Original Article

Annonaceous acetogenins reverses drug resistance of human hepatocellular carcinoma BEL-7402/5-FU and HepG2/ADM cell lines

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Abstract: Hepatocellular carcinoma (HCC) is the most common tumor in worldwide and chemotherapy resistant is a severe obstacle in HCC treatment. Annonaceous acetogenins was a nature compound from Uvaria accuminata and it has show the anti-tumor proliferation activity in many types cancer. In this study, we showed that annonaceous acetogenins is correlated with the drug resistance reversal in human hepatocellular carcinoma BEL-7402/5-FU and HepG2/ADM cell lines. We found that cell apoptosis was improved and cell cycle was arrested, further, multidrug-resistance proteins such as MDR1, MRP1, Topo-IIα, GST-π, cyclin D1, Survivin and bcl-2 are down-regulated, however, intracellular Rh-123 and caspase-3/8 was up-regulated by Annonaceous acetogenins treatment. We also found that there was a decreased activity of NF-κB and Akt in Annonaceous acetogenins treatment groups. Therefore, we demonstrate that Akt/NF-κB pathway was involved in Annonaceous acetogenins reverses drug resistance of human hepatocellular carcinoma cells.

Keywords: Annonaceous acetogenins, hepatocellular carcinoma, drug resistance

Introduction

Hepatocellular carcinoma (HCC) is one of the most common cancers and now it is the third leading cause of cancer deaths in worldwide [1]. The incidence of hepatocellular carcinoma is highest in Asia and Africa, with an estimated half million new cases diagnosed per year globally, and more than 50% of whom are in China [2]. Hepatocellular carcinoma is generally less sensitivity to chemotherapeutic agents and approximately 67% hepatoma patients treated with chemotherapeutic agents develop rapidly acquired resistance [3]. Multi-drug resistance (MDR) refers to the tumor cells with anticancer drug resistance. MDR is one of the major obstacles in HCCs successful chemotherapy treatment [4]. Thus, development of new chemotheraputic drugs reversal of MDR is essential in treating MDR of HCC.

Variety of natural products have been shown to be excellent and reliable sources for pharmaceutical development, such as Paclitaxel and Curcumin [5, 6]. Annonaceous acetogenins (ACGs) are a class of natural polyketides isolated from Annonaceous plants growing in tropical and subtropical regions, which show favorable efficacy in variety of human cancer cell lines in vitro [7]. They have been reported to potently inhibit the activity of NADH-ubiquinone oxidoreductase of the mitochondrial electron transport system [8]. Annonaceous acetogenins phytochemical and pharmacological studies have further revealed a novel therapeutic role as an anticancer agent for these natural products.

Here, we have utilized drug-resistant human hepatocellular carcinoma BEL-7402/5-FU and HepG2/ADM cell lines to investigate the potential reversal drug resistance of human hepatocellular carcinoma, and the nature of the mechanism responsible for reversal multi-drug resistance.

Materials and methods

Cell lines and culture

Human hepatocellular carcinoma cell line Hep-G2 and BEL-7402 were obtained from the

Table 1. Primer lists of real-time PCR

Gene Sequences (5'-3') Tm (°C) MRD1 Forward: GAGCCCATCCTGTTTGACTG 60 Reverse: GCTGCCCTCACAATCTCTTC 58 MRP1 Forward: GTGGAATTCCGGAACTAC 58 Reverse: CGGAGGTCGTGCAGGCCG 60 Top-IIα Forward: GGCTCGATTGTTATTTCCAC 60 Reverse: GGTTGTAGAATTAAGAATAGG 58 GST-π Forward: CTGGAAGGAGGAGGT GGTG 58 Reverse: GACGCAGGATGGTATTGGAC Cyclin D1 Forward: CAA TGACCCCGCACGATTTC 60 Reverse: CATGGAGGGCGGATTGGAA GAPDH Forward: GGAGCGAGATCCCTCCAAAAT 60 Reverse: GAAGATGGTGATGGGATTTC 60 60			
Reverse: GCTGCCCTCACAATCTCTTC MRP1 Forward: GTGGAATTCCGGAACTAC 58 Reverse: CGGAGGTCGTGCAGGCCG Top-IIα Forward: GGCTCGATTGTTATTTCCAC 60 Reverse: GGTTGTAGAATTAAGAATAGG GST-π Forward: CTGGAAGGAGGAGGT GGTG 58 Reverse: GACGCAGGATGGTATTGGAC Cyclin D1 Forward: CAA TGACCCCGCACGATTTC 60 Reverse: CATGGAGGGCGGATTGGAA GAPDH Forward: GGAGCCAGAGATCCCTCCAAAAT 60	Gene	Sequences (5'-3')	Tm (°C)
MRP1 Forward: GTGGAATTCCGGAACTAC 58 Reverse: CGGAGGTCGTGCAGGCCG 60 Top-IIα Forward: GGCTCGATTGTTATTTCCAC 60 Reverse: GGTTGTAGAATTAAGAATAGG 58 GST-π Forward: CTGGAAGGAGGAGGT GGTG 58 Reverse: GACGCAGGATGGTATTGGAC 60 Cyclin D1 Forward: CAA TGACCCCGCACGATTTC 60 Reverse: CATGGAGGGCGGATTGGAA 60 GAPDH Forward: GGAGCGAGATCCCTCCAAAAT 60	MRD1	Forward: GAGCCCATCCTGTTTGACTG	60
Reverse: CGGAGGTCGTGCAGGCCG Top-IIα Forward: GGCTCGATTGTTATTTCCAC 60 Reverse: GGTTGTAGAATTAAGAATAGG GST-π Forward: CTGGAAGGAGGAGGT GGTG 758 Reverse: GACGCAGGATGGTATTGGAC Cyclin D1 Forward: CAA TGACCCCGCACGATTTC 60 Reverse: CATGGAGGGCGGATTGGAA GAPDH Forward: GGAGCGAGATCCCTCCAAAAT 60		Reverse: GCTGCCCTCACAATCTCTTC	
Top-IIα Forward: GGCTCGATTGTTATTTCCAC 60 Reverse: GGTTGTAGAATTAAGAATAGG GST-π Forward: CTGGAAGGAGGAGGT GGTG 58 Reverse: GACGCAGGATGGTATTGGAC Cyclin D1 Forward: CAA TGACCCCGCACGATTTC 60 Reverse: CATGGAGGGCGGATTGGAA GAPDH Forward: GGAGCGAGATCCCTCCAAAAT 60	MRP1	Forward: GTGGAATTCCGGAACTAC	58
Reverse: GGTTGTAGAATTAAGAATAGG GST-π Forward: CTGGAAGGAGGAGGT GGTG Reverse: GACGCAGGATGGTATTGGAC Cyclin D1 Forward: CAA TGACCCCGCACGATTTC Reverse: CATGGAGGGCGGATTGGAA GAPDH Forward: GGAGCGAGATCCCTCCAAAAT 60		Reverse: CGGAGGTCGTGCAGGCCG	
GST-π Forward: CTGGAAGGAGGAGGT GGTG Reverse: GACGCAGGATGGTATTGGAC Cyclin D1 Forward: CAA TGACCCCGCACGATTTC Reverse: CATGGAGGGCGGATTGGAA GAPDH Forward: GGAGCGAGATCCCTCCAAAAT 60	Top-IIα	Forward: GGCTCGATTGTTATTTCCAC	60
Reverse: GACGCAGGATGGTATTGGAC Cyclin D1 Forward: CAA TGACCCCGCACGATTTC 60 Reverse: CATGGAGGGCGGATTGGAA GAPDH Forward: GGAGCGAGATCCCTCCAAAAT 60		Reverse: GGTTGTAGAATTAAGAATAGG	
Cyclin D1 Forward: CAA TGACCCCGCACGATTTC 60 Reverse: CATGGAGGGCGGATTGGAA GAPDH Forward: GGAGCGAGATCCCTCCAAAAT 60	GST-π	Forward: CTGGAAGGAGGAGGT GGTG	58
Reverse: CATGGAGGGCGGATTGGAA GAPDH Forward: GGAGCGAGATCCCTCCAAAAT 60		Reverse: GACGCAGGATGGTATTGGAC	
GAPDH Forward: GGAGCGAGATCCCTCCAAAAT 60	Cyclin D1	Forward: CAA TGACCCCGCACGATTTC	60
		Reverse: CATGGAGGGCGGATTGGAA	
Reverse: GAAGATGGTGATGGGATTTC	GAPDH	Forward: GGAGCGAGATCCCTCCAAAAT	60
		Reverse: GAAGATGGTGATGGGATTTC	

American Type Culture Collection (ATCC, Manassas, VA, USA) and mainained in DMEM medium. 5-fluorouracil (5-FU)-selected drugresistant BEL-7402/5-FU and adriamycin resistant HepG2/ADM cell line was obtained from Nanjing Keygen Biotech. Co. (Nanjing, China). Cells were cultured in RPMI-1640 containing 10% fetal bovine serum (FBS), 100 U/ml penicillin and 100 $\mu g/ml$ streptomycin at in 37°C with 5% CO $_2$ incubator.

MTT assay

BEL-7402/5-FU and HepG2/ADM cell lines were plated in 96-well plates respectively. After over-night incubation, cells were treated with various concentrations of Annonaceous acetogenins. After treated for 72 h, 20 µL MTT solution was added to each well. Incubate in 37°C for 4 h, the supernatant was re moved and 100 µL DMSO was added to each well. Samples were then shaken for 15 min. The optical density (OD) was read a t the wavelength of 540 nm. All experiments were performed in triplicates. Each treatment was performed in sextuplicate and each experiment was repeated three times. The half-maximal inhibitory concentration (IC50) was defined as the concentration that caused 50% inhibition of cell proliferation.

Reverse transcriptase PCR

Total RNA was extracted from cells using Trizol Reagent according to the manufacturer's protocol. RNA was reverse-transcribed using reverse Transcription System (Promega, USA). Primers were obtained from Sangon Biotech (shanghai,

China) and the primers sequences were listed in **Table 1**. The amount of PCR product formed in each cycle was evaluated on the basis of SYBR Green I fluorescence. All amplification reactions were performed using the ABI7500 PCR system with the following cycle conditions: one cycle at 95°C for 5 min, followed by 40 cycles at 95°C for 30 sec, 60°C for 30 sec, and72°C for 30 sec. The mRNA level for e ach gene was normalized to GAPDH.

Apoptosis assay in vitro

Cells were incubated by 5-fluorouracil for BEL-7402/5-FU cells and adriamycin for HepG2/ADM cells with Annonaceous acetogenins for 24 hr. The percentage of apoptosis was determined by AnnexinV-FITC Apoptosis Detection kit (BD Biosciences) following the manufacture's instruction. Briefly, cells were washed with ice-cold PBS and resuspended in binding buffer. Annexin V-FITC and PI (BD Biosciences) were added to cells and incubated at room temperature in the dark for 15 min, and then the cells were analyzed by BD FACS can flow cytometer with 488 nm excitation wavelength and 525 nm emission wavelength.

Cell cycle distribution analysis

The cells were seeded in 6-wellplates (4×10^5) well). For synchronization experiment, cells were treated with 100 ng/mL of nocodazole for 24 h at 37°C prior to beng treated with 10 μM Annonaceous acetogenins. Following 24 h treatment, floating cells were combined with attached cells, then subjected to centrifugation at 3,000 rpm for 5 min at 4°C. The cell pellets were washed with cold PBS, then fixed in ethanolat -20°C. The cells were washed with PBS, rehydrated and resus-pended in 0.2 mL of RNase A in PBS buffer at 37°C for 30 min. The cells were stained with 0.3 mL propidium iodide in a solution containing 0.1% Triton X-100, 0.1 mM EDTA. Samples were analyzed on a BD FACS can flow cytometer and the percentage of cells in the S, GO-G1, and G2-M phases of the cell cycle was determined using WinMDI 2.8.

Western blot analysis

Treated cells (3 \times 10 6) were harvested by cell scraping, washed once in cold PBS and lysed in 300 μ l RIPA lysis buffer (50 mM Tris pH = 8.0, 150 mM NaCl, 0.1% SDS, 1% NP40) containing

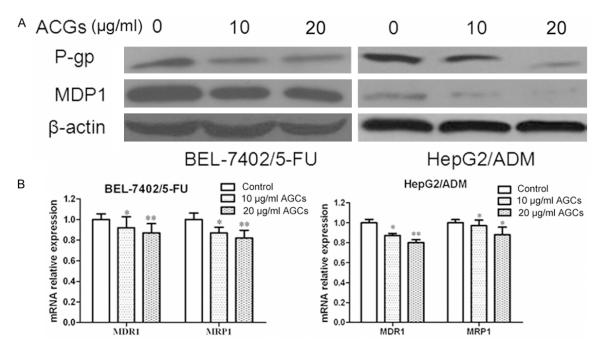


Figure 1. Inhibitory effect of Annonaceous acetogenins on expression of MDR1 and MRP1. A. Western blot analysis of protein extracts obtained from BEL-7402/5-FU and HepG2/ADM cells; B. mRNA expressions of MDR1 and MRP1 genes in BEL-7402/5-FU and HepG2/ADM cells. Data presented are means \pm SD values, N = 3. Bars indicate SD. Significant differences from control were indicated by P<0.05 (*) and P<0.01 (**).

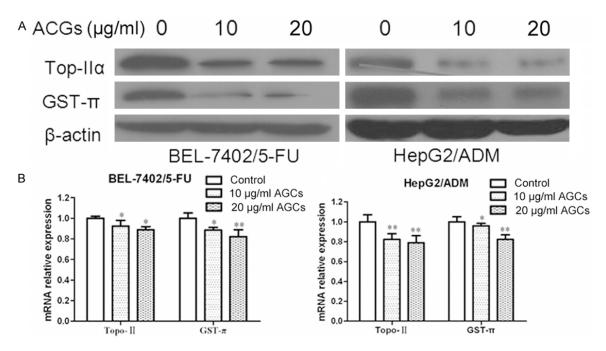
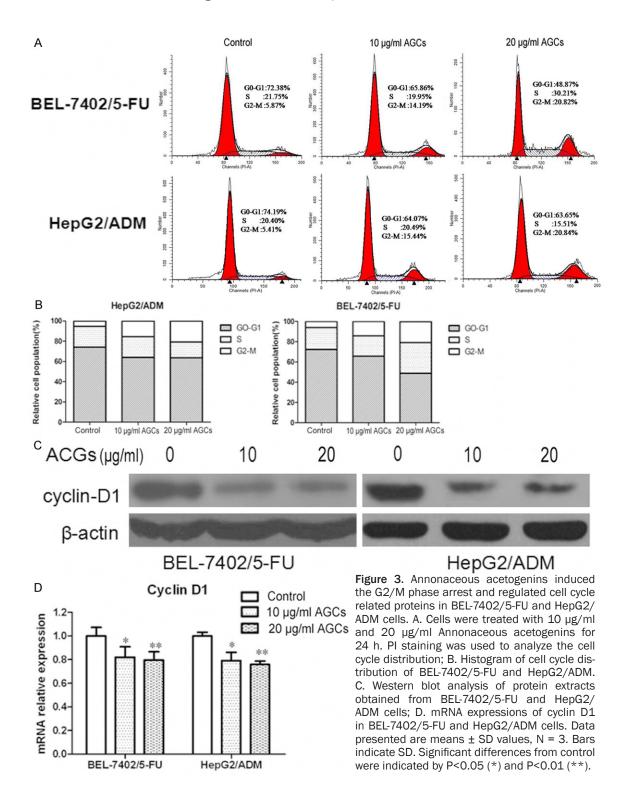


Figure 2. Inhibitory effect of Annonaceous acetogenins on expression of Topollα and GST- π . A. Western blot analysis of protein extracts obtained from BEL-7402/5-FU and HepG2/ADM cells; B. mRNA expressions of Topollα and GST- π genes in BEL-7402/5-FU and HepG2/ADM cells. Data presented are means \pm SD values, N = 3. Bars indicate SD. Significant differences from control were indicated by P<0.05 (*) and P<0.01 (**).

one tablet of Protease Inhibitor Cocktail (Roche Applied Sciences). Lysates were sonicated and

centrifuged at 12,000 r/min at 4°C for 10 min. Protein concentrations were determined by



bicinchoninic acid (BCA) protein assay kit. Per lane 40 µg of whole cell lysate was separated using 10% SDS-acrylamide gels, and transferred on PVDF membranes (Amersham Biosciences) blocked by incubation with 5% bovine serum albumin in tween-tris-buffered saline

(TTBS) at room temperature for 1 hr. Following incubated overnight at 4°C with different primary antibodies [MDR1, 1:500; MDP1, 1:500; Topo-II α , 1:200; GST- π , 1:200; BcI-2, 1:400; Cyclin D1, 1:500; Survivin, 1:400 (all obtained from Santa Cruz); Cleaved caspase-3, 1:1000;

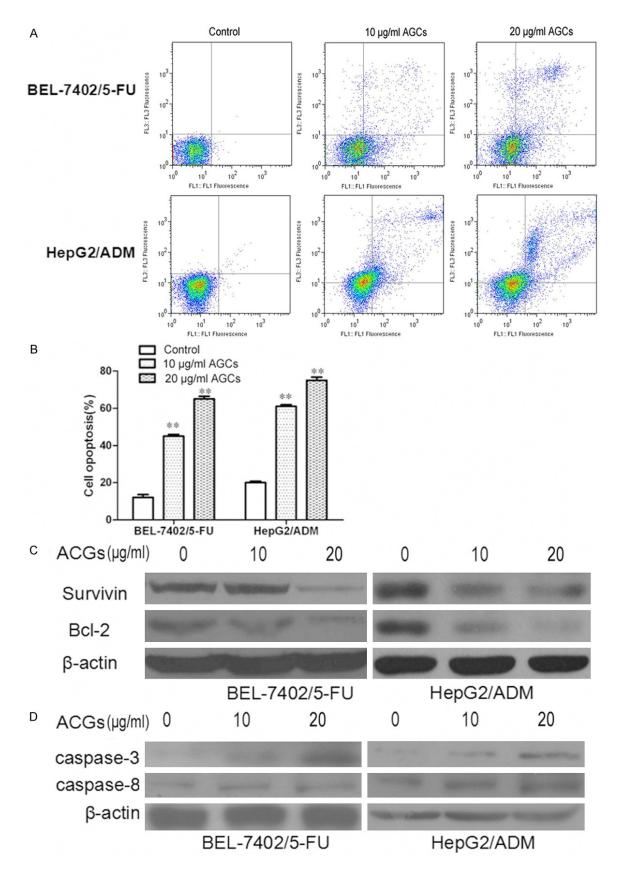


Figure 4. Annonaceous acetogenins induced cell opoptosis in BEL-7402/5-FU and HepG2/ADM cells. A. Cells were treated with 10 μg/ml and 20 μg/ml Annonaceous acetogenins for 24 h. Annexin V-FITC/PI analysis by FCM was

used to analyze the percentage of apoptotic cells in BEL-7402/5-FU and HepG2/ADM cells; B. Histogram of the percentage of apoptotic cells in BEL-7402/5-FU and HepG2/ADM cells. C. The expression of survivin and Bcl-2 proteins were determined by western blot in BEL-7402/5-FU and HepG2/ADM cells; D. mRNA expressions of survivin and Bcl-2 in BEL-7402/5-FU and HepG2/ADM cells. Data were presented as the mean \pm SD and were representative of an average of three independent experiments per concentration. Significant differences from untreated control are indicated by P<0.05 (*) and P<0.01 (**).

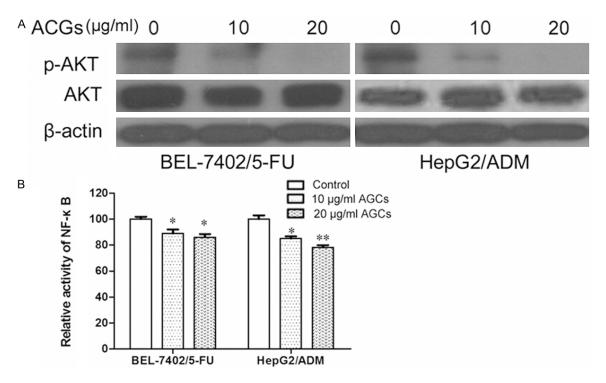


Figure 5. Annonaceous acetogenins affect on AKT/NF-κB Signal Pathway Molecules in BEL-7402/5-FU and HepG2/ADM cells. A. The total cell lysate was separated by SDS-PAGE gel followed by Western blot with anti-p-Akt antibody, anti-total-Akt antibody. The experiment shown is epresentative of three independent experiments with similar results; B. Cells were transfected with pNF-κB-luc and the transcriptional activation was detected by luciferase reporter assay. Data were presented as the mean \pm SD and were representative of an average of three independent experiments per concentration. Significant differences from untreated control are indicated by P<0.05 (*) and P<0.01 (**).

Cleaved caspase-8, 1:1000 (from Sigma); β -actin, 1:5,000 (from Sigma)] in 5% fat-free milk prepared in TBST. After rinsing three times in TBST, filters were incubated with secondary antibodies for 1 h at room temperature in TBST containing 1% fat-free milk. After washing, the blots were visualized by horseradish peroxidase (HRP) Western Blot Detection System.

Rhodamine-123 exclusion assay

The cells were seeded in 6-wellplates and incubated with Annonaceous acetogenins for 24 h. They were trypsinized and resuspended at 1 \times 10 $^6/\text{mL}$, then 10 μ l of the substrate Rh-123 (1 mg/ml) were added to the samples and the cells were incubated for 20 min at 37°C. After

washed twice and resuspended in 0.5 ml of PBS, the samples were then immediately detected by flow cytometry. Data were analyzed using FCS Express V3 (De Novo Software).

NF-kB DNA binding activity assay

NF-κB p65 activity was assayed by reporter gene system according to the instruction manual with moderate modification. Briefly, the assays were performed in a 96-well plate. The cells (1.5 to 2×10^6) were seeded into each well in triplicate for 24 h, then transfected each well of the cells with 0.1 μg pGL4.32 [luc2P/NF-kB-RE/Hypro] plasmids (Promega). After 24 h of transfection, the luciferase activity was analyzed by Bright-GloTM Luciferase Assay System (Promega).

Statistical analysis

Quantitative data are presented as the mean \pm SD determined from the indicated number of experiments. Statistical analysis was based on one-way ANOVA for comparison of multiple groups.

Result

Cytotoxicity of annonaceous acetogenins

The viability of BEL-7402/5-FU and HepG2/ ADM cells were treated with Annonaceous acetogeninsat different concentrations (0, 1, 2, 5, 10, 20, 50 and 100 μ M) for 72 h was determined by MTT assay. Annonaceous acetogenins could inhibited the growth of BEL-7402/5-FU and HepG2/ADM cell lines in a dose-dependent manner in vitro, IC50 of BEL-7402/5-FU cells was 73.7 µg/ml and IC50 of HepG2/ADM cells was 83.7 µg/ml by Annonaceous acetogenins. The results also showed that 10 µg/ml Annonaceous acetogenins was not cytotoxic (inhibition rate < 5%) and 20 µg/ml Annonaceous acetogenins was weakly cytotoxic (inhibition rate 10-15%) for all of BEL-7402/5-FU and HepG2/ADM cell lines. Thus, treatment concentrations of 10 µg/ml and 20 µg/ml Annonaceous acetogenins were chosen to study reversal effect of Annonaceous acetogenins on MDR in BEL-7402/5-FU and HepG2/ADM cells. We found that the IC50 of BEL-7402/5-FU cells by 5-fluorouracil was 311.4 µg/ml and IC50 of BEL-7402/5-FU cells with by 5-fluorouracil 10 µg/ml and 20 µg/ml Annonaceous acetogenins were 22.7 µg/ml and 12.3 µg/ml, the reversal fold (RF = IC50 of MDR cell/IC50 of treatment cell) were 13.7 fold and 25.3 fold, meanwhile, the IC50 of HepG2/ADM cells by adriamycin was 75.7 µg/ml and IC50 of HepG2/ADM cells with by adriamycin 10 µg/ml and 20 µg/ml Annonaceous acetogenins were 7.6 µg/ml and 5.7 μ g/ml, the reversal fold (RF = IC50 of MDR cell/IC50 of treatment cell) were 9.7 fold and 13.3 fold.

Annonaceous acetogenins downregulated the expression of MDR1 and MRP1

MDR frequently results from the incongruous expression of drug-resistant genes, such as multi-drug-resistance gene (MDR1). MDR1 and MRP1 are two major membrane transporter proteins that involved in efflux activities and lead to multiple drug resistance [9]. So we would like to investigate whether Annonaceous

acetogenins could downregulate the expression of these proteins to restore the intercellular accumulation of Rhodamine-123. RT-PCR and western blotting results showed that, oppose to the high expression level observed in the parental cells, the MDR1 and MRP1 were decreased in BEL-7402/5-FU and HepG2/ADM cells (Figure 1). Rh-123 is a substrate for P-gp and commonly used as an indicator for evaluating the activities of P-gp. Thus, the activity of the P-gp drug pump can be gauged by the degree of intracellular accumulation of Rodamine-123, which can in turn be determined by the measurement of intracellular fluorescence [10]. So rhodamine-123 accumulation was detected by flow cytometry. As expect, Annonaceous acetogenins could increased the intracellular accumulation of rhodamine-123 (result did not show).

Annonaceous acetogenins reverse the MDR by inhibition of other MDR related genes expression

Except MDR1 and MRP1, other MDR related genes might also get involved in MDR of HCC. Topoisomerase IIα (Topo IIα) is an essential nuclear enzyme with a role in the maintenance of DNA topology. Topo II α is a target for several anticancer drugs and the activity of topoisomerase $II\alpha$ is important in the development of drug resistance [11]. GST-π is a multifunctional enzyme that plays a critical role in cellular detoxification by catalyzing the conjugation of reduced glutathione to hydrophobic electrophilic compounds and may influence mutagenesis and carcinogenesis [12]. In addition, GST-π is the most important phase II drug-metabolizing enzyme and is involved in the metabolism and detoxification of environmental carcinogens and chemotherapeutics [13]. We next treated BEL-7402/5-FU and HepG2/ADM cells with Annonaceous acetogenins at the concentration of 10 µg/ml and 20 µg/ml for 24 h. Western blot and Real time PCR was used to test genes expression. Topoisomerase $II\alpha$ (TopoII α) and glutathione S-transferase π (GST-π) genes, were observed down-regulated in BEL-7402/5-FU and HepG2/ADM cells after Annonaceous acetogenins incubation (Figure 2).

Annonaceous acetogenins affect cell cycle distribution and downreagulated the expression of cyclin D1

Cell cycle phase distribution was detected by flow cytometry to determine whether there is

any difference in cell cycle kinetics between BEL-7402/5-FU and HepG2/ADM cells and their parental cells. BEL-7402/5-FU and HepG2/ADM cells incubated in Annonaceous acetogenins at the concentration of 10 μ g/ml and 20 μ g/ml for 24 h resulted in had a significant increase in G2/M phase and decrease in the G0/G1 phase compared with control group phase.

Cyclin D1 function as regulators of CDK4 or CDK6, is required for cell cycle G1/S transition, is very important in cell cycle progression [28]. To search for the indication of mechanisms involved in cell cycle,we would like to investigate whether Annonaceous acetogenins could affect cyclin D1 experssion. Real-time PCR and western blotting results showed that, the expression level of cyclin D1 in BEL-7402/5-FU and HepG2/ADM cells obviously decreased compared to their parental cells. These data suggested that Annonaceous acetogenins affect cell cycle distribution may by inhibition of cyclin D1 expression (Figure 3).

Annonaceous acetogenins induced cell apoptosis and downreagulated the expression of Survivin and Bcl-2

To investigate whether the growth inhibition of Annonaceous acetogenins was caused by apoptosis, annexin V-FITC/PI double staining was employed. The percentage of apoptotic cells was 13.5% and 21.3% in Control group of BEL-7402/5-FU and HepG2/ADM cells, respectively. When exposed to 10 μ g/ml and 20 μ g/ml of Annonaceous acetogenins for 24 h, the percentage increased to 45.9% and 65.5%, 61.4% and 75.8%, respectively. In summary, these results demonstrated that Annonaceous acetogenins treatment raised the apoptosis percentage of BEL-7402/5-FU and HepG2/ADM cells.

Survivin, a member of the family of inhibitor of apoptosis proteins, functions as a key regulator of mitosis and programmed cell death. Survivin expression is also highly regulated by the cell cycle and is only expressed in the G2-M phase [14]. Bcl-2 family proteins regulate and contribute to programmed cell death or apoptosis. Bcl-2 suppresses apoptosis in a variety of cell systems including factor-dependent lymphohematopoietic and neural cells [15]. Regulates cell death by controlling the mitochondrial membrane permeability. We are in great interest

to know if these proteins were downreagulated by Annonaceous acetogenins in BEL-7402/5-FU and HepG2/ADM cells. The western blotting results showed that these proteins obviously decreased when treated with Annonaceous acetogenins. Real-time PCR analyses results were in concord with those from western blotting (Figure 4).

Annonaceous acetogenins inhibits AKT/NFκB signaling pathway in BEL-7402/5-FU and HepG2/ADM

It has been reported that activation of NF-κB followed by Akt phosphorylation plays a role in regulation of cell survival, apoptosis and drug resistance [16]. We wondered if intracellular AKT/NF-kB signaling pathway was involved in the action of Annonaceous acetogenins. To determine whether Annonaceous acetogenins could alleviate drug resistance via the AKT/ NF-kB signaling pathway, BEL-7402/5-FU and HepG2/ADM cells were cultured in the presence of Annonaceous acetogenins, Then expressions of the proteins AKT and NF-kB were evaluated. Results indicated that Annonaceous acetogenins could decrease phosphorylation of AKT, but didn't affect Total-Akt expression. And the expression levels of NF-kB were also decreased by ACGs (Figure 5). These results suggest that Annonaceous acetogenins induced apoptosis results in inactivation of NF-kB via Akt/NF-kB pathway.

Discussion

Hepatocellular carcinoma is the fifth most common malignancy worldwide with a poor long time survival. Multidrug resistance (MDR) is one of the major obstacles in the chemotherapeutic treatment of many human cancers [17]. Although, people have been devoted to search for effective and clinically applicable MDR modulators in the past three decades and various mechanisms have been proposed to explain the multidrug resistance of hepatocellular carcinoma, they are complicated and remain unclear [18]. According to reports, the mechanism of resistance to chemotherapy may be associated with the classical drug-resistant pathway, which is mediated by P-gp, and the inhibition on cell apoptosis, are the two mostly investigated mechanisms in research for MDR of tumor. To find more effective, low-toxicity modulators to inhibit MDR become more and more urgent. In this study, we found that drug resistance of BEL-7402/5-FU and HepG2/ADM cells were reversed by Annonaceous acetogenins.

Annonaceous acetogenins (ACGs) are a class of natural polyketides isolated from Annonaceous plants, which show significant cytotoxicity to human cancer cell lines in vitro. However, reports on whether ACGs can reverse thechemo therapeutic drug resistance in hepatocellular carcinoma are rare. In order to investigate the reversal effect of ACGs on BEL-7402/5-FU and HepG2/ADM cells, we chose the weakly cytotoxic concentrations of 10 µM. Firstly, cell viability was detected by MTT. The results showed that the proliferation of drug-resistant cell line BEL-7402/5-FU and HepG2/ADM was significantly inhibited after cells were pretreated with ACGs. FCM results showed that ACGs could promote cell apoptosis in drug-resistant cell line BEL-7402/5-FU and HepG2/ADM cells. Western blotting results showed that the expression of P-gp, MRP1, cyclin D1, Bcl-2, Survivin, NF-kB, phosphorylation of Akt were decreased in drug-resistant cell line BEL-7402/5-FU and HepG2/ADM compared with their control group. Based on this, we further investigated the potential molecule mechanism of reversing the resistance effect of chemotherapeutic drug resistance hepatocellular carcinoma cell line BEL-7402/5-FU and HepG2/ADM by Annonaceous acetogenins, possibly it is via decreasing the function of P-gp and the mitochondrial apoptosis pathway.

P-glycoprotein (P-gp), an ATP-binding cassette (ABC) transporter, which is a 170-kD plasma membrane glycoprotein encoded by the MDR1 gene, is capable of reducing the intracellular accumulation of drugs by excluding a drug outside the cells [19]. P-gp can induce drug resistance due to decrease in the cellular chemotherapeutic drug under effective concentration, through pumping the chemotherapeutic agents out of the cells against the concentration. Its up-regulation helps cancer cells to augment cell survival against the drug and resist the cytotoxic effects of anti-cancer agents. MRP1 is another major membrane transporter protein that involved in efflux activities and lead to multiple drug resistance [20]. Moreover, the ubiquitous NF-kB transcription factor has been reported to induce drug resistance in cancer

cells. The inhibition of NF-kB reduced MDR1 mRNA and P-glycoprotein expression in HCT15 cells [21]. The NF-kB pathway responds actively to MDR1 induction due to its activation by the generation of reactive oxygen species, the activation of IkB kinase, and the degradation of IkB [22]. Besides, NF-kB is bound at nucleotide position-159 of the MDR1 promoter mediating the transcription of MDR1, which suggest NF-kB in the regulation of the MDR1 gene expression in cancer cells and in drug resistance. In our study, the MDR1 and MRP1 expression both in protein and in mRNA level detected by western blot and RT-PCR, and down regulations of MDR1, MRP1 and NF-κB were exhibited after treated with Annonaceous acetogenins. Rh-123 is a substrate of P-gp and can be used as an indicator for testing the P-gp functional activities. The intracellular Rh-123 fluorescence was increase in BEL-7402/5-FU and HepG2/ADM cells treated with ACGs, which perhaps due to the down-regulation of P-gp pump. Thus, we can conclude that ACGs reverses MDR by inhibiting the cellular efflux function of P-gp and the expression of MDR1 and MRP1 both in transcription and translation. Apart from MDR1 and MRP1, other molecules may involved in the tumor resistance. So we also detected the expression levels of Topo-IIα and GST-π after treated with Annonaceous acetogenins.

Another mechanisms of drug resistance refer to insensitivity to drug-induced apoptosis. Cell cycle is closely related with the tumor cell proliferation. There is considerable evidence that the anticancer agent suppressed cell cycle progression in different phases and then induced cell apoptosis. In our experiment, Annonaceous acetogenins (ACGs) could caused cell cycle arrest in G2-M phase, accompanied with a decrease in the GO/G1 phase compared with control group. We also confirmed that ACGs induced apoptosis BEL-7402/5-FU and HepG2/ ADM cells when treated with 10 µg/ml and 20 µg/ml ACGs for 24 h. These results implied that ACGs induce the apoptosis of BEL-7402/5-FU and HepG2/ADM cells by inhibiting cell cycle progression to reverse drug resistance.

Cyclin D is a member of the cyclin protein family that is involved in regulating cell cycle progression. Suppression of cyclin D1 levels has been shown to potentiate the drug sensitivity of cancer cells [23]. Survivin is the smallest member

of the Inhibitor of Apoptosis (IAP) gene family and downregulation of survivin induce apoptosis in tumor cells. Bcl-2, is the founding Bcl-2 family of regulator proteins that regulate apoptosis. Most cellular Bcl-2 is bound to the outer mitochondrial membrane. The down-regulation of Bcl-2 can induce the release of cytochrome-C (cytosol) from mitochondria, trigger the activation of caspases and finally cause cell apoptosis [24], caspase-3 is an important regulating gene and plays a key role in apoptosis, which cleaves and activates caspases 6, 7, and interacts with caspase-8 and caspase-9. Sequential activation of caspases plays a central role in the execution-phase of cell apoptosis [25]. In our experiment, we found that the expression of cyclin D1 and Bcl-2 in BEL-7402/5-FU and HepG2/ADM was significantly lower than their parental cells when treated with ACGs. Suggesting that the mitochondrial receptor pathway may be involved in the formation of drug resistance. ACGs could reverse the chemotherapeutic drug resistant of BEL-7402/5-FU and HepG2/ADM via down-regulation of Cyclin D1, Survivin and Bcl-2 levels.

It is well established that cyclin D1 is a NF-κB target gene. NF-kB can upregulate expression of cyclin D1, thus promoting Rb hyperphosphorylation and cell cycle progression. As known, down-regulation of anti-apoptotic protein is one of the mechanisms that NF-kB takes part in apoptosis and induces apoptosis. The activation of NF-B followed by Akt phosphorylation plays a role in the development of insulin resistance. Akt is also involved in the regulation of NF-kB and apoptosis [26]. Constitutive activation of NF-kB induced by Akt may play a major role in the chemoresistant cells [27]. We postulated that the intracellular AKT/NF-kB signaling pathway was involved in the action of Annonaceous acetogenins. To see whether Akt and NF-kB are the targets of ACGs, their activation in BEL-7402/5-FU and HepG2/ADM cells treated with ACGs was measured. The results showed that Akt phosphorylation and NF-kB transcriptional activation were inhibited by ACGs. And NF-kB was attenuated by Akti-1/2 (an AKT inhibitor) which implies that Akt may play a role in Annonaceous acetogenins. This suggests that ACGs reverses MDR through suppression of the Akt/NF-kB pathway.

In conclusion, Annonaceous acetogenins can effectively reverse the chemotherapeutic drug

resistant of BEL-7402/5-FU and HepG2/ADM. The potential mechanisms refer to multiple aspects including inhibition of cell growth, induction of apoptosis and cell cycle arrest, accumulation of rhodamine-123 and decreased expression of P-gp, MRP1, Cyclin D1, Survivin and Bcl-2. Meanwhile, the inhibition of the Akt/NF-kB pathway is a key role in how ACGs functions.

Disclosure of conflict of interest

None.

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