05).

***Effect of Guggulsterone on*** ***TGF-β1, TNF-α and VEGF*** ***cytokines***

After treatment by 35, 50 and 75 μmol/L of GS for 24 h, TGF-β1 and VEGF levels in HepG2 cells were significantly decreased when compared with the control group (*p* < 0.01, Figure 5A and B). Differently, TNF-α contents in HepG2 cells were significant increased in a dose dependent manner in response to the treatment of 35, 50 and 75 μmol/L of GS when comapred with the untreated cells (Figure 5C). Interestingly, there was no signicant difference in TGF-β1 (Figure 5D), VEGF (Figure 5E) and TNF-α levels (Figure 5F) in L-02 cells after treatment by 35, 50 and 75 μmol/L of GS for 24 h.

**Discussion**

GS has been used for an anticancer agent due to its cell proliferation inhibition and pro-apoptosis effects in tumer cells[10,18]. However, whether GS can be used in HCC treatment is still largely unknown. In this study, cell proferation of HCC cell line HepG2 were significantly inhibited by GS in a dose- and time-dependent manner. More importantly, cell proliferation in normal hepatocytic cell line L-02 was less affected by GS when compared with HepG2, suggesting that normal hepatocytes are significantly more resistant to growth inhibition by GS compared with HCC cells. Therefore, we investigated whether GS treatment has a selective activity to human HCC cells and normal liver cells by conducting experiments on the effects of GS on cell cycles and apoptosis inducing of L-02, a normal human hepatocyte cell line.

G0/G1 fractions of cell cycles were increased but G2/M fractions were decreased in HepG2 cells in response to the treatment of GS, indicating that GS arrested HepG2 cell cycle in G0/G1 phase, and thereby inhibited HCC cell proliferation. The precise mechanism is not clear, however previous studies performed in other tumor cells have illustrated that GS regulates cell cycle *via* downregulated expression level of cell cycle regulatory protein cyclin D1 and induced expression of cyclin dependent kinase inhibitor P21WAF1/CIP1 and P27[24,25].

Furthermore, we found that GS induced HepG2 cell apoptosis in a dose-dependent way, which is also consistent with previous studies[10,24,26]. Our further investigation indicated that GS increased Bax but decreased Bcl-2 gene and protein expression in HepG2 cells, illustrating that the intrinsic mitochondrial pathway was involved in the pro-apoptosis effect of GS. A variety of stress stimuli including growth factors withdrawal, heat shock, and oxidative damage have been shown to activate the apoptosis intrinsic or mitochondrial pathway[27-29]. TGF-β is a multifunctional cytokine involved in the regulation of many cells apoptosis and is implicated in the pathogenesis of human diseases, including carcinogenesis. The VEGF family plays a pivotal role in tumor angiogenesis and is responsible for solid tumor growth and metastasis[29-31]. We therefore investigated the effects of GS on the levels of TGF and VEGF which have been confirmed to be involved in the carcinogenesis during HCC occurring and development. As expected, TGF-β1 and VEGF concentrations were significantly decreased in response to the treatment of GS in a dose dependent manner, indicating that these two growth factors may were negatively invloved in the HCC cell apoptosis. The underlying mechanism is still unknown, however previous studies have shown that the low expression of their receptors on the cell sufrace induced by GS may be involved[32,33].

Accumulating evidence has suggested that GS has chemopreventive and chemotherapeutic potential for cancer treatment. Although the underlying mechanisms are not fully understood, these studies including ours, clearly indicated that anticancer activity of GS is associated with apoptosis induction. Specifically, GS treament on cancer cells induces the alteration and regualtion of Bcl-2 gene family proteins, NF-kB signaling, MAPK pathways, farnesoid X receptor, and the EGFR-STAT3 signaling, tec. However, these resuts are largely from other cancer cells. It is still largely unknown whether GS has similar effects on HCC cells. In the present study, for the first time, we showed that GS treatment significantly inhibited cell growth, induced cell cycle arrest and caused apoptotic cell death in human HCC cells. Our results revealed a novel anti-HCC mechanism of GS, in which the mitochondrial pathway pathway is involved in the GS induced apoptosis in human hepatocarcinoma.

Interestingly, GS treatment did not induce significant cell cycle arrest and apoptosis in L-02 cells, which may explaine the reason why a normal human hepatocyte cell line is more resistant to the growth inhibition by GS as described above. Unsurprisingly, our further investigation showed less alterations in mitochondrial pathway protein bax and Bcl-2, as well as growth factors/cytokines including TGF-β1, VEGF and TNF-α levels in L-02 cells in response to the treatment of GS. There resutls indicated that normal hepetocytes are more resistant to the growth inhibition by GS when compared with the HCC cells, which is consistent to previous resutls in the other cancer cells[34]. The mechanism is largely unkonwn, however, uncharacterized constituent(s) of GS may interact additively or synergistically to inhibit the viability of the caner cells.

In conclusion, we demonstrated GS inhibited the viability of human HCC cells by regulated cell cycle, induced tumor cell apoptosis by activation of the mitochondrial pathway, and decreased TGF-β1and VEGF but increased TNF-α levels. These results suggests that GS influences HCC phenotypes by inhibiting cell proliferation, promoted cell death and regulating carcinogenesis-related growth factors, and thereby is a potential anticancer agent for the treatment of human HCC.

**comments**

***Background***

Hepatocellular carcinoma (HCC) is the fifth most prevalent cancer and the second leading cause of tumor mortality worldwide. Guggulsterone (GS) is a phytosterol extracted from the gum resin of guggul plants that shows strong biological properties of antioxidant, anti- inflammatory, hypolipidemic and hypocholestremic, and has been used for in treatment for several malignant diseases. Apoptosis is known as programmed cell death, which is associated with a distinct set of signaling pathways including the intrinsic mitochondrial pathway and the extrinsic death receptor pathway. Previous studies have shown that GS can induce different cancer cell apoptosis through activation of the extrinsic death receptor pathway. However, it has not been determined whether GS has anti-HCC effect in other signaling pathways, such as the intrinsic mitochondrial pathway.

***Research frontiers***

Due to the strong biological properties of antioxidant, anti-inflammatory, hypolipidemic and hypocholestremic of GS, it has been used for in treatment for several malignant diseases icluding HCC. Pro-apoptotic effect of GS has been found to play a pivotal role in its anti-carcinogenic mechanisms, however its precise mechanism is not fullly understood. Therefore, the relationship between GS and cell apoptosis becomes an important research areas in this field.

***Innovations and breakthroughs***

In this study, the authors demonstrated that GS inhibited the viability of human HCC cells by regulated cell cycle, induced tumor cell apoptosis by activation of the mitochondrial pathway, and decreased TGF-β1 and VEGF but increased TNF-α levels. These results suggests that GS influences HCC phenotypes by inhibiting cell proliferation, promoted cell death and regulating carcinogenesis-related growth factors, and thereby is a potential anticancer agent for the treatment of human HCC. This is the first systematical study demonstrating that GS induces HCC cell apoptosis through activation of by activation of the mitochondrial pathway, and evidence shown here expands our knowledge of the precise mechanism how GS inhibit HCC development.

***Applications***

The study results suggest that the GS induces HCC apoptosis and could be a potential treatment drug for this commonly prevalent tumor in humans in the future.

***Peer- review***

Other reports have shown that GS is pro-apoptotic and modulates various genes. These authors show that GS arrested HepG2 cycle at G0/G1 phase. Bax mRNA was increased together with other changes, suggesting the intrinsic mitochondrial pathways in GS effects. The study indicates that GS is apotential anticancer therapy for HCC.

**Refreences**

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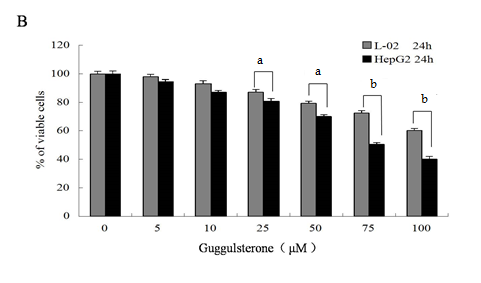
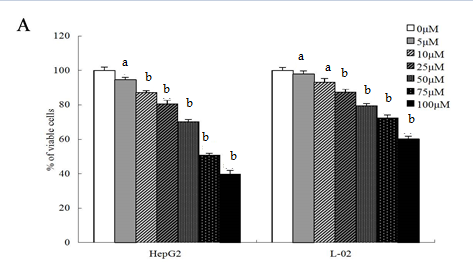
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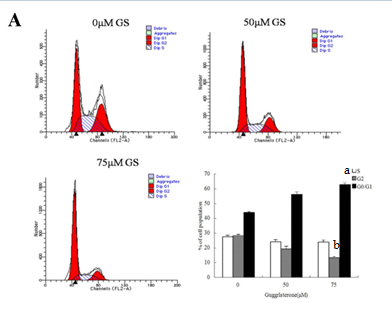
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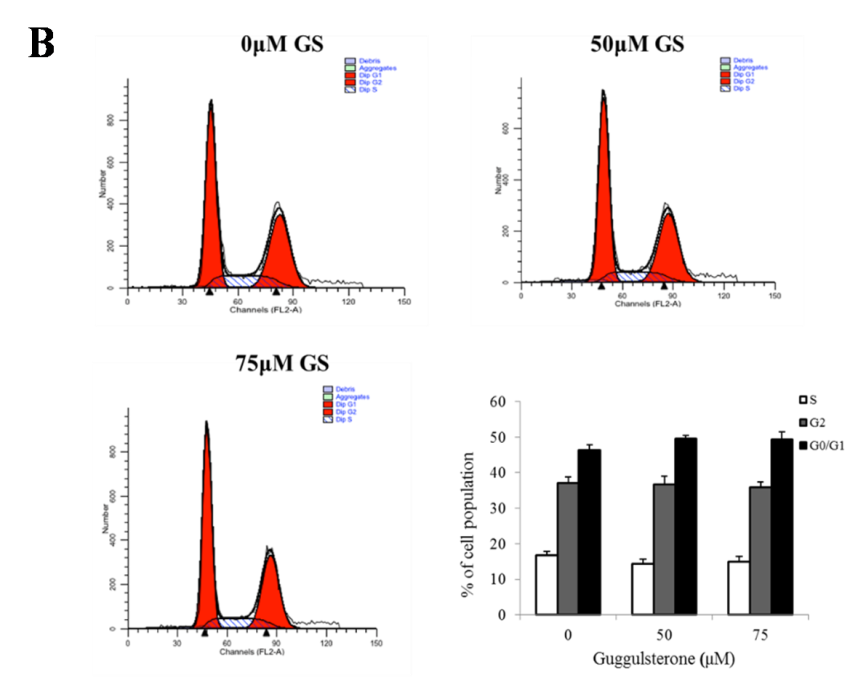
**Table 1 Primers used for real-time PCR analysis**

|  |  |  |
| --- | --- | --- |
| **Name** | **Forward primer (5’-3’)** | **Reverse primer (5’-3’)** |
| Bcl-2 (168 bp) | TGTGTGGAGAGCGTCAAC | GGAGAAATCAAACAGAGGC |
| Bax (164 bp) | ATGCGTCCACCAAGAAGC | CCAGTTGAAGTTGCCGTC |
| GAPDH (138 bp) | GCACCGTCAAGGCTGAGAAC | TGGTGAAGACGCCAGTGGA |

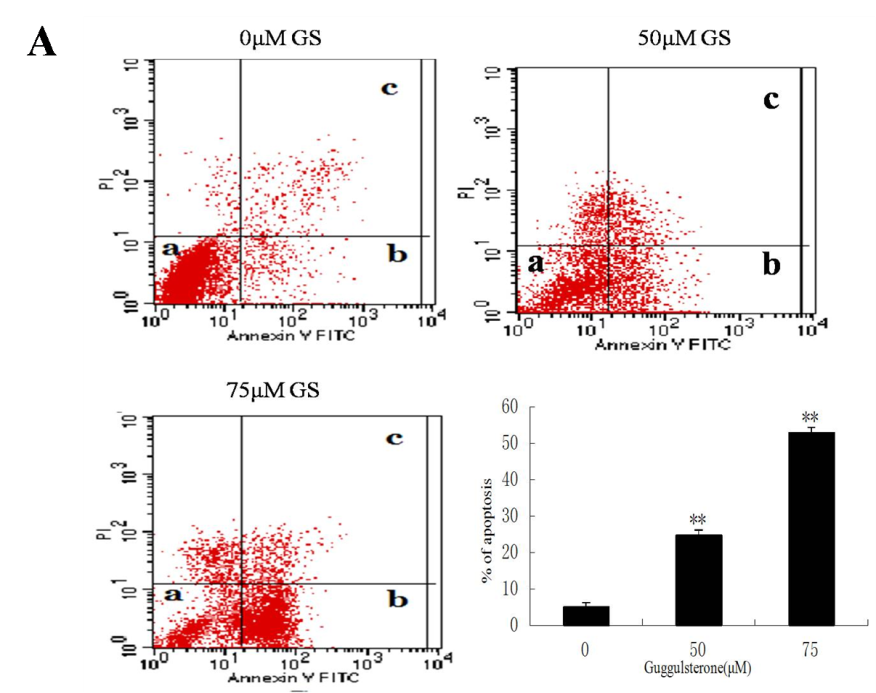


**Figure 1 Human L-02 and HepG2 cells showed different cell viabilities in response to Guggulsterone treatment.** Cell viability in L-02 and HepG2 cells was determined using MTT assay after incubation with 5, 10, 25, 50, 75 and 100 μmol/L guggulsterone respectively for 24 h. Guggulsterone signficnatly reduced cell viability in both L-02 and HepG2 cells in a dose- and time-dependent manner (A). Same concentration of guggulsterone differently reduced cytotoxicity between L-02 and HepG2 cells (B). a*p* < 0.05 or b*p* < 0.01, *vs* the control group.



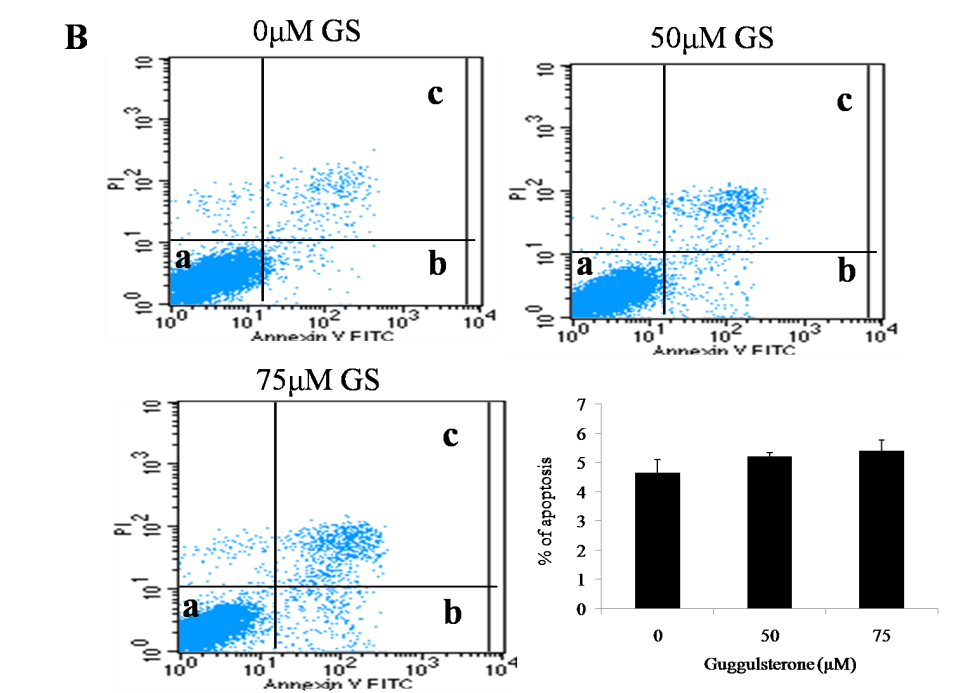
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**Figure 2 Guggulsterone altered cell cycle inhuman HepG2 and L-02cells.** Human HepG2 (A) and L-02 cells (B) were treated with 0 μmol/L, 50 μmol/L and 75 μmol/L of GS for 24 h and the cell cycle was analyzed using flow cytometry respectively. The histogram showed mean % of cell population in each phase of cell cycle. a*p* < 0.05 or b*p* < 0.01, *vs* the untreated control group.

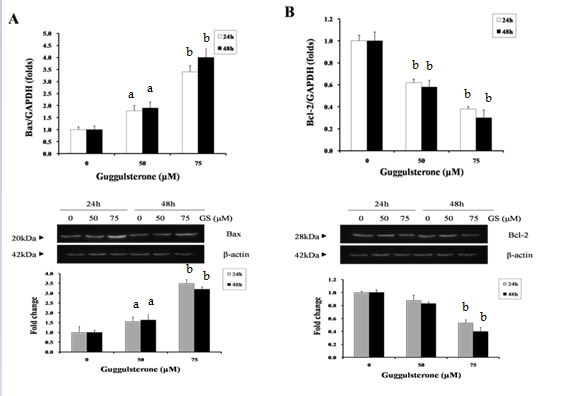


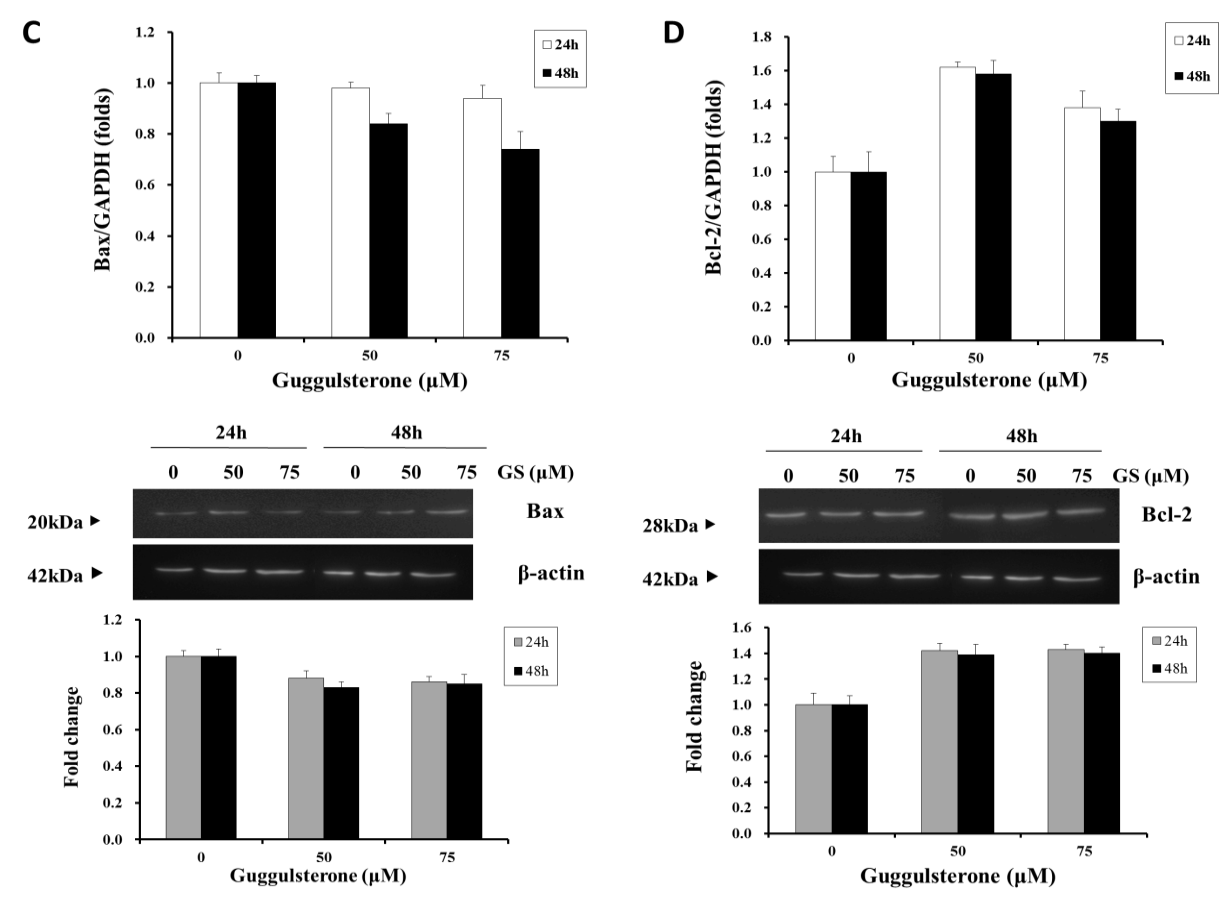
b

b

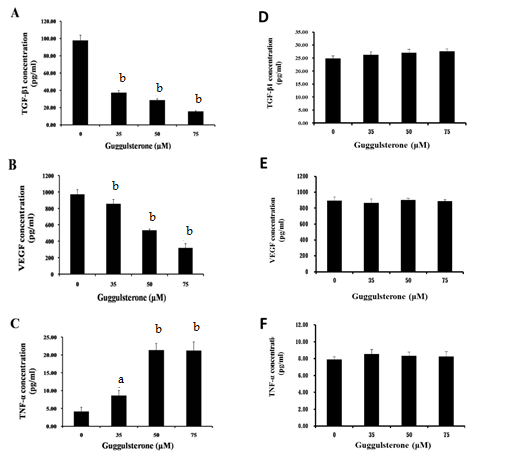


**Figure 3 Guggulsterone induced apoptosis in human HepG2 and L-02 cells.** Human HepG2 (A) and L-02 cells (B) were treated with 0μM, 50μM and 75μM of GS for 24 hours and cell apoptosis was investigated using annexin V-FITC and PI. GS significantly increased HepG2 cell apoptosis with a dose-dependent manner. b*p* < 0.01, *vs* the control group.



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**Figure 4** **Guggulsterone differently altered Bax and Bcl-2 expression in HepG2 cell.**HepG2 (A and B) and L-02 cells (C and D) were treated with 0μM, 50μM and 75μM of GS for 24 and 48 hours respectively. Bax and Bcl-2 mRNA and protein levels were investigated using real-time PCR and Western blotting respectively. a*p* < 0.05 or b*p* < 0.01, *vs* the control group.



**Figure 5 Guggulsterone altered TGF-β1, TNF-α and VEGF levels in HepG2 and L-02 cells.** HepG2 (A, B, and C) and L-02 cells (D, E, and F) were treated with 0, 35, 50 or 75 μmol/L of GS for 24 h respectively. TGF-β1, TNF-α and VEGF levels in the culture medium were investigated using ELISA. a*p* < 0.05 or b*p* < 0.01, *vs* the control group.