

RESEARCH ARTICLE

Alkylglyceronephosphate Synthase (AGPS) Alters Lipid Signaling Pathways and Supports Chemotherapy Resistance of Glioma and Hepatic Carcinoma Cell Lines

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Abstract

Chemotherapy continues to be a mainstay of cancer treatment, although drug resistance is a major obstacle. Lipid metabolism plays a critical role in cancer pathology, with elevated ether lipid levels. Recently, alkylglyceronephosphate synthase (AGPS), an enzyme that catalyzes the critical step in ether lipid synthesis, was shown to be up-regulated in multiple types of cancer cells and primary tumors. Here, we demonstrated that silencing of AGPS in chemotherapy resistance glioma U87MG/DDP and hepatic carcinoma HepG2/ADM cell lines resulted in reduced cell proliferation, increased drug sensitivity, cell cycle arrest and cell apoptosis through reducing the intracellular concentration of lysophosphatidic acid (LPA), lysophosphatidic acid-ether (LPAe) and prostaglandin E2 (PGE2), resulting in reduction of LPA receptor and EP receptors mediated PI3K/AKT signaling pathways and the expression of several multi-drug resistance genes, like MDR1, MRP1 and ABCG2. β -catenin, caspase-3/8, Bcl-2 and survivin were also found to be involved. In summary, our studies indicate that AGPS plays a role in cancer chemotherapy resistance by mediating signaling lipid metabolism in cancer cells.

Keywords: AGPS - Lipid metabolism - cancer chemotherapy resistance

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Introduction

The hallmarks of cancer are keys for cancer treatment. The most fundamental hallmark of cancer cells is their ability to sustain proliferation (Marquardt et al., 2012). This trait is the target of chemotherapy agents, an old but still widely used class of cancer therapeutic medicine. These drugs could be divided into several types. One type is called alkylating agents, the typical drug is cisplatin (Srivastava et al., 2010). Another type is topoisomerase inhibitors, with doxorubicin is the typical one (Pommier et al., 2010). Cisplatin and doxorubicin are extensively used to treatment different types of cancer. However, resistance to these drugs is rather common.

Reprogramming of energy metabolism is an emerging hallmark of cancer in the last decade. Cancer cells reprogram the cellular energy metabolism in order to support continuous cell growth and proliferation (Morandi et al., 2014). One well known such alteration is the Warburg effect (Vander et al., 2009). Recent research suggests that alterations in metabolism could not be viewed as just a response to cell proliferation, but

metabolites themselves can be oncogenic by altering cell signaling and blocking cellular differentiation (Weinstein et al., 2014). Targeting cancer cell metabolism is hence an interesting strategy. Here we focus on the abnormal lipids metabolism. Lipids are not just structural components of cell but can also have important roles in signaling. Deregulation of lipide metabolism is common in cancer and can affect numerous cellular processes, including cell growth and proliferation (Prabhu et al., 2013). Many lipid synthesis related genes have been considered involved in tumor procedure, for example that prostaglandin-endoperoxide synthase 2 (PTGS2) and phospholipase A2 group IIA (PLA2G2A) genes encode enzymes involved in arachidonic acid and prostaglandin biosynthesis, which is associated with carcinogenesis (Liu et al., 2014). Recently, alkylglyceronephosphate synthase (AGPS), an enzyme that catalyzes the critical step in ether lipids synthesis, is shown to up-regulate in multiple types of cancer cell and primary tumor. AGPS alters the balance of structural and signaling lipids to fuel cancer pathogenicity (Benjamin et al., 2013). Nonetheless, whether AGPS is involved in chemotherapy resistance has remained unclear.

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In this study, we showed that AGPS elevated in glioma U87MG/DDP and hepatic carcinoma HepG2/ADM cell lines. Knockdown of AGPS restored the drug sensitivity in these cells. We also showed that silencing of AGPS led to reduced cellular level of lysophosphatidic acid (LPA), lysophosphatidic acid-ether (LPAe) and prostaglandin E2 (PGE2), which could fuel aggressive and progress of cancer (Hull et al., 2004; Brindley et al., 2013). Knockdown of AGPS also results in down-regulation of multi-drug resistance proteins and anti-apoptosis proteins.

Materials and Methods

Cell culture and establishment of drug-resistant cell clones

U87MG and HepG2 cell lines were obtained from Cell Bank of Chinese Academy of Science (Shanghai, China) and maintained at 37°C in a humidified atmosphere with 5% CO₂. The culture medium was DMEM supplemented with 10% fetal bovine serum (FBS) for HepG2 and U87MG. Cisplatin-resistant U87MG/DDP cell line and Doxorubicin-resistant HepG2/ADM cell line were established by continuing exposure of the parental cells to gradient increasing concentration of respective drug.

Silencing of AGPS by shRNA

Lentivirus-shRNA mediated AGPS silencing was employed as previously described (Nomura et al., 2010). Lentiviral plasmids (pLKO.1) containing anti-AGPS shRNA (Open Biosystems) were transfected into cancer cells using Fugene HD transfection reagent (Promega). The transfected cells could be selected out over 3 days with 1 µg/mL puromycin. Control shRNA lentiviral particles were also purchased from Open Biosystems. The sequences of shRNA used to silence AGPS were as follows: shRNA #1: AAGGATTTCTTCTCTAGCAGC; shRNA #2: AATTCGCTCAAACATTCCTTC. Silencing of AGPS was confirmed by western blotting and realtime PCR assay.

MTS assay to determine the drug sensitivity of the cells

0.1 ml of cells suspension (HepG2/ADM: 1×10⁵/ml, U87MG/DDP: 3×10⁴/ml) were seeded into each well of the 96-well microplate, and incubated overnight to get the cells to adherence. Then different concentration of Cisplatin or Doxorubicin was added to each well and the cells were cultured for another 72hrs. CellTiter 96 Aqueous One Solution Reagent (Promega) was used according to the manufacturer's instructions. 4hrs later the cell viability was determined by measuring the absorbance at 490nm using a microplate reader.

Flow cytometry to determine intracellular Rhodamine-123, cell cycle distribution, apoptosis rate and fluorescence-based assay to determine activity of caspase-3/8

About 1×10⁶ of the cells were collected and resuspended in 0.1 ml culture medium. The cells were stained with 10µM rhodamine-123 (Sigma-Aldrich) for 1 hr and the intracellular concentration was determined by flow cytometry (BD, FACSCalibur). For cell cycle analysis, PI (BD Biosciences) was used to stain DNA. For apoptosis analysis, 20 µM cisplatin for U87MG/DDP or

2 µM doxorubicin for HepG2/ADM was added for 24h, then Annexin V-FITC/PI dual stain (BD Biosciences) was employed. For caspase-3/8 activation analysis, 2µM cisplatin for U87MG/DDP or 0.1 µM doxorubicin for HepG2/ADM was added for 24h, then activity of caspase-3/8 was measured by Caspase 3 and Caspase 8 activity assay kits (Promega).

Western blotting

The whole proteins were obtained by RAPI lysis buffer (Millipore) extraction and centrifugation at 12,000 g for 10 min. Total protein concentrations of the supernatants were measured by the BCA kit (Sigma Aldrich). 20 µg proteins were separated by 12% SDS-PAGE and transferred onto PVDF membranes. Target proteins were detected by incubation overnight at 4°C with rabbit anti-AGPS (1:800), MDR1 (1:800), MRP1 (1:800), ABCG2 (1:800), Bcl-2 (1:800), survivin (1:800), β-catenin (1:800), Akt (1:800), p-Akt (1:400), β-actin (1:5000) antibody (Santa Cruz). All primary antibodies were diluted according to the instruction manual. Membranes were washed and incubated for 1 hr with peroxidase-labelled anti-rabbit IgG (Santa Cruz, diluted at 1:2000). Finally, membranes were washed three times in TNT and exposed to the ImmobilonTM Western chemiluminescent HRP substrate (Millipore) for 1 min, and then exposed autoradiography film for 2~3 min in the dark.

Real-time PCR analysis

About 3×10⁶ of cells were harvested for RT-PCR analysis. The total mRNA was extracted from the cells by the Dynabeads mRNA Direct Kit (Invitrogen) according to the manual instruction. Total mRNA was then reverse transcribed for 1hr at 42°C in incubation buffer containing 250 µM of each deoxynucleotide triphosphate, 5 µM oligo (dT)20, 25 units of RNase inhibitor, and 20 units of avian myeloblastosis virus reverse transcriptase (Roche Diagnostics). The transcription level of target proteins was detected by semiquantitative real-time PCR using the icycler iQ detection system (Bio-Rad). The PCR condition was as following: decontamination at 50°C for 2min, denaturation at 95°C for 2min, followed by n cycles at 95°C for 20 sec. and at hybridization T°C for 40 sec. The full details were shown in Table 1.

Table 1. Oligonucleotide Sequences Used for Real-Time PCR

Gene	Primer Sequences (5'-3')	Hybridization T°C	Cycles
AGPS	Forward: ACCAGATTCCTGGAGTTC Reverse: GAACCAACAGGTCTCGATA	65	40
MDR1	Forward: AAAAAAGATCAACTCGTACCCTC Reverse: GCACAAAATACACCAACAA	65	37
MRP1	Forward: CTGGGAACATGATTAGGAAGC Reverse: GAGGATTTCCCAGAGCCGAC	65	42
ABCG2	Forward: GGGTTCTTCTTCTCCTGACGACC Reverse: CAGACAACCAAGTTAGAGTGTGGT	65	38
BCL-2	Forward:ACGGGGTGAACCTGGGGGAGGA Reverse:TGTTTTGGGGCAGGCATGTTGACTT	65	40
Survivin	Forward: GCATGGGTGCCCGACGT TG Reverse: GCTCCGGCCAGAGGCCTCAA	60	38
β-catenin	Forward: GGATTCTGGAATCCATTCTGG Reverse: TCTGAGCCCTAGTCATTGCAT	60	38
β-actin	Forward: TGAGCGCGGCTACAGCTT Reverse: TCCTTAATGTCACGCAGGATT	60	40

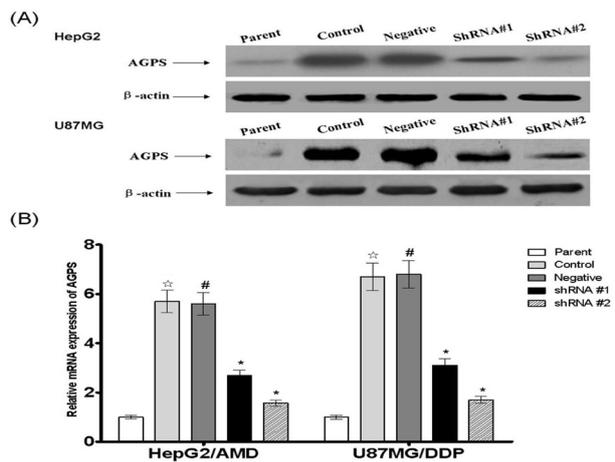


Figure 1. The Expression of AGPS in Human Glioma and Hepatic Carcinoma Cell Lines. A) Western blotting results showed that there was an increased protein expression of AGPS in human hepatic carcinoma HepG2/ADM and glioma U87MG/DDP cell lines compared with HepG2 and U87MG cell lines, meanwhile, there was a decreased protein expression of AGPS in human hepatic carcinoma HepG2/ADM and glioma U87MG/DDP AGPS silencing groups compared with HepG2/ADM and U87MG/DDP cell lines. β -actin was as an internal control. B) Realtime PCR results showed that there was an increased mRNA expression of AGPS in human hepatic carcinoma HepG2/ADM and glioma U87MG/DDP cell lines compared with HepG2 and U87MG cell lines, meanwhile, there was a decreased mRNA expression of AGPS in human hepatic carcinoma HepG2/ADM and glioma U87MG/DDP AGPS silencing groups compared with HepG2/ADM and U87MG/DDP cell lines. β -actin was as an internal control. Bars mean \pm SD. n=5. $\star p < 0.05$, compared to the parent group; $\# p > 0.05$, compared to the control group; $* p < 0.05$, compared to the control group

Extraction and quantization of LPA and LPAe

Cancer cells were starved in serum-free media for 4 h to minimize the contribution of serum-derived metabolites. About 1×10^6 cells were washed twice by PBS and harvested by scraping and centrifugation. Cell pellets were flash frozen at -80°C . Nonpolar lipid metabolites were extracted and analyzed by LC-MS/MS as described previously. Briefly, cell lysis was extracted by 4 mL of mixture of 2 (chloroform): 1 (methanol): 1 PBS with C12:0 dodecylglycerol (10 nmol) and pentadecanoic acid (10 nmol) as internal standards. Organic and aqueous layers were separated by centrifugation. The aqueous layer was acidified by adding 0.1% formic acid and reextracted by chloroform. The organic layer was dried by N_2 and dissolved in 120 μL chloroform and a 10 μL aliquot was analyzed by Agilent 6430 LC-MS/MS.

Analysis of PGE2 by enzyme immunoassay

0.1 ml of cancer cells (HepG2/ADM: $1 \times 10^6/\text{ml}$, U87MG/DDP: $4 \times 10^5/\text{ml}$) were plated in 96-well plates and cultured overnight. The next day, cells were treated with 0.1 ml arachidonic acid (final concentration 15 μM) for 1 h before collecting the culture medium. PGE2 levels in the medium were determined by PGE2 enzyme immunoassay kits (Cayman Chemical).

Measurement of PLD activity

Cancer cells were seeded into 24-well plate and

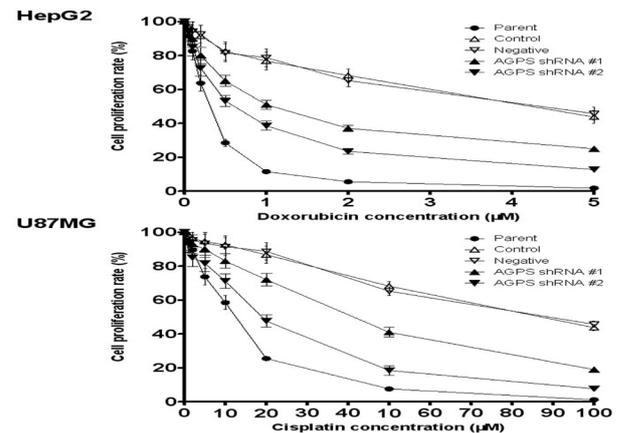


Figure 2. The Effect of AGPS Silencing on Drug Sensitivity in Human Glioma and Hepatic Carcinoma Cell Lines. The MTS results showed that there was an increased drug sensitivity in human hepatic carcinoma HepG2/ADM and glioma U87MG/DDP AGPS silencing groups compared with HepG2/ADM and U87MG/DDP cell lines. Bars mean \pm SD. n=10

allowed to reach to near 90% fusion. Cells were washed three times with PBS. Then 100 μL of an ice-cold Tris 50 mM (pH 8.0) was added to each well and the cells were broken by three freeze and thaw cycles. Samples were collected, mixed with an equal volume of the Amplex Red reaction buffer (Amplex Red Phospholipase D assay kit, Life Technologies), and the PLD activity was estimated after 1 h incubation at 37°C with a Bio-Rad fluorometer. A standard curve was performed with purified PLD from *Streptomyces chromofuscus* (Sigma-Aldrich). Data are given as the mean of six determinations performed on three different cell preparations.

Stimulation of EP receptors by PEG2 and respective agonists

Cancer cells were seeded into 96-well plate as MTS assay mentioned above. On the second day, PEG2 (5 μM) or respective agonists (19(R)-hydroxy PGE1, 1 μM , EP1 and EP3 receptor agonist; CAY10399, 1 μM , EP2 receptor agonist; L-902,688, 100 nM, EP4 receptor agonist; Cayman Chemical) were added to each well 1 h before adding different concentration of drugs to stimulate EP receptors. 72 hrs later, CellTiter 96 Aqueous One Solution Reagent (Promega) was employed to determine the cell viability.

Statistical analyses

Data were expressed as mean \pm SD., statistical analysis was carried out using One Way ANOVA and $p < 0.05$ indicates statistical significance.

Results

The HepG2/ADM and U87MG/DDP cells were resistant to chemotherapy drug and over-expressed AGPS

Clones of drug resistant cancer cells were established by continuing exposure to chemotherapy agents. Significant over-expression of AGPS was found in these drug resistant cancer cells compared to the drug sensitive HepG2 and U87MG cells (parent group).

The drug resistant clones were transfected with

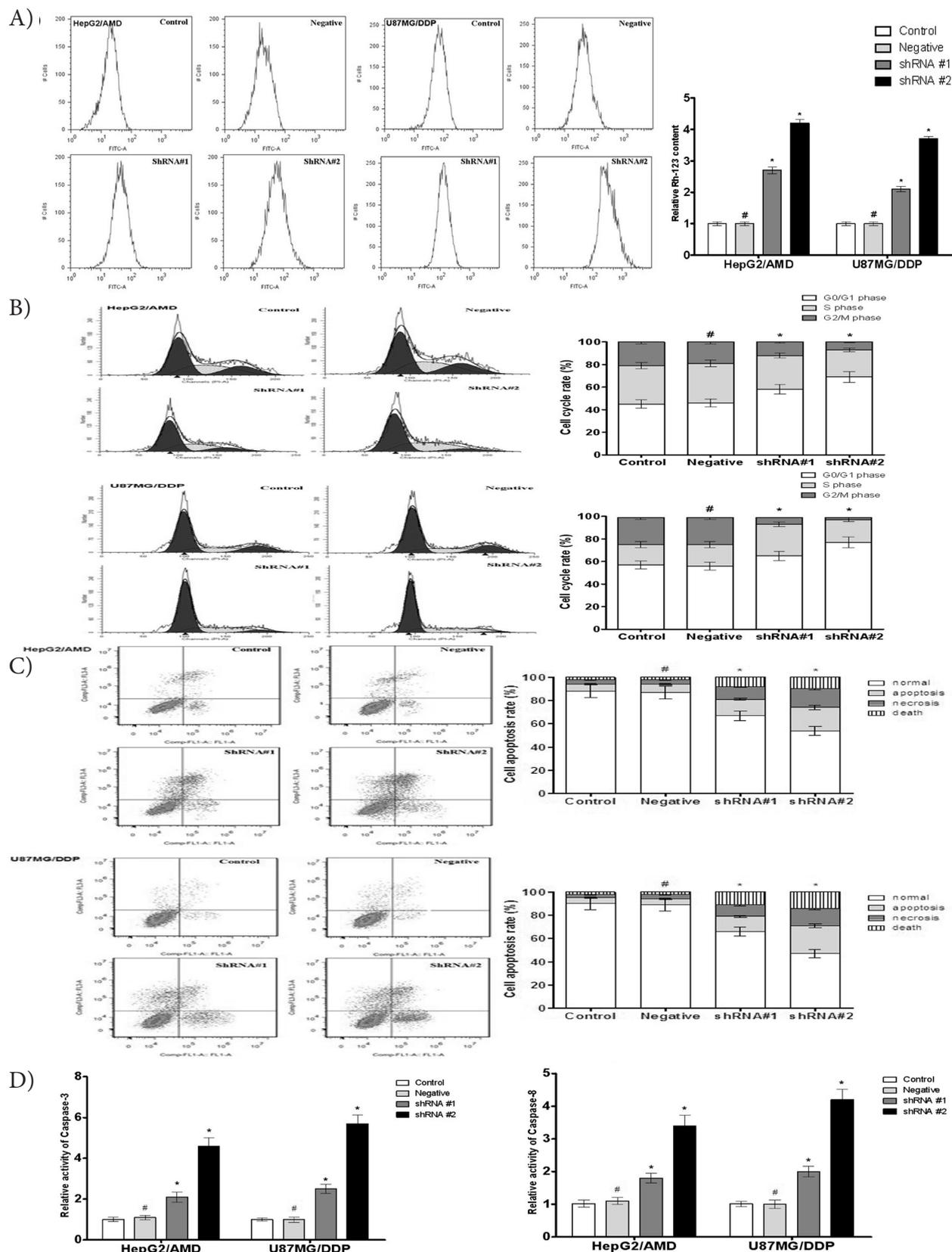


Figure 3. The Effect of AGPS Silencing on Intracellular Accumulation of Rhodamine-123, Cell Cycle Arrest and Increase of Apoptosis Rate and Caspase-3/8 Activities in Human Glioma and Hepatic Carcinoma Cell Lines. A) The FCM results showed that there was an increased intracellular accumulation of Rhodamine-123 in human hepatic carcinoma HepG2/ADM and glioma U87MG/DDP AGPS silencing groups compared with HepG2/ADM and U87MG/DDP cell lines. Bars mean±SD. n=3. #*p*>0.05, compared to the control group; **p*<0.05, compared to the control group. B) The FCM results showed that there was an increased G0/G1 phase in human hepatic carcinoma HepG2/ADM and glioma U87MG/DDP AGPS silencing groups compared with HepG2/ADM and U87MG/DDP cell lines. Bars mean±SD. n=3. #*p*>0.05, compared to the control group; **p*<0.05, compared to the control group. C) The FCM results showed that there was an increased apoptosis rate in human hepatic carcinoma HepG2/ADM and glioma U87MG/DDP AGPS silencing groups compared with HepG2/ADM and U87MG/DDP cell lines. Bars mean±SD. n=3. #*p*>0.05, compared to the control group; **p*<0.05, compared to the control group. D) The fluorescein results showed that there was an increased caspase-3/8 activity in human hepatic carcinoma HepG2/ADM and glioma U87MG/DDP AGPS silencing groups compared with HepG2/ADM and U87MG/DDP cell lines. Bars mean±SD. n=5. #*p*>0.05, compared to the control group; **p*<0.05, compared to the control group.

different sequences of AGPS-shRNA (shRNA #1 and shRNA #2) in order to produce two groups of cells with different AGPS silencing level. Both HepG2/ADM and U87MG/DDP cells transfected with AGPS-shRNA resulted in down-regulation of AGPS mRNA expression in shRNA #1 and shRNA #2 groups compared to the HepG2/ADM and U87MG/DDP cells (control group) ($p < 0.05$). The blank vector HepG2/ADM and U87MG/DDP cells (negative group) had no effect on AGPS mRNA expression when compared to the HepG2/ADM and U87MG/DDP control groups ($p > 0.05$), meanwhile, the AGPS protein expression was showed the similar tendency (Figure 1).

AGPS silencing restored the drug sensitivity

After knockdown of the AGPS, we examined the

drug sensitivity of each cell clone by MTS assay. As shown in Figure 2, we could see the cell proliferation rate of the AGPS silencing cells were lower than that of the control group. The negatively transfected cells were nearly unaffected. The IC_{50} of shRNA#1 and shRNA#2 groups were $0.97\mu M$ and $0.53\mu M$ in HepG2/ADM cell lines for doxorubicin respectively, which was significant sensitive compared with the control group (IC_{50} was $4.4\mu M$) ($p < 0.05$). The reversal factor was 4.5 and 8.3, respectively. The IC_{50} of shRNA#1 and shRNA#2 groups were $31.3\mu M$ and $18.7\mu M$ in U87MG/DDP cell lines for cisplatin respectively, which was significant sensitive compared with the control group (IC_{50} was $94.2\mu M$) ($p < 0.05$). The reversal factor was 3.0 and 5.0, respectively. These results indicated that AGPS play a positive role in

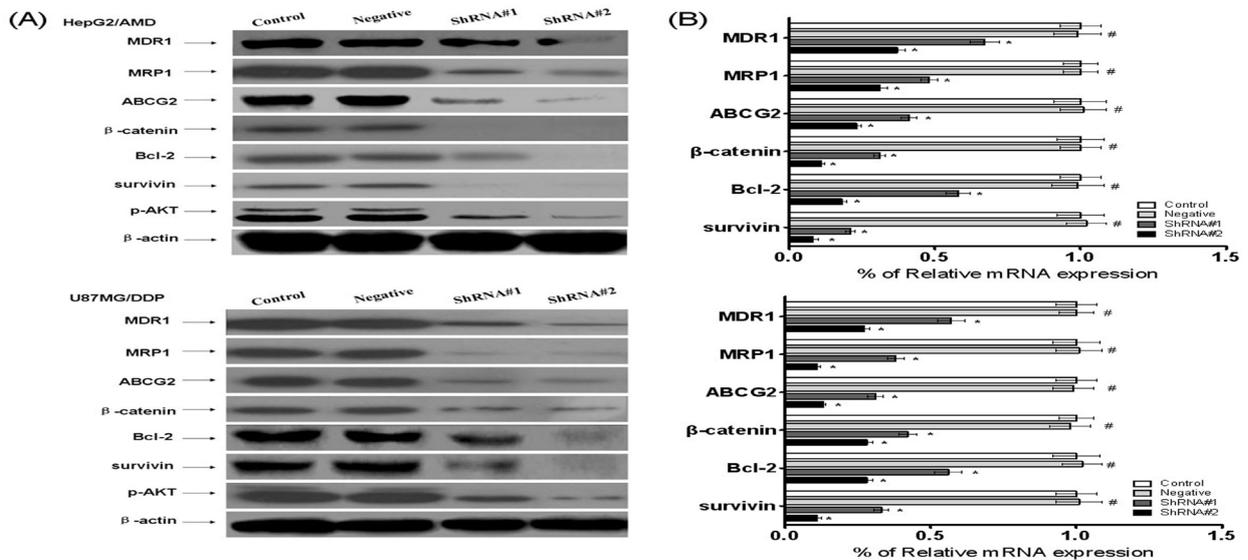


Figure 4. The effect of AGPS Silencing on Expression of Drug Resistance Related Gene in Human Glioma and hepatic Carcinoma Cell lines. A) Western blotting results showed that there was a decreased protein expression of MDR1, MRP1 and ABCG2, β -catenin, Bcl-2, survivin and p-AKT in human hepatic carcinoma HepG2/ADM and glioma U87MG/DDP AGPS silencing groups compared with HepG2/ADM and U87MG/DDP cell lines. β -actin was as an internal control. B) Realtime results showed that there was a decreased mRNA expression of MDR1, MRP1 and ABCG2, β -catenin, Bcl-2, survivin and p-AKT in human hepatic carcinoma HepG2/ADM and glioma U87MG/DDP AGPS silencing groups compared with HepG2/ADM and U87MG/DDP cell lines. β -actin was as an internal control. Bars mean \pm SD. n=5. * $p < 0.05$, compared to the control group

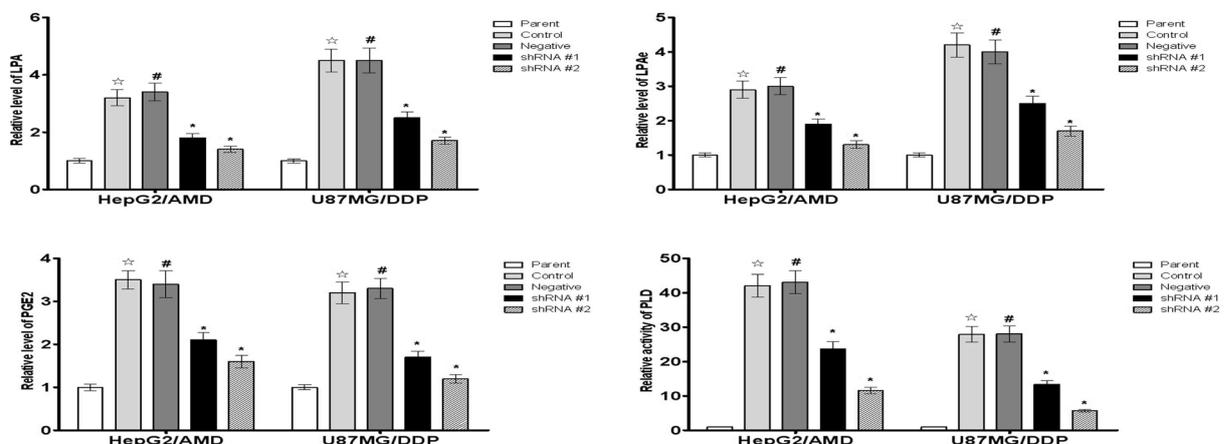


Figure 5. The Effect of AGPS Silencing on Expression of LPA, LPAe, PGE2 and Activity of PLD in Human Glioma and Hepatic Carcinoma Cell Lines. The results showed that there was an increased expression of LPA, LPAe, PGE2 and activity of PLD in human hepatic carcinoma HepG2/ADM and glioma U87MG/DDP cell lines compared with HepG2 and U87MG cell lines, meanwhile, there was a decreased expression of LPA, LPAe, PGE2 and activity of PLD in human hepatic carcinoma HepG2/ADM and glioma U87MG/DDP AGPS silencing groups compared with HepG2/ADM and U87MG/DDP cell lines. Bars mean \pm SD. n=10. $\star p < 0.05$, compared to the parent group; # $p > 0.05$, compared to the control group; * $p < 0.05$, compared to the control group

the chemotherapy resistance of the cancer cells.

Restore of drug sensitivity was accompanied with an intracellular accumulation of Rhodamine-123, cell cycle arrest and increase of apoptosis rate and caspase-3/8 activities

The FCM analysis showed that AGPS silenced cells had a greater intracellular fluorescent activity, which meant that more Rh-123 was retained in the cancer cells. Meanwhile, cell cycle analysis showed that in the AGPS attenuated cancer cells, more cells were distributed in G0/G1 phase, and the percentages of S phase and G2/M phase cells were reduced. After treatment with cisplatin or doxorubicin, the apoptosis rates of AGPS silenced cells were increased significantly when compared to the control cells. Further, the activity of caspase-3/8 was increased significantly in AGPS silenced groups compared to the control group as shown in Figure 3.

AGPS modulated the expression of MDR1, MRP1, ABCG2, β-catenin, Bcl-2, survivin and p-AKT

We investigated whether expression of transporter proteins from ABC family, such as MDR1, MRP1 and ABCG2, which have the potency to transport both Rh-123 and chemotherapy agents, were modulated. Meanwhile, β-catenin could stimulate cell growth and proliferation via its interaction with TCF-4 in the nucleus (Castellone et al., 2005). Bcl-2 and survivin have been implicated in anti-apoptosis, and is also thought to be involved in resistance to conventional cancer treatment (Mita et al., 2008; Aust et al., 2013). As shown in Figure 4, the expression of MDR1, MRP1 and ABCG2, β-catenin, Bcl-2 and survivin was down regulated in AGPS silencing groups compared with control group. The RT-PCR results were consistent

to the results from western blotting (Figure 4).

AGPS modulated drug resistance through LPA, LPAe and PGE2 mediated lipid signaling pathways

LPA, LPAe and PGE2 are several lipids that have been found to be involved in AGPS mediated cancer pathogenicity. Here we demonstrated that in all the three drug resistant cancer cells, LPA, LPAe and PGE2 levels increased when compared to the non-resistant cancer cells. Silencing of AGPS resulted in reduction of LPA, LPAe and PGE2. Since both LPA, LPAe and PLD binds to LPA receptors to drive multiple aspect of cancer (Blackburn et al., 2012; Madan et al., 2013), we tested the downstream signaling activities of LPA receptor. We found that both level of LPA, LPAe and PLD in control group of HepG2/ADM and U87MG/DDP was increased compared to the HepG2 and U87MG cells (parent group) ($p < 0.05$), however, there was not significant modification between control and negative group of HepG2/ADM and U87MG/DDP ($p > 0.05$). Further, both level of LPA, LPAe and PLD in shRNA#1 and shRNA#2 group of HepG2/ADM and U87MG/DDP was decreased to compared to the control group of HepG2/ADM and U87MG/DDP ($p > 0.05$) (Figure 5).

Furthermore, the phosphorylation of AKT was down-regulated as well, which meant the activity of PI3K/AKT pathway was attenuated. PI3K/AKT is a key signaling pathway downstream LPA receptor. To confirm the role of PEG2 in AGPS mediated drug resistance, we supplemented the AGPS silenced cells with PEG2 and EP receptor agonists and found that the drug sensitivity established by AGPS shRNA#1 and shRNA#2 was reversed (Figure 6).

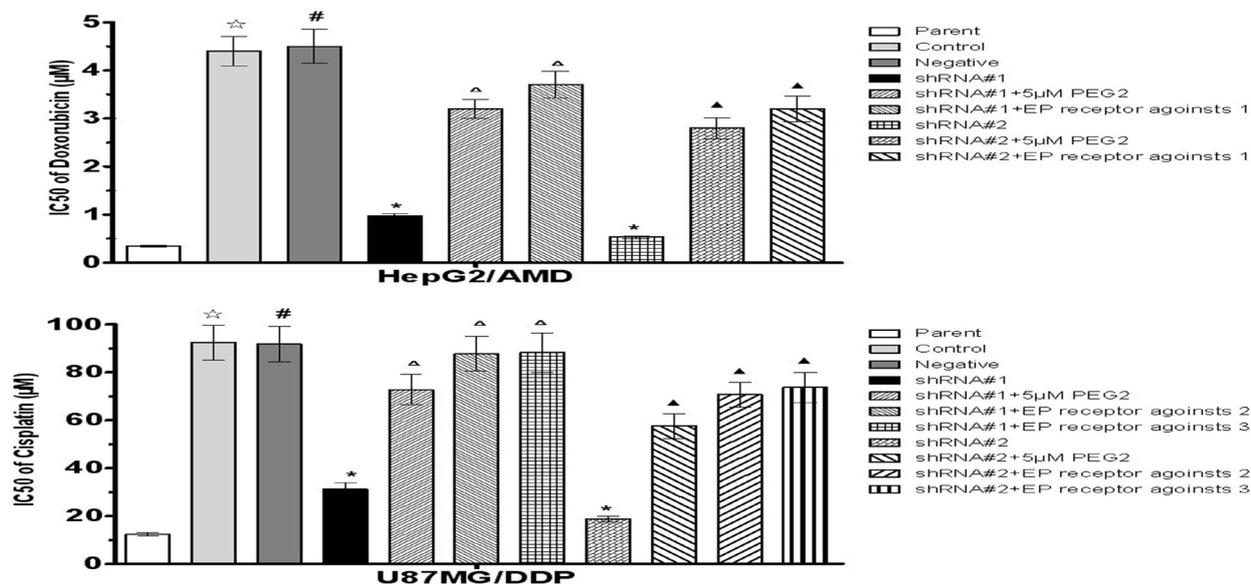


Figure 6. The Effect of AGPS Silencing on Drug Sensitivity Mediated by PEG2 and EP receptor in Human Glioma and Hepatic Carcinoma Cell Lines. The results showed that there was a decreased drug sensitivity in human hepatic carcinoma HepG2/ADM and glioma U87MG/DDP cell lines compared with HepG2 and U87MG cell lines, meanwhile, there was an increased drug sensitivity in human hepatic carcinoma HepG2/ADM and glioma U87MG/DDP AGPS silencing groups compared with HepG2/ADM and U87MG/DDP cell lines. Further, there was a decreased drug sensitivity in human hepatic carcinoma HepG2/ADM and glioma U87MG/DDP silencing cell lines by PEG2 and EP receptor agonists treatment. Bars mean±SD. n=10. ☆ $p < 0.05$, compared to the parent group; # $p > 0.05$, compared to the control group; * $p < 0.05$, compared to the control group; * $p < 0.05$, compared to the AGPS-shRNA#1 group; ▲ $p < 0.05$, compared to the AGPS-shRNA#2 group. 1. 19(R)-hydroxy PGE1, 1µM, EP1 and EP3 receptor agonist; 2. CAY10399, 1µM, EP2 receptor agonist; 3. L-902,688, 100nM, EP4 receptor agonist

Discussion

In order to investigate the role of AGPS in chemotherapy resistance, we used lentivirus to knock-down AGPS in several drug-resistant cancer cell lines in this study. In consistent with previous study that indicate AGPS play a role in the cancer pathogenesis, our MTS assay showed that drug sensitivity of the resistant cancer cells could be restored after AGPS silencing. Further investigations showed that after AGPS silencing, intracellular accumulation of Rh-123 increased, more cells were distributed to G0/G1 phase, and apoptosis rate increased after chemotherapy agent treatment. Therefore, we following these clues to explore the possible mechanism of AGPS mediated resistance.

One major mechanism that involved in drugs resistance is increased expression of multi-drug resistant proteins which efflux drug out of cancer cells, for example MDR1, MRP1 and so on (Louisa et al., 2014). The increase of intracellular concentration of Rh-123 means that drug efflux by the cancer cells is inhibited after AGPS silencing. Western blotting and RT-PCR showed that the expression of ABC family transporters MDR1, MRP1 and ABCG2 were down-regulated by AGPS silencing. Other mechanisms involve cell cycle promotion and anti-apoptosis effects. Cell cycle analysis showed that AGPS silencing resulted in more cancer cells distributing to G0/G1 phase. We found that a key cell cycle modulator protein, β -catenin, was regulated by AGPS. Furthermore, western blotting and RT-PCR showed that the expression of Bcl-2 and survivin, which are involved in cell apoptosis, were modulated by AGPS as well. The mechanisms that AGPS mediated the expression of these proteins could be through the PEG2 pathway as described below.

AGPS is an enzyme that catalyzes the synthesis of lipids. It has been reported that AGPS affects the level of LPA and LPAe, which could promote the cancer. Here we showed that after AGPS silencing, level of LPA and LPAe were all down-regulated in HepG2/ADM and U87MG/DDP. To confirm that LPA and LPAe play a role in the drug resistance of our cancer cells, we determined the downstream pathway. LPA and LPAe could activate the LPA receptor and further activate the PI3K/AKT pathway. Our western blotting assay showed that phosphorylation of AKT was downregulated with AGPS silencing. Furthermore, the activity of PLD, a phospholipase that transduces the activity of LPA receptor, was down-regulated as well. It has been reported that PLD plays a role in the drug resistance of cancer. LPA receptor stimulates PLD activation and then accumulates intracellular PA (phosphatidate). Elevation of PA could activate several pathways such as SK-1/S1P, Raf/ERK, mTOR and PKC- ζ to promote cancer cell migration and survival.

Another lipid-related substrate that is affected by AGPS is PEG2. Our EIA assay showed that intracellular PEG2 was down-regulated after AGPS silencing. Moreover, when supplementing the AGPS silenced cancer cells with PEG2, the drug resistance reemerged, which further valid the role of PEG2 in AGPS mediated drug resistance. In cancer cells, PEG2 could bind to respective receptors to

activate the downstream signaling pathway. There are 4 subtype of receptors, named EP1~4 (Greenhough et al., 2009). It has been reported that in human hepatocellular carcinoma cells EP1 receptor transactivates EGFR/MET receptor tyrosine kinases and enhances invasiveness (Han et al., 2006; Gillibert-Duplantier et al., 2007). U87MG has been reported to express both EP2 and EP4 (Kambe et al., 2008). To confirm the role of EP receptors in respective cancer cells, we used selective agonist to stimulate the respective receptor. Selective EP1 and EP3 agonist 19(R)-hydroxy PGE1 could restore drug resistance of HepG2/ADM, while selective EP2 agonist CAY10399 and selective EP4 agonist L-902,688 had similar effect on U87MG. PGE2 is the key in COX-2 regulating the express of multidrug resistance proteins in cancer cells (Woodward et al., 1993; Tani et al., 2001; Takaoka et al., 2008; Henderson et al., 2011; Hu et al., 2013; Kashiwagi et al., 2013). COX-2/PGE2 pathway could enhance the activation of the APC/ β -catenin pathway to promote cancer cell cycle. Furthermore, PGE2 could regulate the expression of Bcl-2 and survivin to prevent cancer cells from apoptosis (Krysan et al., 2004). These literature reports are consistent with our observation mentioned above. PI3K/AKT is key pathway downstream EP receptors as well. Hence, the downregulation of Akt phosphorylation mentioned above might be the joint action of modulation to LPA receptor and EP receptor.

In summary, we found that AGPS overexpression could be a causation of chemotherapy agent resistance of cancer cells. AGPS silencing could lead to drug uptake and cell cycle arrest and apoptotic cell death. Our study suggested that lipid signaling pathways was critical for AGPS mediated drug resistance.

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