

CD13 downregulation mediated by ubenimex inhibits autophagy to overcome 5-FU resistance by disturbing the EMP3/FAK/NF-κB pathway in gastric cancer cells

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Background: Gastric cancer (GC) is one of the most common malignant tumours in China, but the efficacy of chemotherapy on GC is significantly reduced due to the occurrence of drug resistance. Some studies have shown that the expression level of CD13 is associated with tumour resistance, but whether ubenimex, as a CD13 inhibitor, reverses GC drug resistance and the underlying mechanism remain unclear. **Methods:** Herein, resistance to 5-fluorouracil (5-FU) was reversed in GC by ubenimex, and the underlying mechanism was determined using Cell Counting Kit-8 (CCK-8) assays, gene chip analysis, high content screening (HCS), transmission electron microscopy, flow cytometry, immunofluorescence and western blot

Results: Flow cytometry, transmission electron microscopy and immunofluorescence analyses indicated that ubenimex, an inhibitor of CD13, regulated the autophagy and apoptosis of SGC7901/5-FU cells by downregulating CD13 expression. In addition, Gene chip analysis and HCS demonstrated that epithelial membrane protein 3 (EMP3)/focal adhesion kinase (FAK) was a putative signalling pathway downstream of CD13. Furthermore, western blot analyses showed that ubenimex not only inhibited EMP3, FAK and nuclear factor- κ B (NF- κ B) expression but also suppressed GC autophagy and activated apoptosis by targeting CD13. These findings indicated a potential mechanism via the CD13/EMP3/FAK/NF- κ B pathway and that the activity of which was restrained.

Conclusions: Ubenimex affects autophagy and apoptosis to reverse GC cell resistance by targeting the CD13/EMP3/FAK/NF-κB pathway. These results showed that ubenimex is a promising agent that may inhibit GC autophagy to improve chemotherapeutic drug sensitivity and thereby reverse drug resistance.

Keywords: Gastric cancer (GC); CD13; ubenimex; autophagy; epithelial membrane protein 3 (EMP3)

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Introduction

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As the fourth most common cancer, gastric cancer (GC) is ranked as the second leading cause of cancer-related death worldwide (1). Patients with early GC are usually asymptomatic, and the rate of early diagnosis of GC is

low; most patients (>70%) present with advanced GC (2). Currently, 5-fluorouracil (5-FU)-based chemotherapy is widely used to enhance the quality of life and extend the survival cycle of patients with advanced GC. However, experimental studies in the clinic have revealed that the sensitivity of GC patients to 5-FU is gradually decreasing, making the treatment for GC patients end in failure (3,4). 5-FU resistance has been demonstrated to be a fundamental obstacle in the chemotherapy of GC, and identifying ways to reverse 5-FU resistance in GC has become an important issue to be addressed.

Autophagy is a process by which cells use lysosomal enzymes to degrade harmful substances and maintain a stable intracellular environment (5). The traditional view indicates that autophagy is an adaptive response made by normal cells to tolerate metabolic stress environments that are not conducive to their own survival to maintain the normal physiology of cells (6). However, with the evolving understanding of the molecular mechanisms involved in autophagy, it has been found that autophagy produces a marked effect on the development of drug resistance. To relieve chemotherapy-induced stress, tumour cells activate autophagy to defend against stress, leading to cytoprotective effects called chemoresistance (7). Studies have shown that autophagy mediates tumour escape from anoikis and promotes tumour cell survival; therefore, cell autophagy induces the development of drug resistance in GC, which is closely related to its triggering of apoptosis tolerance (8,9).

CD13, also known as aminopeptidase N (APN), belongs to the class II membrane-bound metalloproteinases. CD13 is overexpressed in liver, stomach, pancreas, colon, prostate and thyroid tumours, and it utilizes the action of proteolytic enzymes to accelerate tumour proliferation, vascularization, metastasis and infiltration (10). A previous study has found that ubenimex, a CD13 inhibitor, synergizes with 5-FU to improve the antitumour activity of 5-FU in liver, kidney and breast cancer cells (11). Ubenimex also targets CD13, promotes apoptosis induced by chemotherapeutic agents and reverses multidrug resistance (MDR) in liver cancer cells (12). However, as the only marketed CD13 inhibitor, ubenimex is only used as an immunomodulatory adjuvant for the treatment of haematological malignancies (13). There are no reports on the reversal of 5-FU resistance mediated by ubenimex.

Initially discovered to undergo hypermethylationmediated transcriptional silencing in glioma, non-smallcell lung cancer (NSCLC) and oesophageal squamous cell carcinoma (ESCC), epithelial membrane protein 3 (EMP3) has attracted attention as a well-established tumour suppressor (14-16). However, accumulating evidences suggest a tumour-promoting role for EMP3 in the breast, urothelium and hepatocytes (14,17,18). Although there is evidence showing that the mRNA of EMP3 is upregulated in GC-derived cell lines (19), the role of EMP3 in GC malignancy development and its impact on GC treatment are currently unknown.

Focal adhesion kinase (FAK) is a cytoplasmic nonreceptor tyrosine kinase that mediates the adhesion of cells to the extracellular matrix (ECM) and plays a crucial role in regulating the function of basic cells with overexpression in many cancers, including glioblastoma, breast, colorectal, pancreatic, lung and ovarian cancers (20,21). Evidence has indicated that the expression and phosphorylation of FAK play an important role in the initiation and transmission of autophagy signals as well as the induction of apoptosis tolerance in GC cells (22). Blockade of FAK expression and phosphorylation activity inhibits the entry of FAK into the nucleus, which attenuates the transcriptional activity of nuclear factor- κ B (NF- κ B), thereby downregulating the expression of autophagy proteins, such as Beclin-1, and inhibiting the occurrence of autophagy (23). Among autophagy proteins, Beclin-1 plays a key role in the process of apoptosis resistance induced by autophagic "addiction". Inhibition of Beclin-1 expression activates caspase-mediated apoptosis in tumour cells, whereas activated caspases induce apoptosis in tumour cells by affecting Beclin-1, thereby interfering with Bcl-2 and Bax protein levels (24,25).

In the present study, we confirmed that ubenimex alleviates GC autophagy to reverse 5-FU resistance by attenuating the expression of CD13. Additionally, we utilized high content screening (HCS) and found that EMP3 is a key molecule in the targeting of CD13 by ubenimex in suppressing GC, and through a series of experiments, we verified that ubenimex may attenuate GC autophagy by affecting the CD13/EMP3/FAK/NF- κ B pathway. Collectively, these results indicated that ubenimex is a prospective agent that may attenuate GC autophagy to reverse 5-FU resistance, thus improving GC sensitivity to 5-FU. We present the following article in accordance with the MDAR reporting checklist (available at https://tcr. amegroups.com/article/view/10.21037/tcr-22-345/rc).

Methods

Reagents and antibodies

Ubenimex was purchased from Shenzhen Main Luck Pharmaceutical, Inc. (Shenzhen, China), and 5-FU was obtained from Xudong Haipu Pharmaceutical Co., Ltd. (Shanghai, China). Rabbit monoclonal antibodies against ANPEP (CD13; Abcam Biotechnology, Cambridge, USA;

Translational Cancer Research, Vol 11, No 8 August 2022

cat. no. ab108382), LC3B (Abcam; cat. no. ab192890), Bax (ABclonal Technology Co., Ltd., Wuhan, China; cat. no. A19684), p-FAK (Abcam; cat. no. ab81298), Rabbit polyclonal antibodies against EMP3 (Abcam; cat. no. ab236671), PTK2 (Sangon Biotech Co., Ltd., Shanghai, China; KleanAB; cat. no. P102163), NF-κB P65 (Proteintech, Rosemont, IL, USA; cat. no. 10745-1-AP), P62/SQSTM1 (Proteintech; cat. no. 18420-1-AP), Beclin-1 (Proteintech; cat. no. 11306-1-AP), ATG5 (Proteintech; cat. no. 10181-2-AP), Bax (Proteintech; cat. no. 50599-2-Ig), Bcl-2 (ABclonal; cat. no. A0208 and Proteintech; cat. no. 12789-1-AP), GAPDH (ABclonal; cat. no. AC001). All secondary antibodies (cat. no. SA00001-2) were purchased from Proteintech Biosciences.

Establishment of 5-FU-resistant GC cells

The human GC cell line, MKN-45, was purchased from the Cell Bank of the Chinese Academy of Sciences (Shanghai, China). When MKN-45 cells were in logarithmic growth phase, medium containing 5-FU (2 µg/mL) or oxaliplatin (0.1 µg/mL) was added. The above steps were repeated over 7 months until induction of MDR in cells (designated MKN-45/X) was confirmed. The human GC cell line, SGC-7901, was purchased from the Cell Bank of the Chinese Academy of Sciences (Shanghai, China). We generated SGC-7901/5-FU cells by exposing the parental cells to increasing concentrations of 5-FU, ranging from 0 to 80 µg/mL, for more than 3 months. The resistance index (RI) was estimated by the following formula: $RI = IC_{50}$ (resistant cells)/IC₅₀ (parental cells). The morphology and RI of drugresistant cells are shown in Figure S1A,S1B. Furthermore, CD13 expression was up-regulated in SGC-7901/5-FU cells, compared to the parental GC cells (Figure S1C).

Plasmid construction and gene amplification

Enhanced green fluorescent protein (EGFP)-labeled ANPEP (CD13; GenBank number NM_001150), EMP3 (GenBank number NM_001425) overexpression plasmids and negative control plasmids (empty vector control plasmid; CMV-MCS-3FLAG-IRES-EGFP-SV40-Neomycin) were used in this study. Detailed information is provided in Appendix 1.

Gene chip analysis and HCS of cell function

The Affymetrix $Clariom^{TM} S$ gene chip and the HCS were

used to were used to screen the key genes involved in the reversal of resistance to ubenimex. Detailed information is provided in Appendix 1.

Proliferative activity and cell sensitivity assays

The proliferative activity and sensitivity of GC cells to 5-FU were determined by the Cell Counting Kit-8 (CCK-8) assay. Detailed information is provided in Appendix 1.

Western blot analysis

After cells were incubated with different treatments, protein expression levels were determined. The detailed procedure is provided in Appendix 1.

Transmission electron microscopy

The human 5-FU-resistant cell line, SGC-7901/5-FU, was treated with 0.12 mg/mL ubenimex for 24 h. Cells were then fixed with 2.5% glutaraldehyde in PBS buffer for 1.5 h followed by fixation in 1% osmium tetroxide for 1.5 h. Cells were then washed, stained with 3% aqueous uranyl acetate, dehydrated with an increasing concentration gradient of ethanol and embedded in Araldite. Ultrathin sections were observed under a CM-120 electron microscope (Philips, USA).

Annexin V-FITC/PI staining

Annexin V-FITC/PI dual staining was used to determine the effect of ubenimex on apoptosis. Briefly, the indicated cells were treated with different concentrations of 5-FU and ubenimex for 48 h and then harvested. Cells were resuspended in 300 μ L of 1× binding buffer, incubated with Annexin V-FITC and PI in the dark for 15 min and then analysed using an EPICS XL flow cytometer (BD Biosciences, USA).

Immunofluorescence analysis

Cell concentration smears were made, and cells were fixed with 4% paraformaldehyde for 20 min. Cells were thoroughly washed with cold phosphate-buffered saline (PBS) (3 min \times 3), fixed with 4% paraformaldehyde for 20 min at room temperature and washed with cold PBS (3 min \times 3). Goat serum blocking solution (100 µL/sample) was added for 60 min at room temperature followed by incubation



Figure 1 Ubenimex reverses the drug resistance of SGC-7901/5-FU cells. (A) SGC-7901/5-FU cells were incubated in the presence or absence of ubenimex (0.12 mg/mL) for 24 h, followed by treatment with increasing concentrations of 5-FU (0.5, 1, 2, 4, 16, 32 µg/mL) for 48 h. (B) SGC-7901/5-FU cells treated with or without ubenimex (0.12 mg/mL) for 24 h were treated with 5-FU (12 µg/mL) for 0–72 hours. Cell viability was determined by the CCK-8 method. The results are expressed as the means ± SD of three replicates. *P<0.05 and **P<0.01. OD, optical density; 5-FU, 5-fluorouracil; Ube, ubenimex; CCK-8, Cell Counting Kit-8; SD, standard deviation.

with primary antibodies diluted 1:200 in blocking solution at 4 °C overnight in the dark. Cells were then washed in cold PBS (3 min ×3) and then incubated 1:200 fluorescent secondary antibody at 100 μ L/sample (diluted in PBS) at room temperature in the dark for 1 h. Cells were washed with cold PBS (5 min ×3) and then incubated with 4',6-diamidino-2-phenylindole (DAPI) for 15 min. Cells were washed with cold PBS (3 min ×3) and then incubated with an anti-fluorescent quencher.

Statistical analysis

GraphPad Prism 7.0 software (La Jolla, CA, United States) was utilized for statistical analysis (GraphPad Prism, RRID:SCR_002798). Each group of independent experiments was performed at least three times. All data are presented as the means \pm standard deviation (SD) and were evaluated by one-way analysis of variance. P<0.05 indicated statistical significance. Statistically significant P values are presented as *P<0.05, **P<0.01, ***P<0.001, and ****P<0.0001.

Results

Ubenimex reverses the drug resistance of SGC-7901/5-FU cells

Ubenimex acts as a CD13 inhibitor to induce immune activation by stimulating CD16⁺CD56⁺ NK cells and CD3⁺CD4⁺ T lymphocytes. Ubenimex can be used as an immunopotentiator for the treatment of leukaemia and multiple myeloma. However, there are currently few studies on the use of ubenimex for the treatment of GC or reversal of drug resistance in GC-resistant cells. We next investigated the effects of ubenimex on 5-FU sensitivity. SGC-7901/5-FU cells were incubated with various concentrations of 5-FU (0.5, 1, 2, 4, 16 and 32 µg/mL) for 48 h in the absence or presence of ubenimex (0.12 mg/mL) or were incubated with 5-FU (12 µg/mL) for 0–72 hours in the absence or presence of ubenimex (0.12 mg/mL). The CCK-8 results showed that ubenimex significantly increased the sensitivity of SGC-7901/5-FU cells to 5-FU in a timeand dose-dependent manner (*Figure 1A*,1B). Therefore, these findings suggested that ubenimex may reverse the drug resistance of acquired drug-resistant GC cells.

Ubenimex inhibits autophagic death in SGC-7901/5-FU cells

Previous experimental evidence has demonstrated that ubenimex inhibits autophagic cell death (26). In the present study, western blot analyses, cellular immunofluorescence and transmission electron microscopy were used to examine the autophagy level in SGC-7901/5-FU cells after 48 h of exposure to ubenimex following pretreatment with 5-FU. With increasing doses of ubenimex, LC3B, Beclin-1 and ATG5 levels decreased in SGC-7901/5-FU cells, whereas the levels of P62 showed a tendency to increase (*Figure 2A*). These findings indicated that the increased dose of ubenimex plays a key role in inhibiting a higher level of autophagic cell death in GC-resistant cells. LC3B attachment to the membrane is widely regarded as a sign of autophagosome formation (27). Hence, SGC-7901/5-FU cells were treated the same as described above and further



Figure 2 Ubenimex inhibits autophagic death in SGC-7901/5-FU cells. (A) Western blotting of autophagy associated protein expression in SGC-7901/5-FU cells after treatment for 48 h with 0.06 or 0.12 mg/mL ubenimex. GAPDH is shown as a loading control. Quantification was performed by a densitometry analysis and normalized to GAPDH levels. The bars indicate the means ± SD from the 3 independent experiments (*P<0.05, **P<0.01, ***P<0.001, and ****P<0.0001). (B) Immunofluorescence staining of LC3B with 0.06 or 0.12 mg/mL ubenimex treatment on SGC-7901/5-FU cells. (C) SGC-7901/5-FU cells were treated with 0.12 mg/mL ubenimex for 12 h. Then the cells were collected and prepared for electron microscopy analysis. The arrows indicate the appearance of autophagosomes (scale bar, 2 µm). Untreated cells were used as control. (D) Quantification of the autophagosomes of 3 independent experiments is shown. ***P<0.001 vs. control untreated cells. Ube, ubenimex; 5-FU, 5-fluorouracil; SD, standard deviation.

examined LC3B expression levels by immunofluorescence. We found that the results were consistent with previous conclusions (*Figure 2B*). In addition, transmission electron microscopy was used to observe a decrease in the number of autophagosomes in ubenimex-treated cells (P<0.001; *Figure 2C,2D*), which suggested that autophagy was reduced after ubenimex treatment. Together, these data indicated a critical role of autophagy in ubenimex-regulated 5-FU sensitivity in GC-resistant cells.

Ubenimex induces apoptosis mediated by chemotherapeutic drugs in SGC-7901/5-FU cells

Cells were treated with Annexin V-FITC/PI double staining to quantify the level of apoptosis. After pretreatment with 5-FU, ubenimex induced apoptosis of SGC-7901/5-FU cells in a dose-dependent manner. After 48 h of exposure to ubenimex, the increased dose (0.12 mg/mL) of ubenimex resulted in a more than fourfold increase in the proportion of total apoptotic cells compared to control cells, and the 2492



Figure 3 Ubenimex induces apoptosis mediated by chemotherapeutic drugs in SGC-7901/5-FU cells. (A) Flow cytometric analysis of Annexin V/PI double-stained SGC-7901/5-FU cells pre-treated for 24 h with 5-FU followed by the induction of ubenimex (0.06 or 0.12 mg/mL). Representative histograms are shown (left panels), as well as the means \pm SD of the proportions of apoptotic cells in three independent experiments (right panel). ****P<0.0001. (B) Western blotting of apoptosis associated protein expression in SGC-7901/5-FU cells after treatment for 48 h with 0.06 or 0.12 mg/mL ubenimex. GAPDH is shown as a loading control. Quantification was performed by a densitometry analysis and normalized to GAPDH levels. The bars indicate the means \pm SD from the 3 independent experiments (*P<0.05, **P<0.01, ***P<0.001, and ****P<0.0001). (C,D) Immunofluorescence staining of Bax and Bcl-2 with 0.06 or 0.12 mg/mL ubenimex treatment on SGC-7901/5-FU cells. 5-FU, 5-fluorouracil; Ube, ubenimex; SD, standard deviation.

majority of SGC-7901/5-FU cells were characterized as a late apoptotic population (*Figure 3A*). In addition, the expression of major apoptotic proteins namely, cleaved caspase-3, cleaved caspase-9, Bax and Bcl-2, was further examined by western blot analysis to explore the effect of ubenimex on apoptosis. As the concentration of ubenimex



Figure 4 Screening of target genes involved in the reversal of drug resistance by ubenimex. (A) Heat map was generated using R package to depict transcripts that were significantly differentially expressed in MKN-45/X cells after ubenimex treatment. The green and red colors indicate up-regulated and down-regulated transcripts, respectively. (B) Through HCS experiment, the RI value fold changes of MKN-45/X cells after overexpression or silencing of related genes were evaluated. (C) The column graph shows the changes in the RIs fold change after silencing of the 16 genes. **P<0.01 and ***P<0.001. (D) GO function and KEGG pathway enrichment analysis of the above genes. Ube, ubenimex; RI, resistance index; FDR, false discovery rate; GO, Gene Ontology; KEGG, Kyoto Encyclopedia of Genes and Genomes; HCS, high content screening.

increased, the expression of the Bcl-2 antiapoptotic protein decreased, while the levels of cleaved caspase-3, cleaved caspase-9 and Bax were significantly upregulated (*Figure 3B*), which further corroborated that ubenimex treatment induced apoptotic effects. The immunofluorescence results also showed that the expression of Bax was elevated and that the expression of Bcl-2 was decreased (*Figure 3C,3D*). These results revealed that ubenimex plays a key role in inducing apoptosis by 5-FU in GC-resistant cells.

Screening of target genes involved in the reversal of drug resistance by ubenimex

To obtain the target gene related to GC drug resistance,

we used microarray analysis to identify the gene expression profile of MKN-45/X cells after ubenimex treatment. Ubenimex caused the downregulation of 264 genes, including CD13, and the upregulation of 228 genes (*Figure 4A*). We evaluated the changes in RI values of MKN-45/X cells after overexpression or silencing of relevant genes by HCS experiments. We screened the top 16 genes, including CD13, that were related to the reversal of the resistance effect of ubenimex in GC (*Figure 4B,4C*). Through GO function and KEGG pathway enrichment analyses, we found that the above genes were all enriched in the NF- κ B pathway and involved in the regulation of cell autophagy and apoptosis (*Figure 4D*). After silencing the CD13 gene, we utilized HCS to observe its effect on the expression of

Xiu et al. Ubenimex reverses 5-FU resistance in GC cells



Figure 5 Ubenimex inhibits CD13/EMP3/FAK/NF-κB signalling pathway activity in SGC-7901/5-FU cells. (A) Associated protein expression was identified by western blot assay in SGC-7901/5-FU cells after CD13 was over-expressed. (B) Associated protein expression was identified by western blot assay in SGC-7901/5-FU cells after EMP3 was over-expressed. (C) Western blot analyses of associated protein expression in SGC-7901/5-FU cells treated with 0.06 or 0.12 mg/mL of ubenimex. (D) SGC-7901/5-FU cells were pretransfected with pEGFP-N1-CD13 plasmid for 24 h, and then stimulated with ubenimex (0.12 mg/mL) for another 24 h. Associated protein expression were detected by western blot assay. (E) SGC-7901/5-FU cells were pretransfected with pEGFP-N1-EMP3 plasmid for 24 h, and then stimulated with ubenimex (0.12 mg/mL) for another 24 h. Associated protein expression were detected by western blot assay. (F) Indicated cells were pretreated with pEGFP-N1-CD13 plasmid or pTZU-CD13-shRNA plasmid for 24 h, followed by treatment with ubenimex (0.12 mg/mL) for another 24 h, indicated associated channel protein expression in SGC-7901/5-FU cells. GAPDH is shown as a loading control. Quantification was performed by a densitometry analysis and normalized to GAPDH levels. The bars indicate the means ± SD from the 3 independent experiments (*P<0.05, **P<0.01, ***P<0.001, ***P<0.001, and *P>0.05). EMP3, epithelial membrane protein 3; FAK, focal adhesion kinase; NF-κB, nuclear factor-κB; EGFP, enhanced green fluorescent protein; Ube, ubenimex; 5-FU, 5-fluorouracil; SD, standard deviation.

other related genes and ranked them by the magnitude of the effect. We found that EMP3 may be a key molecule in the targeting of CD13 by ubenimex and reversing drug resistance in GC (Figure S2A). Similarly, we investigated the impact of silencing EMP3 on related genes and found that the mRNA level of PTK2 (the encoding gene of FAK) was downregulated (Figure S2B), indicating the presence of a potential CD13/EMP3/FAK/NF-KB regulatory axis.

Ubenimex inhibits CD13/EMP3/FAK/NF-κB signalling pathway activity in SGC-7901/5-FU cells

To validate the potential CD13/EMP3/FAK/NF- κ B regulatory axis identified by microarray analysis, GO functional analysis and KEGG pathway enrichment analysis, we explored the relationship between CD13 and EMP3 by western blot analysis after overexpression of these proteins. *Figure 5A* shows that the expression levels

2494

of EMP3, FAK and NF-KB were increased after CD13 overexpression. EMP3 overexpression had no significant effect on the expression of CD13 protein, but EMP3 overexpression significantly increased the expression levels of FAK and NF- κ B (*Figure 5B*). These results suggested that CD13 has a one-way regulatory effect on EMP3 expression. Furthermore, western blot assays confirmed that the expression of CD13, EMP3, FAK and NF-κB was downregulated in GC-resistant cells treated with ubenimex at different concentrations (Figure 5C). Interestingly, CD13 overexpression reversed the inhibitory effect of ubenimex on CD13, EMP3, FAK and NF-KB. However, the inhibitory effect of ubenimex on CD13 was not significantly affected after EMP3 overexpression, but EMP3 overexpression reversed the inhibitory effect of ubenimex on EMP3, FAK and NF- κ B (*Figure 5D*, 5E). Western blot analysis further confirmed that ubenimex affects the expression of downstream pathway proteins by targeting CD13/EMP3. Moreover, CD13 silencing and ubenimex treatment both attenuated the expression of EMP3, FAK and NF-KB, but the effect mediated by ubenimex was suppressed after CD13 overexpression (Figure 5F). Taken together, these findings showed that ubenimex inhibits EMP3/FAK/NFκB signalling pathway activity by targeting CD13 in SGC-7901/5-FU cells.

Ubenimex suppresses autophagy to promote 5-FU-induced apoptosis by inhibiting the activation of the CD13/EMP3/ FAK/NF- κ B pathway

Accumulating evidence suggests that the expression and phosphorylation of FAK play important roles in the initiation and transmission of autophagy signals as well as the induction of apoptosis tolerance in GC cells (22). Immunofluorescence staining of SGC-7901/5-FU cells was performed to test whether ubenimex affects the phosphorylation of FAK by targeting CD13. In both the ubenimex-treated and CD13-deficient groups, the phosphorylation of FAK was significantly downregulated. Notably, the inhibition by ubenimex was attenuated upon CD13 overexpression (Figure 6A). We next investigated whether ubenimex attenuates autophagy in GC-resistant cells by modulating the CD13/EMP3/FAK/NF-KB pathway. Figure 6B shows that there was a significant increase in SQSTM1 expression but a significant reduction in LC3B, Beclin-1 and ATG5 expression in the ubenimex-treated 5-FU-resistant GC cells compared to the control cells, which indicated the downregulation of autophagy. As expected, the results in

the CD13-deficient group were consistent with this finding. More importantly, CD13 overexpression reversed the inhibition of autophagy by ubenimex. We completed further validation by detecting LC3B by immunofluorescence (Figure 6C). Together, these results suggested a critical role of autophagy in ubenimex-increased 5-FU sensitivity in GC-resistant cells. It is well known that dysregulated expression of the Bcl-2 family and subsequent inactivation of caspase-3 or caspase-9 are fundamental mechanisms for apoptosis resistance in GC cells (28,29). In the present study, ubenimex increased the expression of Bax, but it downregulated the expression of Bcl-2. Similar to the CD13-deficient group, ubenimex also significantly elevated the expression of the cleaved forms of PARP, caspase-3, and caspase-9, but this effect was attenuated by CD13 overexpression (Figure 6D). In addition, we used Annexin V/PI staining to assess the effects of CD13 silencing, CD13 overexpression and ubenimex treatment on cell apoptosis rates after 5-FU treatment. Ubenimex enhanced the 5-FUinduced apoptosis in GC-resistant cells, while the effect was attenuated by upregulating CD13, which indicated a role for CD13 in 5-FU-induced apoptosis (Figure 6E). Collectively, these results suggested that ubenimex inhibited autophagy to promote 5-FU induced apoptosis, in which the activity of the CD13/EMP3/FAK/NF-KB pathway is alleviated.

Ubenimex renders GC-resistant cells sensitive to 5-FU by alleviating the activity of the CD13/EMP3/FAK/NF- κ B pathway

To determine whether ubenimex reverses drug resistance by targeting CD13 and regulating the EMP3/FAK/NFκB pathway to enhance cellular sensitivity to 5-FU, we downregulated CD13 expression in SGC-7901/5-FU cells. Control cells, ubenimex-treated cells and CD13-deficient cells were subjected to the CCK-8 cell viability assay after pretreatment with different concentrations of 5-FU. Compared to control cells, GC-resistant cells showed increased 5-FU sensitivity after knockdown of CD13 or treated with ubenimex (P<0.05; Figure 7A). However, there was no significant difference between the CD13-deficient and ubenimex-treated groups. In addition, the 5-FU pretreatment group was also tested in the same manner to corroborate whether CD13 knockdown has the same effect on ubenimexenhanced 5-FU sensitivity and correlates with treatment time. All three groups were treated with 5-FU (12 µg/mL) for 0-72 hours (P<0.05; Figure 7B), and the results indicated that ubenimex enhanced the chemosensitivity of SGC7901/5-



Figure 6 Ubenimex suppresses autophagy to promote 5-FU-induced apoptosis by inhibiting the activation of the CD13/EMP3/FAK/NF-κB pathway. (A) Immunofluorescence staining of p-FAK with 0.12 mg/mL ubenimex treatment or pretransfected with pEGFP-N1-CD13 or pTZU-CD13-shRNA on SGC-7901/5-FU cells. (B) Western blotting analysis of autophagy related proteins with 0.12 mg/mL ubenimex treatment or pretransfected with pEGFP-N1-CD13 or pTZU-CD13-shRNA after 5-FU pretreated on SGC-7901/5-FU cells. GAPDH is shown as a loading control. Quantification was performed by a densitometry analysis and normalized to GAPDH levels. The bars indicate the means ± SD from the 3 independent experiments (*P<0.05, **P<0.01, ***P<0.001, and [#]P>0.05). (C) Immunofluorescence staining of LC3B with 0.12 mg/mL ubenimex treatment or pretransfected with pEGFP-N1-CD13 or pTZU-CD13-shRNA after 5-FU pretreated on SGC-7901/5-FU cells. (D) Western blotting analysis of apoptosis related proteins with 0.12 mg/mL ubenimex treatment or pretransfected with pEGFP-N1-CD13 or pTZU-CD13-shRNA after 5-FU pretreated on SGC-7901/5-FU cells. (D) Western blotting analysis of apoptosis related proteins with 0.12 mg/mL ubenimex treatment or pretransfected with pEGFP-N1-CD13 or pTZU-CD13-shRNA after 5-FU pretreated on SGC-7901/5-FU cells. (D) Western blotting analysis of apoptosis related proteins with 0.12 mg/mL ubenimex treatment or pretransfected with pEGFP-N1-CD13 or pTZU-CD13-shRNA after 5-FU pretreated on SGC-7901/5-FU cells. (D) Western analysis and normalized to GAPDH levels. The bars indicate the means ± SD from the 3 independent experiments (*P<0.05, **P<0.01, ***P<0.01, and [#]P>0.05). (E) Flow cytometric analysis of Annexin V/PI double-stained SGC-7901/5-FU cells with 0.12 mg/mL ubenimex treatment or pretransfected with pEGFP-N1-CD13 or pTZU-CD13-shRNA after 5-FU pretreated. Representative histograms are shown (left panels), as well as the means ± SD of the proportions of apoptotic cells in three independent experiments (right panel). *P<0.05, **P<0.01,



Figure 7 Ubenimex renders GC-resistant cells sensitive to 5-FU by alleviating the activity of the CD13/EMP3/FAK/NF- κ B pathway. (A) SGC7901/5-FU cells were pretreated with ubenimex (0.12 mg/mL) and pTZU-CD13-shRNA both for 24 h, followed by incubation with 5-FU at increasing concentrations (0, 0.5, 1, 2, 4, 16, 32 µg/mL) for another 48 h. (B) SGC7901/5-FU cells were pretreated with ubenimex (0.12 mg/mL) and pTZU-CD13-shRNA both for 24 h, followed by incubation with 5-FU at increasing concentrations (0, 0.5, 1, 2, 4, 16, 32 µg/mL) for another 48 h. (B) SGC7901/5-FU cells were pretreated with ubenimex (0.12 mg/mL) and pTZU-CD13-shRNA both for 24 h, followed by incubation with 5-FU (12 µg/mL) for 0–72 hours. Cell viability was determined by the CCK-8 method. The results are expressed as the means \pm SD of three replicates. *P<0.05 and **P<0.01. OD, optical density; 5-FU, 5-fluorouracil; Ube, ubenimex; EMP3, epithelial membrane protein 3; FAK, focal adhesion kinase; NF- κ B, nuclear factor- κ B; CCK-8, Cell Counting Kit-8; SD, standard deviation.

FU cells and that CD13 is an essential key target, which was consistent with previous findings.

Discussion

GC is identified as the second leading cause of cancer incidence and mortality, resulting in a serious threat to public health (30). 5-FU is one of the most widely used chemotherapeutic agents for the treatment of GC, which inhibits tumour proliferation and DNA replication by inhibiting thymidylate synthase to synthesize thymine, ultimately inducing apoptosis (31). However, the response rate to 5-FU chemotherapy in patients with advanced GC is less than 32% not satisfied (32). The development of drug resistance in cancer is a complex process in which multiple factors participate, including drug resistanceassociated proteins, dysfunction of DNA damage repair, cell autophagy, decreased drug accumulation, metabolic detoxification, alterations in drug targets and signal transduction molecules. This is the first study to suggest that role of CD13-induced autophagy in the 5-FU resistance of GC cells is to protect tumour cells against apoptosis and that autophagy induced by 5-FU is a protective mechanism conducive to cell survival. Consistently, we also found that ubenimex, a CD13 inhibitor, reverses 5-FU resistance and significantly increases the sensitivity of SGC-7901/5-FU cells to 5-FU in a dose- and time-dependent manner.

As recognized, FAK expression and phosphorylation activity are important for the initiation and transmission of autophagy signals as well as the induction of apoptosis tolerance in GC cells, which can affect the expression of NF- κ B, whereas the NF- κ B, p65 and E2F transcription factors upregulate BECN1 gene expression and increase the level of autophagy (23). As one of the key molecules in autophagosome nucleation, Beclin-1 is an important target in the regulation of autophagy. ATG5 is one of the critical regulators of autophagic cell death and is a protective molecule in tumour cells during the course of chemotherapy. Upregulated expression of ATG5 in GC tissues is associated with chemoresistance in GC and has an important protective role in autophagy (33). p62, another important autophagy-related protein, is an autophagy substrate that is degraded in autophagosomes, and it is considered an indicator of autophagic flux (34). It has been shown that high p62 expression is closely related to poor prognosis in oral squamous cell carcinoma and triplenegative breast cancer but not in colon and breast cancers (35).

Apoptosis refers to the process of genetically controlled, autonomous and orderly death of multicellular organisms under physiological or pathological changes in which caspase-dependent apoptotic pathway is well documented. In particular, the caspase family-mediated proteasome cleavage cascade, ultimately leading to apoptosis. Evidence has shown that the signals of each apoptotic pathway eventually converge to caspase-3 to execute apoptosis, and the activation of caspase-3 may destroy its substrate, PARP, leading to the loss of DNA repair function and cell apoptosis (36). The Bcl-2 gene family is one of the main regulators of the mitochondrial pathway, and its family members are divided into two main categories as follows: anti-apoptotic proteins (such as Bcl-2) and pro-apoptotic proteins (such as Bax). Both Bcl-2 and Bax interact with each other on mitochondria, and their regulation of apoptosis is not only dependent on the expression level of the gene but also related to the ratio between the two.

Apoptosis and autophagy are two important cellular biological behaviours that maintain the stability of the internal environment, and they exist in the physiological process of normal cells and are also an important defence mechanism for cells under injury. However, once a tumour is formed or when the external environment disturbs cellular regulation, autophagy provides abundant nutrition and energy for cancer cells to promote cell survival. Recent evidence has suggested that autophagy plays a dual role in promoting or inhibiting apoptosis. Autophagy mainly plays an anti-apoptotic role by degrading apoptosis-related proteins, which contribute to the occurrence of acquired drug resistance. As a caspase substrate, Beclin-1 is an important molecule in autophagy because it can be cleaved by a caspase. After cleavage, Beclin-1 loses autophagy function but promotes cell apoptosis (37). As an important tumour suppressor protein, Beclin-1 not only interacts with the apoptosis-related genes, Bcl-2 and caspase-9, but also participates in the formation of the autophagy bilayer membrane and jointly participates in the regulation of apoptosis and autophagy (38).

Here, the genes related to the ubenimex-induced reversal of GC resistance were screened using microarray analysis and HCS. EMP3 and FAK were identified to be the key molecules involved in the ubenimex-induced reversal of GC resistance by targeting CD13, and a positive correlation between FAK and EMP3 expression was also found in GC patient tissues. However, studies on the relationship between EMP3 and 5-FU-resistant GC cells as well as the key mechanism by which EMP3 regulates FAK activity are limited. Previous evidence has demonstrated that the aberrant expression of EMP3 in GC cells is partially attributed to hypermethylation impairment of its silencing mechanism (39). Additionally, CD13 may be the initiator of harm in this process by further regulating NF-KB expression, nuclear import and its transcriptional activity by affecting FAK expression and phosphorylation activity. The possibility of a CD13/EMP3/FAK/NF-KB regulatory axis was also further supported by western blot analysis in the present study using CD13 and EMP3 overexpression and silencing vectors, which showed that ubenimex increased its inhibitory effect on pathway activity in a dosedependent manner. Therefore, these findings suggested that

ubenimex enhances 5-FU sensitivity and thus reverses drug resistance by inhibiting the activation of the EMP3/FAK/ NF- κ B pathway in GC cells by targeting CD13. To the best of our knowledge, this is the first study to investigate the underlying molecular mechanisms of ubenimexinduced chemosensitivity of GC cells to 5-FU via CD13/ EMP3/FAK/NF- κ B-mediated transcriptional activation. Furthermore, the changes in the ATG5, Beclin-1, LC3B and p62 autophagy-related proteins after treatment with ubenimex, CD13 overexpression and CD13 silencing were determined using western blot and immunofluorescence analyses. Finally, the number of autophagosomes was examined by transmission electron microscopy. These results demonstrated that ubenimex suppresses the occurrence of autophagy in GC drug-resistant cells by targeting CD13.

Consistently, the apoptosis of GC drug-resistant cells was detected by Annexin V/PI double staining, and flow cytometry analysis showed that SGC-7901/5-FU cells treated with ubenimex had increased apoptosis compared to control cells, and the effect of apoptosis increased with the concentration of ubenimex. Western blot and immunofluorescence analyses also showed increased expression of Bax, cleaved PARP, cleaved caspase-3 and cleaved caspase-9 as well as decreased expression of Bcl-2, indicating a mixed phenotype of autophagic cell death and apoptosis by targeting CD13 in response to treatment with ubenimex in SGC-7901/5-FU cells. Importantly, we found that ubenimex significantly attenuated the autophagic properties of 5-FU in GC-resistant cells. Therefore, we hypothesized that ubenimex may be a reliable autophagy inhibitor and promote 5-FU-induced apoptosis.

The present study showed that FAK expression and phosphorylation play a key role in regulating autophagy and apoptosis. In the present study, immunofluorescence detection of p-FAK expression indicated that FAK phosphorylation was significantly decreased after ubenimex treatment and CD13 silencing, but these effects were reversed after CD13 overexpression. These findings indirectly demonstrated that ubenimex inhibits autophagy by regulating the EMP3/FAK/NF-κB pathway by targeting CD13 to promote chemotherapeutic drug-induced apoptosis and reverse chemoresistance. However, the detailed mechanism by which ubenimex induces apoptosis by inhibiting autophagy needs to be further studied.

Conclusions

Our findings demonstrated that autophagy may be

Translational Cancer Research, Vol 11, No 8 August 2022

suppressed, at least in part, by interfering with the CD13/ EMP3/FAK/NF- κ B axis, which attributes to the poor response to 5-FU-based chemotherapy in patients. Overall, the present study suggested that the combination of 5-FU and ubenimex may represent an innovative therapeutic strategy for GC patients and provide attractive insights regarding the combined use of Ubenimex as a GC chemotherapeutic intervention.

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Footnote

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Conflicts of Interest: All authors have completed the ICMJE uniform disclosure form (available at https://tcr.amegroups.com/article/view/10.21037/tcr-22-345/coif). The authors have no conflicts of interest to declare.

Ethical Statement: The authors are accountable for all aspects of the work in ensuring that questions related to the accuracy or integrity of any part of the work are appropriately investigated and resolved.

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Appendix 1

Methods

Gene chip analysis and HCS of cell function

The Affymetrix ClariomTM S gene chip was used to determine the gene expression spectrum of MKN-45/X cells before and after Ubenimex treatment and assess changes in gene expression. In this study, the fold change of RI values of MKN-45/X cells was evaluated by the HCS, from which genes with the most significant differential fold changes were screened out. The effect of silencing CD13 expression on the mRNA level of the above genes was observed by HCS experiment, and the key genes downstream of CD13 were selected.

Plasmid construction and gene amplification

EGFP-labeled ANPEP (CD13; GenBank number NM_001150), EMP3 (GenBank number NM_001425) overexpression plasmids and negative control plasmids (empty vector control plasmid; CMV-MCS-3FLAG-IRES-EGFP-SV40-Neomycin) were used in this study. The following primer pairs were designed based on the ANPEP and NAB1 mRNA sequences: ANPEP-p1, 5'-TACCGG ACTCAGATCTCGAGCGCCACCATGGCCAAGGGCCTTCTATATTTCCAAG-3'; ANPEP-p2, 5'-TCCTTGTAGTCCATGGATCCTTTGCTGTTTTCTGTGAACCACTGGAGCAC-3'; EMP3-p1, 5'-GAGGATCCCGGGGTACCGGGTCGCCACCATGGCGGAGCCGAGCGGGC-3'; EMP3-p1, 5'-TCACCATGGTG GCGACCGGGGCTGACACTCAACTGAGCA-3'. The PCR mixture contained 2 µL of cDNA template, 1 µL of primer (10 µmol/L), 10 µL of PCR mix and dH₂O to a total volume of 20 µL. The PCR program was as follows: 94 °C for 5 min; 40 cycles of 94 °C for 30 s, 58 °C for 30 s and 72 °C for 5 min; and a final extension at 72 °C for 5 min. The final PCR products were separated by electrophoresis using 1% polyacrylamide gels, and the target fragment was purified and recovered using an agarose gel extraction kit (Watson Biomedical Inc., Shanghai, China).

Proliferative activity and cell sensitivity assays

Human GC parental cell line and 5-FU resistant cell were dispensed into 96-well culture plates (5×10^3 cells/well) and incubated at 37 °C for 0–48 h to evaluate proliferative activity. In addition to evaluate the resistant cell sensitivity to 5-FU, SGC-7901/5-FU cells were incubated with various concentrations of 5-FU (0.5, 1, 2, 4, 16, 32 µg/mL) for 48 h, in the presence or absence of ubenimex (0.12 mg/mL); or were incubated with 5-FU (12 µg/mL) for 0–72 hours in the presence or absence of ubenimex (0.12 mg/mL). Finally, 10 µL CCK-8 solution was added into each well. the cells were then incubated for another 4 h, and the absorbance at 450 nm was read by a microplate reader (EL340; Bio-Tek Instruments, Hopkinton, MA, USA). The resulting absorbance is used to generate the growth curve to assess the proliferative activity or cell sensitivity.

Western blotting

Before lysis, cells were incubated with different treatments to determine the expression level of protein, then, cold PBS washed the cells and proteins were extracted from the cells by resuspension in cold radioimmunoprecipitation assay (RIPA) buffer containing 1% phenylmethylsulfonyl fluoride (PMSF) and 1% phosphatase inhibitor for 15 min. The samples were centrifuged at 12,000 rpm for another 15 min at 4 °C, and the supernatants were recovered for continued analysis. The bicinchoninic acid (BCA) protein assay kit (Solarbio, Beijing, China) was used for the measurement of protein concentration. Subsequently, SDS-PAGE was used to separate proteins and which were transferred to polyvinylidene difluoride (PVDF) membranes. After incubating with different primary antibodies overnight at 4 °C, the membranes incubated for another 1.5 h at room temperature with horseradish peroxidase (HRP)-conjugated secondary antibodies. Subsequently, immunoblots were visualized using enhanced chemiluminescence (LAS-4000), the experiments were repeated at least three times.



Figure S1 Establishment of 5-FU resistant GC cells. (A) SGC7901/5-FU cells and their parental cells. (B) SGC7901/5-FU cells and their parental cells were treated with increasing concentrations of 5-FU (0.5, 1, 2, 4, 16, 32 µg/mL) for 48 h. Cell viability was determined by the CCK-8 method. The results are expressed as the means \pm SD of three replicates. **P<0.01. (C) Western blotting analysis of CD13 in parental and SGC-7901/5-FU cells. GAPDH is shown as a loading control. Quantification was performed by a densitometry analysis and normalized to GAPDH levels. The bars indicate the means \pm SD from the 3 independent experiments (*P<0.05). 5-FU, 5-fluorouracil; RI, resistance index; GC, gastric cancer; CCK-8, Cell Counting Kit-8; SD, standard deviation.

A	Gene Symbol	mRNA Fold Change(After CD13 down-regulation	Regulation	В	Gene Symbol	mRNA Fold Change(After EMP3 down-regulation	Regulation
	EMP3	8.699537426	Down		PTK2	8.408637398	Down
	PTK2	8.408637398	Down		RELA	7.714647979	Down
	RELA	7.714647979	Down		BECN1	7.047199391	UP
	BAK1	7.613932323	Down	-	BCL2L1	6.666254896	Down
	BECN1	7.032987828	UP		BCL-2	6.658808486	Down
	BCL2L1	7.025541418	Down		BAK1	6.613932323	Down
	BCL-2	7.023929478	Down	Fold change	BCL2L4	6,500690446	Down
	BCL2L4	6.824932378	Down	rold change	DIABLO	6 301693346	LIP
	DIABLO	6.808841425	UP		CASP3	6 285602303	Doum
	CASP3	6.799591928	Down		CASES	0.285002595	Down
	APAF1	6.762938928	UP	ŧ	APAFI	5.776352896	UP
	ATG12	6.389432328	Down		ATG12	5.739699896	Down
	ATG5	5.906001318	UP		ATG5	5.366193296	UP
	MAP1LC3	5.785109163	Down		MAP1LC3	4.882762286	Down
	HMGB1	5.619668501	Down		HMGB1	4.761870131	Down

Figure S2 The effect of *CD13* silence and *EMP3* silence on the expression of indicated genes. (A) The *CD13* gene was silenced to observe its effect on the expression of other related genes and ranked. (B) The *EMP3* gene was silenced to observe its effect on the expression of other related genes and ranked. EMP3, epithelial membrane protein 3.