Effects of proton pump inhibitors on reversing multidrug resistance via downregulating V-ATPases/PI3K/Akt/mTOR/HIF-1α signaling pathway through TSC1/2 complex and Rheb in human gastric adenocarcinoma cells in vitro and in vivo

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Background: Our study aimed to explore the effects of PPIs on reversing multidrug resistance (MDR) to chemotherapy in gastric cancer by inhibiting the expression of V-ATPases and the PI3K/Akt/mTOR/HIF-1α signaling pathway.

Methods: The gastric cancer cell lines SGC7901 and the multidrug resistance cell lines SGC7901/MDR were pretreated by the pantoprazole or the esomeprazole, respectively. Real-time PCR was used to determine mRNA levels, and western blotting and immunofluorescent staining analyses were employed to determine the protein expressions and intracellular distributions of the V-ATPases, PI3K, Akt, mTOR, HIF-1α, P-gp and MRP1 before and after PPIs pretreatment. SGC7901/MDR cells were planted on the athymic nude mice. Then the effects of PPZ pretreatment and/or ADR were compared by determining the tumor size, tumor weight and nude mouse weight.

Results: PPIs pretreatment could inhibit mRNA levels of V-ATPases, MDR1 and MRP1, PI3K, Akt, mTOR and HIF-1α. PPIs inhibited V-ATPases and down-regulated the expressions of P-gp and MRP1. And further to block the expression of mTOR by Rapamycin could obviously inhibit the expressions of HIF-1α, P-gp and MRP1 in a dose-dependent manner. Therefore, PPIs inhibited the expressions of V-ATPases and then reversed MDR of the chemotherapy in gastric cancer by inhibiting P-gp and MRP1, and it could be speculated that the mechanism might be closely related to down-regulating the PI3K/Akt/mTOR/HIF-1α signaling pathway. Meanwhile, PPIs also could inhibit the expressions of TSC1/TSC2 complex and Rheb which might be involved into regulating the signaling pathway intermediately. The weight growth rate of the mice bearing tumor in the treatment group was lower than that of the nude mice in the normal group, while the weight growth rate of the mice in control group was significantly lower than that of the normal group and the treatment group, presenting a downward trend.

Conclusion: Therefore, PPIs inhibited the expressions of V-ATPases and then reversed MDR of the chemotherapy in gastric cancer by inhibiting P-gp and MRP1, and it could be speculated that the mechanism might be closely related to down-regulating the PI3K/Akt/mTOR/HIF-1α signaling pathway, and also to inhibiting the expressions of TSC1/TSC2 complex and Rheb which might be involved into regulating the signaling pathway intermediately.

Keywords: gastric adenocarcinoma, V-ATPases, tumor acidity, hypoxia, proton pump inhibitors, multidrug resistance, signaling pathway
Introduction

Gastric cancer is one of the most common digestive malignancies with lower detection rate during the early stage. Most of patients are diagnosed after cancer has advanced rather than at early stage. Curative effects are often difficult to achieve through only surgery. Chemotherapy plays an important role in comprehensive therapy of cancer. However, many cancers failed to respond to chemotherapy by acquiring multidrug resistance (MDR). Although MDR had several causes, one major form of resistance to chemotherapy was closely correlated with the presence of at least three molecular “pumps” that actively transported drugs out of the cancer cells. The most prevalent of those MDR transporters were P-glycoprotein (P-gp) and the multidrug resistance protein 1 (MRP1), two members of the ATP Binding Cassette superfamily.

Tumor microenvironment, characterized by reversed pH gradient with an acidic extracellular pH (pHe) and an alkaline intracellular pH (pHi) compared to the normal cells, closely affected tumor sensitivity to chemotherapy. Evidence was accumulating that hypoxia and acidity were deeply involved in tumor microenvironment, resulting in cancer progression, invasiveness, metastasis, and tumor chemosensitivity. The hypoxic and acidic tumor microenvironment might conversely induce selection of tumor cells being able to survive in such an unfavorable condition. Furthermore, a common feature of MDR cells is net decrease in the intracellular accumulation of drugs. The acid pHe will effectively impair the entry of weakly basic antitumor drugs into cancer cells since it could neutralize or sequester drugs into the acidic intracellular vesicles.

Vacuolar H⁺-ATPases (V-APTases), as a specific proton type of the cells, mainly function in the control of pH and pHi. V-APTases are involved in maintaining a relatively neutral pHi, luminal pH, and an acidic pHe. Although cancer cells with low metastatic potential preferentially use Na⁺/H⁺ exchangers and HCO₃⁻-based H⁺-transporting mechanisms, highly metastatic cells preferentially use plasma membrane V-APTases. The expressions of V-APTases obviously are highly expressed in chemoresistant cancer cells and can be induced by chemotherapeutics. Recent results showed that a molecular inhibition of V-APTases by small interfering RNA (siRNA) led to tumor cytotoxicity and marked inhibition of human tumor growth in xenograft model.

PPIs, mainly treating acid-related diseases, could inhibit the expressions of V-APTases and then reverse the transmembrane pH gradient to sensitize the SGC7901 cells to the antitumor drugs, which was reflected in the downregulated expressions of MRP1 and P-gp. PPIs treatment reversed the pH gradient by blocking the V-APTases-mediated H⁺ efflux, in turn allowing anticancer drugs to penetrate and function within tumor cells. However, so far, the concrete mechanism for PPIs on reversing the MDR of chemotherapy in gastric cancer was not yet known. Hence, the molecular characterization of pathways regulating chemosensitivity is a central prerequisite to improve chemotherapy.

As originally reported by Otto Warburg, most cancers are characterized by aerobic glycolysis-wasteful glycolytic conversion of glucose to lactic acid, even when sufficient oxygen is available to support efficient mitochondrial respiration. In fact, an important determinant of tumor acidity is the anaerobic metabolism that allows the selection of cells able to survive in a hypoxic-anoxic environment through the upregulation of hypoxia-inducible factor 1α (HIF-1α) and the adaptation of a glycolytic phenotype with generation of lactate. HIF-1α, upregulated by hypoxia, plays an important role in determining chemosensitivity focused on responsible molecular pathway. Blocking the expression of HIF-1α by siRNA in gastric cancer cells could effectively reverse MDR phenotype induced by hypoxia. Although oxygen tension plays a determinant role in the process of HIF-1α activation, the amplitude of this response is modulated by growth factor-dependent signaling pathways, including the Ras-Erk and phosphoinositide 3-kinase (PI 3-kinase)/AKT cascades. Furthermore, HIF-1α located the downstream of mTOR (mammalian target of rapamycin) complex 1 (mTORC1), which was regulated by the PI3K/Akt signaling pathway. Amplified signaling through phosphoinositide 3-kinase, and its downstream target, mTORC1, enhances HIF-1-dependent gene expression in certain cancer types. Pretreatment with the mTOR inhibitor, Rapamycin, inhibited both the accumulation of HIF-1α and HIF-1-dependent transcription induced by hypoxia.

The primary function of the TSC1-TSC2 complex is considered as a critical negative regulator of mTORC1 activation. As with the TSC-mTOR connection, the GTP-bound Rheb regulated by the TSC1-TSC2 complex to control mTORC1 activation was identified and characterized through a combination of genetics and biochemistry. Purified Rheb loaded with GTP, but not GDP, stimulates mTORC1’s in vitro kinase activity in a dose-dependent manner, suggesting that the regulation is rather direct to a certain extent. Nowadays, molecular characterization of signaling pathways on regulating chemosensitivity is a central prerequisite to improve cancer therapy. The PI3K/Akt/mTOR/HIF-1α signaling pathway has been linked to chemosensitivity, while the underlying molecular mechanisms remain largely obscure.
Our study therefore aimed to investigate whether PPIs could improve the chemosensitivity on the gastric cancer cells via downregulating the expressions of V-ATPases, and then PI3K/Akt/mTOR/HIF-1α signaling pathway by inhibiting the expressions of P-gp and MRP1. Among them, TSC1/2 complex and Rheb might be involved into regulating the signaling pathway immediately.

**Materials and methods**

**Reagents and drugs**

V-ATPases 6V, A mouse polyclonal antibody was purchased from Abnova company (Taipei City, Taiwan, P.R. China). PI3 kinase p110α antibody, Akt antibody, mTOR antibody, Phospho/TSC2(s933) antibody, Phospho/TSC2(Thr1462) antibody, phosphor(Ser/Thr) Akt substrate antibody were purchased from Cell Signaling Technology, Inc. (Danvers, MA, USA). Hamartin (TSC1), Tuberin (TSC2), HIF-1α, P-gp, MRP1, RHEB (Ras homolog enriched in brain) were purchased from Abcam PLC (Cambridge, MA, USA). Pan-toprazole (PPZ) Sodium salts (Altana Pharma AG D-78467, Wesel, Germany) and Esomeprazole Sodium for Injection (AstraZeneca PLC, Cambridge, UK) were resuspended in the RPMI-1640 (pH 6.65, in order to activate PPIs) at 1 mg/mL immediately before use. Rapamycin was purchased from LC Laboratories (Woburn, MA, USA).

**Cell lines and cell culture**

Human gastric adenocarcinoma cell line, SGC7901, was kindly gifted by the Department of Oncology, the Affiliated Drum Tower Hospital of Nanjing University, Medical School. The multidrug-resistant SGC7901 cells (SGC7901/MDR) were induced by our lab with the Adriamycin and Cisplatin step by step. The average values of IC50 to the Adriamycin in the SGC7901 and the SGC7901/MDR cells lines, detected by Cell Counting Kit-8 (CCK-8, MedChem Express, NJ, USA) assay, were 0.654 ± 0.103 and 3.203 ± 0.302 µg/mL, respectively (P < 0.05) (Figure 1). Cells lines were cultured in RPMI-1640 medium (Hyclone, Thermo Fisher Scientific, Waltham, MA, USA) supplemented with 10% fetal bovine serum (Hangzhou Sijiqing Biological Engineering Materials Co., Ltd., Zhejiang, People’s Republic of China) and antibiotics (100 U/mL penicillin and 100 µg/mL streptomycin) in a humidified air with 5% CO2 atmosphere at 37°C (Thermo Direct Heat CO2, Thermo Direct Inc., Garner, NC, USA). Multidrug resistant (Adriamycin and Cisplatin) SGC7901 cell strains were cultured in the Medium containing Adriamycin (800 ng/mL) maintaining their drug-resistant phenotype. The in vivo experiments strictly followed the ethical principles and national guidelines for scientific experiments on animals. Our in vitro experiments were approved by our institutional review board and ethics committee of the Affiliated Drum Tower Hospital of Nanjing University, Medical School.

**PPIs pretreatment**

The SGC7901 and SGC7901/MDR cells were treated with Esomeprazole in the medium at pH 6.65 (RPMI-1640) for 24 hours with the concentration of 0, 10, 20, 50, 80, 100 µg/mL, respectively. SGC7901/MDR cells also were treated with PPZ for 24 hours with the same concentration gradient under the same condition.

**V-ATPases siRNA interference assay**

The V-ATPases siRNA Assay kit (catalog 4390824, Ambion, Thermo Fisher Scientific) with Life Technologies (Thermo Fisher Scientific) was used according to the instructions of the manufacturer. For transfection, cells in exponential growth phase were plated in six-well plates containing antibiotic-free medium at 30% confluence and incubated overnight, then transfected with V-ATPases siRNA using lipofectamine 2000 (Invitrogen, Thermo Fisher Scientific), according to the manufacturer’s protocol. The final concentration of siRNA was 50 nM. After 24 hours transfection, mediums were replaced with RPMI-1640 supplemented with 10% fetal bovine serum 24 hours. Total proteins were extracted from cells for Western blotting.

**Rapamycin inhibition assay**

The SGC7901/MDR cells were seeded into six-well plates in 2 mL of standard growth medium. After an overnight culture,
the cells were washed and then transferred into low-serum (2%) medium. After starvation for 12 hours, the cells were pretreated with 20, 40, 80 µg/mL rapamycin for 24 hours, respectively.

RNA extraction, reverse transcription, and quantitative real-time PCR

Total RNA was extracted from cultured cells using TRIzol reagent (Sigma-Aldrich Co., St Louis, MO, USA) and reverse transcription was carried out with 1 µg RNA in a total 20 µL reaction volume using PrimeScript™ RT Master Mix (Takara Bio Inc., Kusatsu, Japan) according to the manufacturer’s instructions as described previously. cDNA was used as a template in Mix (Bio-Rad Laboratories Inc., Hercules, CA, USA). Quantitative real-time PCR experiments were done with the 7500 Real-time PCR system (Applied Biosystems, Thermo Fisher Scientific) by using SYBR Premix Ex Taq reagents (Takara Bio Inc.). Primers were designed and validated by Invitrogen Biotechnology Co., Ltd (Thermo Fisher Scientific). The sequences of primers used in real-time PCR are listed in Table 1. All data were normalized to the human β-actin. All assays were done in triplicate.

Western blotting analysis

The SGC7901 and SGC7901/MDR Cells were gathered after various kinds of PPI pretreatments. And total protein was extracted on ice in lysate (containing 0.01% PMSF, 150 mM NaCl, 50 mM Tris (pH 8), 0.1% SDS, 0.2% EDTA, 1% Triton X-100, 1% sodium deoxycholate) supplemented with protease inhibitors (aprotinin, leupeptin, phenylmethylsulfonyl fluoride, sodium orthovanadate; Hoffman-La Roche Ltd., Basel, Switzerland) and phosphatase inhibitors (Cell Signaling Technology, Inc.). Protein concentrations were measured by the BCA Protein Assay Kit, following the manufacturer’s instructions. Equal quantities were separated by SDS-PAGE, transferred to nitrocellulose membranes using a semidyrid transfer system (Bio-Rad Laboratories Inc.). And probed with antibodies against V-ATPase 6V, A, PI3 kinase p110α, Akt, mTOR, TSC1, TSC2, HIF-1α, P-gp, MRPI, Rheb, Phospho/TSC2(Ser939), Phospho/TSC2(Thr1462), Phospho(Ser/Thr) Akt substrate, a monoclonal mouse antibody to β-actin (Cell Signaling Technology, Inc.) and a monoclonal rabbit antibody to Tubulin (Bioworld Technology, Inc., St Louis Park, MN, USA) as the controls for protein loading. Anti-mouse or anti-rabbit IgG, conjugated with horseradish peroxidase (Cell Signaling Technology, Inc.) was used for secondary detection. Antibody staining was visualized by enhanced chemiluminescence (Santa Cruz Biotechnology Inc., Dallas, TX, USA). The images of Western blot products were collected and analyzed by Quantity One V4.31 (Bio-Rad Laboratories Inc.).

Immunofluorescent staining analysis

Dispersed single cells (2 × 10⁶ cells per well) were grown on 22 × 22 × 1 mm³ glass cover slips (pretreated with 0.3% gelatin) in six-well culture plates. After 36–48 hours incubation and 24 hours Esomeprazole pretreatment, cells were fixed in ice-cold acetone for 10 minutes at 4°C. The cells were blocked with 10% normal goat serum (Boster Biological Technology, Pleasanton, CA, USA) for 30 minutes and probed with V-ATPase 6V, A antibodies, PI3 kinase p110α antibodies, Akt antibodies, mTOR antibodies, HIF-1α antibodies, and P-gp antibodies (1:100) at 4°C overnight, respectively. Alexa Fluor Dye Conjugated secondary antibodies (1:75, Alexa Fluor 488 goat anti-mouse IgG (H + L) highly cross-absorbed), 2 mg/mL, Invitrogen Biotechnology Co., Ltd (Thermo Fisher Scientific) were used to incubate for 1 hour to be visualized under a fluorescent microscope (Imager A1, Axio, Carl Zeiss Meditec AG, Jena, Germany). DAPI (2 mg/mL, Invitrogen Biotechnology Co., Ltd, Thermo Fisher Scientific) was used to stain the nuclei.

Effects of PPZ and/or antitumor agents on tumor growth in vivo

Solid tumor models were developed from the SGC7901/MDR cells. Forty-five male SCID immunodeficient BALB/C nude mice aged 4–6 weeks weighing 18–22 g (Model Animal Research Center of Nanjing University, People’s Republic of China) were randomly divided into four groups (n = 10 each in the control, PPZ, ADR, and PPZ + ADR groups; n=5 in each group). The mice were treated with PPZ (20, 40, 80 µg/mL) for 24 hours, and then tumor volumes were measured weekly by using a planimeter. The tumor volumes were calculated as follows: tumor volume = (length × width²)/2. The tumor volumes were measured weekly by using a planimeter. The tumor volumes were calculated as follows: tumor volume = (length × width²)/2. The experiment was terminated at day 21, at which point the mice were sacrificed and the tumors were excised and weighed. The data were analyzed by one-way ANOVA followed by post-hoc Dunnett’s test. *p < 0.05, **p < 0.01, and ***p < 0.001 were considered significant differences.

Table 1 The sequences of primers for real-time PCR

<table>
<thead>
<tr>
<th>Gene</th>
<th>Primer sequence</th>
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<tbody>
<tr>
<td>Actin</td>
<td>Forward 5′-AGCGGACATCCCCCAAAAGTT-3′</td>
</tr>
<tr>
<td></td>
<td>Reverse 5′-GGGCAAGGCTCATATT-3′</td>
</tr>
<tr>
<td>V-ATPases</td>
<td>Forward 5′-GTCCTAACCGGGGATATC-3′</td>
</tr>
<tr>
<td>(ATP6V0E)</td>
<td>Reverse 5′-AAAGAGAGGTTGTAGGCGGC-3′</td>
</tr>
<tr>
<td>MDR1</td>
<td>Forward 5′-TGCCGACAGGAGATAGCTG-3′</td>
</tr>
<tr>
<td></td>
<td>Reverse 5′-GCCAAATCACAAGGTTAGCTT-3′</td>
</tr>
<tr>
<td>MRPI</td>
<td>Forward 5′-CGGAAACCATCACCAGCCCTATCC-3′</td>
</tr>
<tr>
<td></td>
<td>Reverse 5′-ACCTTCTCCTGCACTCACCCTTGG-3′</td>
</tr>
<tr>
<td>PI3K</td>
<td>Forward 5′-TTAGCTTTCCCCACGAGGA-3′</td>
</tr>
<tr>
<td>(PIK3CA)</td>
<td>Reverse 5′-CACAATAGTGTCTGACCTC-3′</td>
</tr>
<tr>
<td>AKT</td>
<td>Forward 5′-CTGAGATTTGTCGACGCTGTA-3′</td>
</tr>
<tr>
<td>(AKT1)</td>
<td>Reverse 5′-CACAGCAAGTCTGATGTCACTA-3′</td>
</tr>
<tr>
<td>mTOR</td>
<td>Forward 5′-ATGGACGTCTGTCCGTTTTC-3′</td>
</tr>
<tr>
<td></td>
<td>Reverse 5′-AATCGACAGGCGCAAAGG-3′</td>
</tr>
<tr>
<td>HIF-1α</td>
<td>Forward 5′-TTTGGCAGAAGACACAGAC-3′</td>
</tr>
<tr>
<td></td>
<td>Reverse 5′-TGATTGAGTGCAGGTCAGC-3′</td>
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the normal group without bearing tumor), and were kept in a germ-free environment and their food, water, and bedding were autoclaved prior to use. Forty mice were subcutaneously injected with $3 \times 10^6$ SGC7901/MDR cells resuspended in 0.2 mL of RPMI-1640 containing 10% FCS in the right scapula. Once tumors became evident (at least $0.30 \times 0.30$ cm, approximately 10 days after the tumor cell injection), PPZ resuspended in normal saline (15 mg/mL) immediately before use was orally administered by gavage at a dose of 75 mg/kg. ADR was administered by intraperitoneal injection at a dose of 1.25 mg/kg simultaneously with PPZ oral treatment, 24 hours after PPZ treatment, or in mice that did not receive any PPZ treatment, or in mice that did only receive PPZ treatment, or in mice that even did not receive any treatment after inoculation, and or in mice that did not receive inoculation. Tumor dimensions were measured once a day with calipers and body weight were also monitored. Tumor volume was estimated by using the following formula: tumor volume (mm$^3$) = length (mm) × width$^2$ (mm$^2$)/2$^{25}$. The relative tumor volume or body weight was expressed as the $V_t$/ $V_o$ or $W_t$/ $W_o$ index, where $V_t$ and $W_t$ are the tumor volume or body weight on the day of measurement and $V_o$ and $W_o$ are the same indices at the start of the treatment. All mice were sacrificed at the end of the experiments, within 3 weeks after the injection of the human tumor cells. Tumor specimens were fixed in 10% (v/v) neutral formalin solution for 24 hours and processed routinely by embedding in paraffin. Tissue serial sections were cut at 4 mm. Then the TUNEL assay apoptotic cells in sections of mice tumor tissue were detected using an in situ apoptosis detection kit (KEYGEN, Nanjing City, Jiangsu Province, People’s Republic of China) as instructed by the manufacturer. Cells were visualized with a light microscope (Olympus IX70, Olympus Corporation, Tokyo, Japan). The apoptotic index was calculated as follows: the apoptotic index number of apoptotic cells/total number of cells. Meanwhile, the protein expressions of V-ATPases, PI3K, AKT, mTOR, HIF-1α, P-gp, and MRP1 also were determined in those tissues of bearing tumor of BALB/C nude mice. All in vivo experiments were approved by the ethical committee in the Affiliated Drum Tower Hospital of Nanjing University, Medical School.

Statistical analysis
All data were presented as the mean ± SD for at least three independent experiments. Statistical analysis was performed with SPSS software (version 22.0; IBM Corporation, Armonk, NY, USA). The significant differences between the two groups were then evaluated by One-way ANOVA. Statistical significance was defined as $P < 0.05$.

Results

PPIs pretreatment could inhibit mRNA levels of V-ATPases, MDR1, and MRP1
Under a medium with pH 6.65, the mRNA levels of V-ATPases, MDR1, and MRP1 significantly decreased after PPZ pretreatment (more than 20 µg/mL) for 24 hours on SGC7901/MDR cells when compared with those without PPZ pretreatment, which did not depend on a concentration manner (Figure 2A). As to SGC7901 cells, after PPZ pretreatment for 24 hours, the mRNA levels of MDR1 began to decrease when the concentration of PPZ was 40 µg/mL. Meanwhile, the mRNA levels of V-ATPases and MRP1 decreased when PPZ concentration was 80 µg/mL (Figure 2B). The amplification curve and dissolution curve are indicated in Figure 2C and D.

PPIs pretreatment could inhibit the mRNA levels of PI3K, AKT, mTOR, and HIF-1α
Under a medium with pH 6.65, PPZ pretreatment (20 µg/mL) for 24 hours on the SGC7901/MDR cells could inhibit mRNA of PI3K significantly compared with those without PPZ pretreatment ($P < 0.05$). PPZ pretreatment (40 µg/mL) for 24 hours on the SGC7901/MDR cells could inhibit the mRNA levels of AKT, mTOR, and HIF-1α ($P < 0.05$) (Figure 3A). As to the SGC7901 cells, PPZ pretreatment (40 µg/mL) for 24 hours could obviously inhibit the mRNA levels of PI3K than those without PPZ pretreatment ($P < 0.05$). The levels of mTOR mRNA began to decrease when PPZ pretreated for 24 hours with a concentration of 60 µg/mL ($P<0.01$). However, the levels of HIF-1α decreased when the PPZ concentration was 100 µg/mL ($P < 0.05$) (Figure 3B). The amplification curve and dissolution curve are indicated in Figure 3C and D.

PPIs inhibited V-ATPases and downregulated the expressions of P-gp and MRP1
After 24-hour treatment by Esomeprazole with different concentrations on the SGC7901/MDR cells in the medium of pH 6.65, compared with those in the control group (0 µg/mL), the expressions of V-ATPases and P-gp began to decrease when the concentration of Esomeprazole increased to 20 µg/mL ($P < 0.05$), and then, their expressions continued to decrease in a dose-dependent manner as the concentration of the Esomeprazole increased ($P < 0.05$), when the concentration was up to 100 µg/mL, their expressions were the lowest. Furthermore, compared to those in the
Figure 2 PPZ pretreatment on the SGC7901 cells and SGC7901/MDR cells could inhibit mRNA levels of V-ATPases, MDR1, and MRPI.

Notes: (A) PPZ pretreatment could inhibit mRNA levels of V-ATPases, MDR1 and MRPI of the SGC7901/MDR cells; (B) PPZ pretreatment could inhibit mRNA levels of V-ATPases, MDR1 and MRPI of the SGC7901 cells; (C) Rn vs. cycle; (D) Dissociation curve. (**P < 0.005; *P < 0.01; +P < 0.05).

Abbreviation: PPZ, pantoprazole.

control group, when the concentration of Esomeprazole was up to the 50 µg/mL, the expression of MRPI began to decrease (P < 0.05) and also continued to decrease in a dose-dependent manner as the concentration of the Esomeprazole increased (P < 0.05) (Figure 4A). However, under the same conditions, after the SGC7901 cells were treated by using the same way, the expressions of V-ATPases did not decrease as the concentration of the Esomeprazole increased (P > 0.05), and the expressions of MRPI and P-gp were not affected either (P > 0.05) (Figure 4B). PPZ with different concentrations also was used to treat the SGC7901/MDR cells in the same medium for 24 hours. Compared with those in the control group, the expression of V-ATPases started to decline when the PPZ concentration was up to 10 µg/mL (P < 0.05). And with the increasing concentration of PPZ, its expression continued to decrease in a dose-dependent manner (P < 0.05) (Figure 4C).

Subcellular localization of V-ATPases and P-gp in the SGC7901/MDR cells, before and after PPZ (100 µg/mL) pretreatment for 24 hours, were detected with the immunofluorescent staining analysis (Figure 5). And similar results were indicated in Figure 6 that subcellular localization of V-ATPases, HIF-1α, and MRPI in SGC7901/MDR cells before and after PPZ treatment (100 µg/mL) for 24 hours. It could be inferred that PPIs could change and further inhibit the subcellular localization of V-ATPases, HIF-1α, P-gp, and MRPI.

PPIs inhibited the expressions of PI3K/Akt/mTOR/HIF-1α signaling pathway

SGC7901/MDR cells were dealt with Esomeprazole with different concentrations in the medium of pH 6.65 for 24 hours, compared with the control (0 µg/mL), the expressions of the PI3K and HIF-1α began to decrease (P < 0.05) and as the concentration of the PPZ increased, their expressions continued to decrease dose dependently (P < 0.05) (Figure 4C).
Esomeprazole was up to the 20 µg/mL ($P < 0.05$), and then, their expressions continued to be inhibited in a dose-dependent manner as the concentration of the Esomeprazole increased ($P < 0.05$). When the concentration was up to 100 µg/mL, however, their expressions were the lowest. Furthermore, compared to the control group, when the concentration of Esomeprazole went to 50 µg/mL, the expression of Akt and mTOR began to decline ($P < 0.05$) and also continued to decrease dose dependently as the Esomeprazole concentration increased ($P < 0.05$) (Figure 7A).

However, under the same conditions, the expression of PI3K decreased only when the Esomeprazole concentration was up to 100 µg/mL ($P < 0.05$) by using the same way to treat the SGC7901 cells, and the expressions of the Akt, mTOR, and HIF-1α were not affected ($P > 0.05$) (Figure 7B).

Furthermore, the SGC7901/MDR cells were dealt with the PPZ in the same medium for 24 hours. The expressions of PI3K, Akt, mTOR, and HIF-1α started to decrease when the PPZ concentration was up to 50 µg/mL ($P < 0.05$). And as the concentration of PPZ increased, their expressions continued to decrease dose dependently ($P < 0.05$) (Figure 7C).

At the same time, after Esomeprazole with different concentrations was used to treat the SGC7901/MDR cells, our study demonstrated that the expressions of TSC1 and TSC2 and Rheb started to decline when the Esomeprazole concentration up to 50 µg/mL ($P < 0.05$). And as the concentration of the Esomeprazole increased, its expression continued to decrease in a dose-dependent manner (Figure 8A). However, under the same conditions by using the same way to treat the SGC7901 cells, the expression of the TSC1 declined only when the Esomeprazole concentration was up to 100 µg/mL ($P < 0.05$), and the TSC2 expression decreased when the concentration was up to 80 µg/mL, but as the Esomeprazole concentration increased, TSC2 expression did not decline any more. However, the Esomeprazole did not affect the

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**Figure 3** PPIs pretreatment on the SGC7901 and the SGC7901/MDR cells for 24 hours could affect the mRNA levels of PI3K, AKT, mTOR, and HIF-1α.

**Notes:** (A) PPZ pretreatment could inhibit mRNA levels of PI3K, AKT, mTOR and HIF-1α in the SGC7901/MDR cells; (B) PPZ pretreatment could inhibit mRNA levels of PI3K, AKT, mTOR and HIF-1α in the SGC7901 cells; (C) Run vs. cycle; (D) dissociation curve. (***$P < 0.005$; **$P < 0.01$; *$P < 0.05$).

**Abbreviation:** PPIs, proton pump inhibitors.
expression of the Rheb at any concentration (P > 0.05) (Figure 8B).

After pretreatment by Esomeprazole or PPZ for 24 hours in the SGC7901/MDR cells, the expressions of the phosphor-Ser939 and phosphor-Thr1462, which were the two phosphorylation sites of Akt to activate TSC2, also downregulated dose dependently. And the expression of the phosphor-Akt substrate decreased gradually as the concentration of the Esomeprazole or the PPZ increased (Figure 9).

Inhibition of the PI3K/Akt/mTOR/HIF-1α signaling pathway and P-gp protein expression by V-ATPases siRNA interference

V-ATPases siRNA was used to process SGC7901/MDR cells for 24 hours, intracellular V-ATPases expression was significantly inhibited (P < 0.05), and the intracellular protein expressions of the PI3K, Akt, mTOR, and HIF-1α could be significantly inhibited (P < 0.05), and then, the final resistance
protein P-gp expression decreased ($P < 0.05$). However, the intracellular expressions of the TSC1 and TSC2 also were inhibited by V-ATPases siRNA ($P < 0.05$) (Figure 10).

**Inhibition of mTOR expression could downregulate the intracellular HIF-1α and P-gp expression**

After pretreatment by Rapamycin on the SGC7901/MDR cells for 24 hours, compared with the control group, the intracellular expression of the mTOR was significantly inhibited when the concentration of Rapamycin was up to the 20 $\mu$g/mL, and as the concentration of Rapamycin increased, the expression of mTOR continued to decrease, and when the concentration of Rapamycin was up to 80 $\mu$g/mL, the intracellular mTOR expression could not be detected anymore. And at the same time, HIF-1α and P-gp, the downstream protein of mTOR, significantly decreased when the mTOR expression was inhibited. And their intracellular
protein expressions gradually dropped with the rise of the Rapamycin’s concentration, showing the remarkable concentration dependent manner (Figure 11A and B).

### PPIs on TSC1/2 complex expressions time dependently

The SGC7901/MDR cells were treated by Esomeprazole (80 µg/mL) for 3, 6, 12, 18, 24 hours, respectively. And the SGC7901/MDR cells were treated by Esomeprazole (80 µg/mL) for 15 minutes as the control. Compared with the control, after pretreatment for 3 hours, the intracellular expression of the TSC2 began to increase (*P* < 0.05 vs control). For 24 hours, the TSC2 expression significantly decreased compared with the control group (*P* < 0.05). Simultaneously, Esomeprazole (80 µg/mL) pretreated the SGC7901/MDR cells for 3 hours, compared to the control group, the intracellular expression of the TSC1 began to increase (*P* < 0.05), and reached the peak after pretreatment.
for 12 hours. And as the expression of TSC2, after 18 hours, the expression of TSC1 had no significant difference with the control group ($P > 0.05$). But for 24 hours, the TSC1 expression also significantly decreased compared with the control group ($P < 0.05$) (Figure 12).

Inhibitory effects of LY294002 on protein expressions of PI3K, AKT, HIF-1α, P-gp, and MRPI
LY294002 with various concentrations (including 0, 10, 20, 50 µmol/L) was used to treat the SGC7901/MDR cells for 12 hours. And as the expression of TSC2, after 18 hours, the expression of TSC1 had no significant difference with the control group ($P > 0.05$). But for 24 hours, the TSC1 expression also significantly decreased compared with the control group ($P < 0.05$) (Figure 12).

**Figure 8** Effects of PPIs pretreatment for 24 hours on the expressions of the TSC1, TSC2, and Rheb in the SGC7901/MDR and SGC7901 cell lines.

**Notes:** (A) After Esomeprazole with different concentration to treat the SGC7901/MDR, the expressions of the TSC1, TSC2, and the Rheb decreased in a dose-dependent manner as the concentration of the Esomeprazole increased. (B) Under the same conditions, using the same way to treat the SGC7901 cells, the expression of the TSC1 declined only when the Esomeprazole concentration was up to 100 µg/mL, and the TSC2 expression decreased when the concentration was up to 80 µg/mL, but as the Esomeprazole concentration increased, the TSC2 expression did not decline any more ($^*P < 0.05$ vs the control).

**Abbreviation:** PPI, proton pump inhibitor.

**Figure 9** Effects of PPIs pretreatment for 24 hours on the expressions of the phosphor-Ser939, phosphor-Thr1462, and phosphor-Akt substrate in the SGC7901/MDR cells.

**Notes:** (A) Effects of the Esomeprazole on the expressions of the phosphor-Ser939, phosphor-Thr1462, and the phosphor-Akt substrate. (B) Effects of the Pantoprazole on the expressions of the phosphor-Ser939, phosphor-Thr1462, and the phosphor-Akt substrate.
**Figure 10** Inhibitory effect on the PI3K/Akt/mTOR/HIF-1α signaling pathway in the SGC7901/MDR cell lines by interference of V-aTPases.

**Notes:** (A) Western blot analysis of V-aTPases by siRNA-mediated inhibition. Interference system could effectively inhibit V-aTPases and positive control GAPDH. (B) After the expression of V-aTPases was downregulated by siRNA, the expression of the PI3K/Akt/mTOR/HIF-1α signaling pathway was significantly inhibited simultaneously. And the expression of P-gp also was downregulated (*P < 0.05 vs the blank group).

**Abbreviation:** V-aTP, V-aTPases.

**Figure 11** Effects of the Rapamycin inhibition on the protein expressions of mTOR, HIF-1α, and P-gp in the SGC7901/MDR cells.

**Notes:** (A) Inhibitory effects of Rapamycin on the protein expressions of mTOR, HIF-1α, and P-gp. (B) Relative optical density value of protein expressions of mTOR, HIF-1α, and P-gp after Rapamycin treatment (*P < 0.05 vs the control).

**Figure 12** Effects of PPis on the Tsc1/2 complex expressions time-dependently.

**Notes:** (A) Effects of PPis on the Tsc1/2 complex expressions in a time-dependent manner. (B) Relative optical density value of Tsc1/2 complex expressions after PPis treatment. (*P < 0.05 vs the control, showing the expression increase, ^P < 0.01 vs the control, showing the expression decrease.)

**Abbreviations:** PPis, proton pump inhibitors; V-aTP, V-aTPases.
tumor volume of athymic mice, and the changes of relative body weight of athymic mice, relative to the Ctrl group since there was a relatively higher apoptotic index in the PPZ + ADR group (Figure 15A). According to the protein expressions of bearing-tumor tissues, the expressions of PI3K, mTOR, HIF-1α, and P-gp were significantly inhibited in the PPZ + ADR group, which also further supported our results in vitro. However, the expressions of V-ATPases, AKT, and MRP1 were not inhibited in the PPZ + ADR group, which were inconsistent with the data in our in vitro experiments (Figure 15B and C).

Discussion

Intrinsic and acquired MDR are the primary causes for limited efficacy of chemotherapy in the majority of gastrointestinal malignancies, including gastric cancer. Drug resistance represents a complex and multifactorial phenomenon related to tumor microenvironment, for example, hypoxia, acidosis and inflammation, as well as the neoplastic cell. Tumor cell metabolism is being nowadays the object of renewed consideration for better understanding of cancer biology and therapeutic strategies. In fact, the glucose metabolism in hypoxic conditions by the neoplasms leads to a pH drift toward acidity. The acid microenvironment is modulated through the over-expression of H+ transporters that also are involved in tumor progression, invasiveness, distant spread, and chemoresistance. Several strategies to block/downmodulate the efficiency of these transporters are currently being investigated. Among them, PPIs have been utilized to successfully block the H+ transporters in vitro and in vivo, leading to apoptotic death. Furthermore, their action seems to synergize with conventional chemotherapy protocols, leading to chemosensitization and reversal of MDR. Evidence is demonstrated that hypoxia and acidity are involved in cancer progression and the MDR of tumor to chemotherapy. Besides, hypoxia and acidity also may contribute to the progression from benign to malignant growth. Tumor acidity, in particular, has role in resistance chemotherapy and metastatic behaviour. V-ATPases play a critical role in regulating the transmembrane pH gradient. The subcellular localization of V-ATPases is mainly on the membrane of internal acidic vesicles and plasma membrane. Compared to the normal cells, in which V-ATPases only expressed on the acidic vesicle.
are very active in malignant cells, correlating with a high expression at both the plasma membrane and acidic vesicle level.\(^8,^9,^{31}\) PPIs were the powerful tools in the treatment of human solid tumors because they could induce sensitivity to a wide range of antitumor drugs by inhibiting the expression of V-ATPases.\(^10,^{11,^{32}}\) One major cause of resistance to chemotherapy has been correlated with the presence of the “pumps” that actively transport drugs out of the cell. The most prevalent transporters of MDR are p-gp and MRP1.\(^{33}\) Our studies demonstrated that both PPZ and Esomeprazole could inhibit the intracellular expressions of the V-ATPases with the simultaneous suppression of expressions of P-gp.

Figure 14 Effects of PPZ on sensitivity of human tumors to antitumor agents in athymic mice engrafted with the SGC7901/MDR cells in vivo.

Notes: (A–E) Effects of PPZ and/or ADR on tumor size of athymic mice xenografted with the SGC7901/MDR cells. (F) Effects of PPZ and/or ADR on relative body weight of athymic mice xenografted with the SGC7901/MDR cells. (G) Effects of PPZ and/or ADR on relative tumor volume of athymic mice xenografted with the SGC7901/MDR cells. (H) Effects of PPZ and/or ADR on tumor volume and relative tumor volume of athymic mice xenografted with the SGC7901/MDR cells. PPZ resuspended in normal saline (15 mg/mL) immediately before use was orally administered by gavage at a dose of 75 mg/kg. ADR was administered by intraperitoneal injection at a dose of 1.25 mg/kg. RTV means relative tumor volume representing the ratio of tumor volume at the end and tumor volume at the beginning.

Abbreviations: ADR, adriamycin; PPZ, pantoprazole.
and MRP1 in a dose-dependent manner. And the subcellular localization of the V-ATPases and P-gp in SGC7901/MDR cells differed greatly before and after PPIs pretreatment. Furthermore, downregulating the intracellular expression of V-ATPases by siRNA interference also could inhibit the expression of P-gp. These results implied that PPIs could inhibit the expression of V-ATPases, reverse the intracellular and extracellular pH gradients, and then downregulate the expression of the resistance protein so as to improve the chemosensitivity and reverse MDR. However, so far, the exactly intracellular signal mechanism of PPIs to reverse the drug resistance has been not yet known.

Evidences suggested that the acid pH could activate the PI3K/AKT signaling pathway and could significantly enhance phosphorylation of AKT. Many of these cellular growth and metabolism involve signaling through the PI3K/AKT/mTOR pathway. In particular, the mTOR pathway appears to play a central role in the development of multiple cancers. Although often called “a master regulator”, mTOR is one signaling target in an intricate signaling cascade that controls cell growth and angiogenesis in both normal and cancerous conditions. Other important factors in this pathway include upstream activators such as PI3K and AKT, negative regulators such as the
tuberous sclerosis complex 1/2 (TSC1/2), and downstream effectors such as the transcription factor hypoxia-inducible factor 1α (HIF-1α). Evidence demonstrated that amplified signaling through phosphoinositide 3-kinase, in turn, phosphorylating and activating Akt and its downstream target, mTOR, enhances HIF-1-dependent gene expression in certain cell types; and rapamycin, as the specific mTOR inhibitor, inhibits both the accumulation of HIF-1α and HIF-1-dependent transcription induced by hypoxia. HIF-1 constitutes a pivotal regulator of cellular adaptation to hypoxia and has been implicated in MDR. HIF-1α is centrally involved in multiple aspects of tumorigenesis including tumor cell proliferation, angiogenesis, metastasis, as well as the response to chemotherapy and radiotherapy. HIF-1α is overexpressed in a vast number of solid tumors, and tumorous HIF-1α expression is often associated with poor prognosis. Therefore, the PI3K/Akt/mTOR/HIF-1α signaling pathway has been reported closely to be linked to chemosensitivity. Our studies clarified that PPIs could inhibit the intracellular protein expressions of PI3K/AKT/mTOR/HIF-1α signaling pathway and also could decrease the phosphorylation of Akt substrate. Akt was the first kinase demonstrated to directly phosphorylate the TSC1-TSC2 complex in cells in response to growth factors. And the main sites of the human TSC2 to be phosphorylated by Akt were Ser939 and Thr1462. And after the pretreatment by PPIs, the phosphorylations of the two sites, Ser939 and Thr1462, were significantly inhibited as the concentration of PPIs in the SGC7901/MDR cells increased. After the V-ATPases were knocked down by the siRNA in the SGC7901/MDR cells, the intracellular protein expressions of PI3K/Akt/mTOR/HIF-1α signaling pathway also were significantly inhibited, suggesting that PPIs could inhibit the PI3K/Akt/mTOR/HIF-1α signaling pathway by the way of downregulating the expression of V-ATPases.

TSC1 and TSC2 are the tumor-suppressor genes mutated in the tumor syndrome TSC (tuberous sclerosis complex). The TSC1–TSC2 (hamartin–tuberin) complex, through its GAP (GTPase-activating protein) activity toward the small G-protein Rheb (Ras homologue enriched in brain), is a critical negative regulator of mTORC1 (mammalian target of rapamycin complex 1). PPIs could inhibit the expression of Rheb dose dependently, and simultaneously the expression of the TSC1-TSC2 complex increased significantly compared with the control group. So, PPIs could inhibit the expression of mTOR through the TSC1-TSC2 complex signaling pathway. However, our results showed that after 24-hour pretreatment by PPIs on the SGC7901/MDR cells, the V-ATPases siRNA interference could inhibit the expressions of TSC1–TSC2 complex, and at the same time, the intracellular expression of mTOR also decreased with the V-ATPases, suggesting negative feedback involving various mechanisms of activation. The conclusion was confirmed through the inhibitory effects of PPIs on TSC1/2 complex expressions not only dose dependently but also time dependently. The TSC1/2 complex expressions kept on increasing from the beginning and then decreasing while the underlying molecular mechanisms remained largely elusive. Also, PPIs treatment inhibited the expressions of Rheb dose dependently. Therefore, we speculated that TSC1/2 complex and Rheb might be greatly involved in the effects of PPIs on regulating the PI3K/AKT/mTOR/HIF-1α signaling pathway.

At this time, we found that downregulating the expression of V-ATPases could inhibit the expressions of PI3K/Akt/mTOR/HIF-1α signaling pathway as well as of P-gp and MRP1. Meanwhile, siRNA interference on the expression of V-ATPases also could downregulate the expressions of TSC1/2 complex. But it was not yet clarified whether the PPIs inhibited the P-gp and MRP1 was via the PI3K/Akt/mTOR/HIF-1α signaling pathway or not. mTORC1 is strongly sensitive to inhibition by the naturally occurring compound rapamycin. And the mTOR protein function has relied heavily on the use of Rapamycin and are therefore probably specific to mTORC1, which was located the downstream of PI3K/Akt, and upstream of HIF-1α. Our data showed that mTOR protein expression, as well as the HIF-1α and P-gp, decreased in a dose-dependent manner as the concentration of the rapamycin increased. Therefore, we predicted that P-gp might locate the downstream of mTOR. Evidence had demonstrated that HIF-1α determined gastric cancer chemosensitivity via modulation of p53 and NF-κB. However, the relationship between drug-resistance proteins such as P-gp or MRP1 and HIF-1α was not yet been clarified. The underlying molecular mechanisms as well as the role of HIF-1α for MDR under hypoxic conditions remain largely elusive.

Conclusion
PPI inhibits the expression of V-ATPases, downregulates the MDR so as to sensitize the SGC7901/MDR cells to the antitumor drugs. Our study added some important information on the preclinical setting-up for the use of PPIs in the treatment of a poorly treatable tumor such as human gastric cancer. And simultaneously, the PI3K/Akt/mTOR/HIF-1α signaling pathway has been linked to chemosensitivity. Downregulation of V-ATPases expression by PPIs could
inhibit the PI3K/AKT/mTOR/HIF-1α signaling pathway to decrease the expressions of P-gp and MRP1 and facilitate the chemosensitivity of SGC7901/MDR to the anticancer drugs. Among them, PPIs might also downregulate TSC1/TSC2 complex and Rheb which might be involved into the regulation mechanism of PI3k/AKT/mTOR/HIF-1α signaling pathway. However, the possible relevant signaling pathway of PPIs was involved into enhancing the chemosensitivity and reversing MDR also needs further research in vivo.

Conclusion
PPIs could reverse MDR via downregulating V-ATPases/PI3K/Akt/mTOR/HIF-1α signaling pathway through TSC1/2 complex and Rheb in human gastric adenocarcinoma cells in vitro and in vivo.

Data sharing statement
All the data involved in our study are available.

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Author contributions
All authors made substantial contributions to conception and design, acquisition of data, or analysis and interpretation of data; took part in drafting the article or revising it critically for important intellectual content; gave final approval of the version to be published; and agree to be accountable for all aspects of the work.

Disclosure
The authors report no conflicts of interest in this work.

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