

# Effects of *Helicobacter pylori* infection and interleukin 6 on the expression of ITIH4 in human gastric cancer cells

### Dadao Jing, Jie Jin, Zhu Mei, Qingqing Zhu, Yingying Lu, Xingpeng Wang

Department of Gastroenterology, Shanghai General Hospital, Shanghai Jiao Tong University School of Medicine, Shanghai, China *Contributions:* (I) Conception and design: D Jing, X Wang; (II) Administrative Support: D Jing; (III) Provision of study materials or patients: J Jin, Z Mei; (IV) Collection and assembly of data: Q Zhu, Y Lu; (V) Data analysis and interpretation: D Jing, Z Mei, X Wang; (VI) Manuscript writing: all authors; (VII) Final approval of manuscript: All authors.

*Correspondence to:* Dadao Jing; Xingpeng Wang. Department of Gastroenterology, Shanghai General Hospital, Shanghai Jiao Tong University School of Medicine, 650 Xinsongjiang Rd, Songjiang, Shanghai, China. Email: dadaojing@126.com; richardwangxp@163.com.

**Background:** Human inter-α-trypsin inhibitor heavy chain 4 (ITIH4) is an acute phase response protein that is positively regulated by interleukin-6 (IL-6), suggesting that ITIH4 can be used as an early serological biomarker of gastric cancer.

**Methods:** The mechanism of ITIH4 elevation in gastric cancer cells was investigated via *Helicobacter pylori* (*H. pylori*) with or without recombinant human interleukin-6 (rhIL-6) in human gastric cell line AGS cells. The messenger RNA (mRNA) expressions of ITIH4 and IL-6 in AGS cells were detected with real-time quantitative polymerase chain reaction (qPCR), while the expression of ITIH4 protein and IL-6 in AGS cells was detected with Western blot and enzyme-linked immunosorbent assay (ELISA) methods.

**Results:** We found that the mRNA and protein expressions of ITIH4 started to increase synchronously at 3 hours (h), peaked at 12 h, and then declined to normal a level at 24 h in AGS cells after *H. pylori* infection. Interestingly, the expression patterns of ITIH4 were positively correlated to IL-6 expression levels. In the ITIH4 expression experiment with IL-6 administration, the expression level of ITIH4 began to increase at a concentration of 6.25 ng/mL with rhIL-6 administration, reach a peak, and plateau at a concentration of 25 ng/mL of rhIL-6. The expression patterns of ITIH4 in AGS cells induced by *H. pylori* infection were closely related to the infection of *H. pylori* and the level of IL-6.

**Conclusions:** The detection of ITIH4 and IL-6 during *H. pylori* infection may be useful for the screening of early gastric cancer.

**Keywords:** Gastric cancer; *Helicobacter pylori* (*H. pylori*); human inter-α-trypsin inhibitor heavy chain 4 (ITIH4); interleukin-6 (IL-6)

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#### Introduction

Human inter- $\alpha$ -trypsin inhibitor heavy chain 4 (ITIH4) is a 120 kDa serum glycoprotein mainly secreted by the liver and is extensively involved in liver development and regeneration (1,2). One study showed that ITIH4 circulates in the blood at a concentration of about 100 ng/mL (3). It is an acute phase responsive protein and is positively regulated by interleukin 6 (IL-6) (4,5). Previous studies have shown that the expression of ITIH4 in cancer patients is related to tumorigenesis and cancer progression, suggesting that ITIH4 may be used as a serological marker for diagnosis cancer (6-8). Our cytological studies have also shown that 70 kDa ITIH4 protein fragments are expressed in gastric cancer cells. In addition, our study also found that elevated serum ITIH4 in patients with early gastric cancer was strongly associated with *Helicobacter pylori* (*H. pylori*) infection and serum IL-6 levels. So far, however, the effects of ITIH4 on the occurrence and development of gastric cancer, and the mechanism of its elevation in early gastric cancer serum, remain unclear. A large number of studies have demonstrated that *H. pylori* infection can cause chronic inflammation, atrophy, intestinalization, dysplasia and metaplasia of gastric mucosal epithelial cells, and canceration (9,10). Other studies have shown that IL-6 receptor expression is closely related to the emergence and development of gastric cancer (11).

To evaluate the exact functions of IHIT4 expression in gastric cancer, we treated human AGS gastric cancer cells with recombinant human interleukin 6 (rhIL-6) and *H. pylori*, and observed the effects of IL-6 and *H. pylori* on ITIH4 expression in gastric cancer cells. The results yielded from this experiment elucidated the mechanism of ITIH4 elevation in early gastric cancer serum and could be valuable in further clarifying the possibility of ITIH4 as a serological marker of early gastric cancer.

#### Methods

#### Cell culture

Human gastric cancer cell line AGS (Shanghai institute of cell research, Chinese academy of sciences) was selected and cultured in RPMI 1640 medium (Gibco, USA) containing 10% newborn bovine serum (Hyclone, USA), 100 unit (U)/mL penicillin, and 100 U/mL streptomycin, and placed in 5% CO<sub>2</sub> cell incubator at 37 °C. When the cells adhered to the 80–90% confluence, they were digested and subcultured with 0.25% trypsin solution detachment. This study was approved by ethical committee of Shanghai General Hospital (Approval#: 2017KY210) and complied with the Declaration of Helsinki (As revised in 2013). Informed consent was not required because we only used human cancer cell line in the entire experiment.

#### H. pylori strain culturing

*H. pylori* bacteria were resuscitated and inoculated into brain heart infusion containing 7% defibrinated sheep blood, 0.4% IsoVitaleXEnrichment (Becton, Dickinson, and Co., Franklin Lakes, NJ, USA), 0.08% amphotericin B, 0.2% vancomycin, and 0.5% sulfa synergist infusion agar medium (USA, Becton, Dickinson, and Co., Franklin Lakes, NJ, USA). This was cultured at 37 °C in a three-gas incubator containing 5% O<sub>2</sub>, 10% CO<sub>2</sub>, 85% N<sub>2</sub>, and 95% relative humidity. After 3–5 days, the typical colonies were needle-point-like (diameter 1–2 days, the typical colonies were needle-point-like (diameter 1nergihad increased, the colonies fused on the surface of the culture medium to form a translucent bacterial moss. The motor status of H. *pylori* was observed under a dark-field microscope. The morphological identification of H. *pylori* strains was verified by Gram staining, Giemsa staining, and Warthin-Starrysilver staining.

#### Co-culture of H. pylori and AGS cells

To evaluate IHIT4 gene and protein expression in H. pylori-infected AGS cells, we performed the following steps for AGS cells culture. First, AGS cells were taken in logarithmic growth phase, routinely digested with sterile 0.25% trypsin, and were counted. Cells were adjusted with antibiotic-free RPMI 1640 medium with 10% fetal bovine serum (FBS) at 5bov<sup>5</sup> cells density/ml and inoculated into a 12-well cell culture plate at 200 µL per well. Then, each well was supplemented with 0.8 mL RPMI 1640 culture medium with 10% FBS, and the final volume of each well was 1 ml. These cells were placed in a 5% CO<sub>2</sub> incubator at 37 °C to culture for 24 hours (h). Second, 4-5 colonies with a diameter of about 1 mm that had been cultured for 5-7 days on the brain heart infusion agar medium with an inoculating loop and saline were sterilized with 3-5 mL of antibiotic-free 10% FBS RPMll640 medium; this adjusted the bacterial concentration to  $1 \times 10^8$  colony-forming units (CFU)/mL. Third, the cell culture plate was taken out for 2 h, the culture supernatant was removed, and then divided into five groups: A, B, C, D, and E with 6 duplicate wells in each group. Separately, groups A, B, C, D, and E were combined with 100 L of bacterial solution, AGS cells, and H. pylori that were incubated at a ratio of 1:100 multiplicity of infection (MOI). The AGS cells served as the control group. Then, 1 ml of fresh, serum-free, antibiotic-free RPMI1640 culture solution was added into each well and cultured in a 5% CO<sub>2</sub> incubator at 37 °C for 1, 3, 6, 12, and 24 h for groups A, B, C, D, and E, respectively. The culture conditions are described in Table 1. Cells were harvested for ITIH4 gene and protein analysis according to the set time.

#### rhIL-6-induced ITIH4 expression

To assess the effects of IL-6 on ITIH4 gene expression in AGS cells, we harvested AGS cells in logarithmic growth phase with sterile 0.25% trypsin digestion. Then, cells were counted and cell density was adjusted at  $5 \times 10^5$  cells/mL

Variable	А	В	С	D	Е	Control			
AGS cell	1×10 <sup>5</sup>	1×10⁵	1×10 <sup>5</sup>	1×10 <sup>5</sup>	1×10⁵	1×10 <sup>5</sup>			
H. pylori (CFU/mL)	1×10 <sup>7</sup>	0							
Total volume (mL)	1.0	1.0	1.0	1.0	1.0	1.0			
Time of culture (hour)	1	3	6	12	24				

Table 1 Co-culture of H. pylori and AGS cell lines

H. pylori, Helicobacter pylori; CFU, colony-forming-unit.

Table 2 Recombinant human interleukin 6 induces ITIH4 expression

Variable	А	B	С	D	E
AGS cell	1×10 <sup>5</sup>	1×10 <sup>5</sup>	1×10⁵	1×10 <sup>5</sup>	1×10 <sup>5</sup>
rhIL-6 (ng/mL)	0	6.25	12.5	25.0	50.0
Total volume (mL)	1.0	1.0	1.0	1.0	1.0
Time of culture (hour)	24	24	24	24	24

rhIL-6, recombinant human interleukin 6; ITIH4, inter-α-trypsin inhibitor heavy chain 4.

with antibiotic-free 10% FBS RPMI 1640 medium. The cells were inoculated into a 24-well cell culture plate, 200 in AGS cells, we harvested AGS cells in logarithmic growth phase cultured in RPMI 1640 medium (Gibco, USA) which was then incubated in a 5% CO<sub>2</sub> incubator at 37 °C for 24 h. Previous cells were treated with 6.25, 12.5, 25.5, and 50.0 ng/mL rhIL-6 (PeproTech EC, UK) and antibiotic-free RPMI 1640 cell culture medium, respectively (*Table 2*). After the cells were incubated for 24 h at 37 °C, 5% CO<sub>2</sub> incubator and were digested, the cells with 0.25% sterile trypsin, RNA and protein were extracted and stored in –70 °C low temperature refrigerator for later use.

### Determination of IL-6 and ITIH4 mRNA expression in cells by real-time quantitative polymerase chain reaction (qPCR)

### qPCR

Total RNA was isolated from AGS cells with *H. pylori* and IL-6 administration using TRIzol and was then used for complementary DNA synthesis using random primers. Reverse transcriptasequantitative polymerase chain reaction (PCR) was performed following standard procedures. PCR conditions for ITIH4 were 35 cycles of 95 °C for 15 seconds (s); 58 °C for 15 seconds; and 72 °C for 10 seconds, conditions for IL-6 were 33 cycles of 95 °C for 30 seconds; 56 °C for 30 seconds; and 72 °C for 30 seconds, and conditions for the house keeping gene

GAPDH were 24 cycles of 95 °C for 30 seconds; 58 °C for 30 seconds; and 72 °C for 30 seconds. The primer pairs of the expected products are displayed in Table 3. For each specimen in duplicate wells, a sample without cDNA was set as a negative control, and a blank buffer without cDNA and no probe was used as a blank control. The ABI7900 fluorescence quantitative PCR instrument collected the genes to be tested, and the internal reference GAPDH was used to amplify the fluorescence signals of each cycle, using SDS2.0 software (Applied Biosystems, USA) for fluorescence collection and data analysis. The detection mode was absolute quantification. Sequence Detector 1.7 software analysed the cycle threshold (Ct) value and Ct threshold value (threshold value = standard deviation of baseline signal ×10). The cDNA t value reflects the number of reaction cycles required when the template is amplified to a certain number of copies (in an exponentially rising period): the larger the Ct value, the smaller the amount of starting template participating in the reaction, and vice versa. Additional calculation included  $\Delta\Delta$ Ct and relative quantity (RQ) value. This value was arrived at by calculating the average Ct value of each specimen, and taking the target gene Ct value minus GAPDH Ct value as the d t value as follows:  $\Delta\Delta Ct = \Delta Ct$  (experimental group) –  $\Delta Ct$  (control group); RQ =  $2^{-\Delta\Delta Ct}$ . Amplified products were resolved by 2-3% agarose gel electrophoresis and 100 V electrophoresis for 45 min; the gel was observed under UV light, and a photo of the bottom was taken.

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Table 9 Tolymerase chain reaction primer sequence of this experiment						
Objects	Primer sequences	Amplified fragment (bp)	Annealing temperature (°C)			
ITIH4	Forward 5'-GGTCTCATCCCGATTTGC-3', reverse 5'-TGATCCCTGGGTAGGTCAT-3'	155	58			
IL-6	Forward 5'-GACAGCCACTCACCTCTTCA-3', reverse 5'-CGCAGAATGAGATGAGTTGT-3'	457	56			
GAPDH	Forward 5'-AACGGATTTGGTCGTATTG-3', reverse 5'-GGAAGATGGTGATGGGATT-3'	208	58			

Table 3 Polymerase chain reaction primer sequence of this experiment

ITIH4, inter-α-trypsin inhibitor heavy chain 4; IL-6, interleukin-6; GAPDH, glyceraldehyde 3-phosphate dehydrogenase.

# Western blot method to detect the expression of ITIH4 and IL-6 protein in gastric cancer cells

Cell lysates were prepared, and protein concentration measured by bicinchoninic acid (BCA) protein assay reagent. An equal amount of protein (20 ng) were loaded on a 6% or 10% SDS-polyacrylamide gel and transferred to a polyvinylidene fluoride (PVDF) membrane. The transferred protein was blocked in 5% fat-free dry milk in phosphate buffered saline (PBS) containing 0.1% Tween-20 (PBST) for 1 h at room temperature. Then, the membrane was incubated with a primary antibody (rabbit anti-human ITIH4 antibody, Abcam, Cat#ab180139, USA; rabbit anti-human IL-6 antibody, Abcam, Cat# 1457-1, RRID:AB\_562150, USA) in 5% skimmed milk powder in 3% PBS at 4 °C overnight and incubated with the respective horseradish peroxidase (HRP)-conjugated secondary antibodies (goat anti-rabbit IgG-HRP, Cat#G-21234 Invitrogen, USA) at 1:2,000 for 2 h and washed again with PBST. An enhanced chemiluminescence (ECL) detection kit (Shanghai Shengsheng Biotechnology Co., Ltd.) was used to visualize the expressed proteins. The Gene Genius imaging system was used for scanning and imaging, and each procedure was repeated at least three times.

#### Expression levels of ITIH4 and IL-6 in cell lysate

ITIH4 and IL-6 levels in cell lysate were measured by enzyme-linked immunosorbent assay (ELISA) according to the manufacturer's protocol (Wuhan Huamei Biological Engineering Co., Ltd. and R & D).

#### Image analysis

Gel-Pro image analysis software was used to analyse the Western blot results. The detection results were described in terms of relative absorbance. The relative absorbance was the ratio of the grey level of the target band to the grey level of the internal reference GAPDH band.

#### Statistical analysis

The obtained data are expressed as mean  $\pm$  SD. The relevant data were analysed by SPSS 24.0 statistical software. The data between the two groups were analyzed by paired *t*-test. The data between multiple groups were analyzed by a single factor analysis of variance. A P value <0.05 was considered statistically significant.

#### **Results**

### ITIH4 mRNA expression of AGS cells induced by H. pylori infection (amplification curve of ITIH4 gene by real-time qPCR)

After harvesting H. pylori-infected AGS cells at the set time, RNA was isolated and extracted, and real-time quantitative PCR was performed to analyze the differential expression of ITIH4 mRNA in AGS cells. The phase of the fluorescence intensity detected by the PCR instrument was uniformly set to the annealing period of the amplification reaction. In this experiment, ITIH4 and gene amplification were performed at 60 °C with 30 seconds as the annealing period. After 35 cycles of amplification reactions had elapsed, the system analyzed the fluorescence intensity growth index (DRn) of each reaction tube collected during each cycle of reaction and drew the amplification curve of each reaction tube (Figure 1A,B,C). The amplification curve of each sample had reached the plateau stage. In contrast, none of the melting curves detected primer dimers and other non-specific fluorescent signals (Figure 1D,E,F).

# Differential expression of ITIH4 mRNA in AGS cells with H. pylori infection

The expression level of ITIH4 mRNA in AGS cells began to increase 3 h after *H. pylori* infection, reached a peak at 6 h, and decreased to maintain low levels at 24 h. The statistics of ITIH4 in three independent experiments showed that the



Figure 1 The dynamic and melting curve of the ITIH4 and IL-6 genes by real-time PCR. (A, B, and C) are the amplification curves of ITIH4, IL-6, and GAPDH. (D, E, and F) are the melting curve of ITIH4, IL-6, and GAPDH, respectively. PCR, polymerase chain reaction; ITIH4, inter- $\alpha$ -trypsin inhibitor heavy chain 4; IL-6, interleukin-6.

expression of ITIH4 in AGS cells was highest at 6h after *H*. *pylori* infection (P<0.01) (*Figure 2A,B*).

# Differential expression of IL-6 mRNA in AGS gastric cancer cells after H. pylori infection

To further clarify the mechanism of enhanced ITIH4 mRNA expression in AGS cells after *H. pylori* infection, we also analyzed the expression of IL-6 in *H. pylori*-infected gastric cancer cells. The results are shown in *Figure 2B,C*. The expression level of IL-6 mRNA in AGS cells began to

increase 3 h after *H. pylori* infection, and reached a peak at 6 h IL-6 values in three independent experiments showed that the expression of IL-6 mRNA in AGS cells increased at the same time as ITIH4, reached a peak at 6 h, and decreased after 12 h, but remained at a high level (P<0.05) (*Figure 2B,C*). We also performed correlation analysis of ITIH4 and IL-6 expression. The results revealed that the change of ITIH4 expression level in AGS cells was significantly correlated with the increase of IL-6 expression level (r=0.897, P<0.01) (*Figure 2D*). These data indicated that *H. pylori* could induce the expressions of ITIH4 and IL-6 genes in a time



**Figure 2** Expressions of ITIH4 and IL-6 genes and proteins in AGS gastric cancer cells after *H. pylori* infection. (A) is the qPCR product of ITIH4 and IL-6 genes in AGS cells with *H. pylori* infection by agarose gel at different time points. (B) is the comparison of ITIH4 gene expression quantity in AGS gastric cancer cells after *H. pylori* infection. (C) is the expression of IL-6 mRNA in AGS gastric cancer cells after *H. pylori* infection. (D) is the correlation analysis of ITL-6 and ITIH4 mRNA changes in AGS cells after *H. pylori* infection. ITIH4, inter- $\alpha$ -trypsin inhibitor heavy chain 4; IL-6, interleukin-6; qPCR, quantity polymerase chain reaction; *H. Pylori*, *Helicobacter pylori*.

dependent manner. Moreover, the expression pattern of ITIH4 showed a strong positive correlation with IL-6expression.

# Changes of ITIH4 and IL-6 protein expression in gastric cancer cells after H. pylori infection

To further clarify the relationship between ITIH4 expression and *H. pylori* infection and IL-6 expression, Western blot and ELISA techniques were used to detect the expression levels of ITIH4 and IL-6 proteins in lysates of AGS cells infected with *H. pylori*. AGS cells were collected at a predetermined time to obtain the cell lysates. Western blot analysis showed that ITIH4 protein increased 3 h after *H. pylori* infection and reached a peak at 12 h. In contrast, the IL-6 protein also started to increase at 3 h after infection, but reached its peak at 6 h (*Figure 3A*). ELISA

results revealed that the expression of ITIH4 and IL-6 proteins was similar to the Western blot pattern, but the level of IL-6 was higher than that of ITIH4 at several time points (*Figure 3B*). This analysis confirmed the expression pattern of ITIH4 and IL-6 in AGS cells with *H. pylori* infection.

# *Effect of recombinant human interleukin 6 (rbIL-6) on the expression of ITIH4 in gastric cancer cells*

Different concentrations of rhIL-6 were used to induce AGS gastric cancer cells. Real-time quantitative PCR showed that as the concentration of rhIL-6 increased, the expression of ITIH4 mRNA increased in a dose-dependent manner (*Figure 4A*). ELISA also showed that the level of ITIH4 protein in the AGS cell lysate increased linearly as the concentration of rhIL-6 increased. At a concentration



Figure 3 Expression of IL-6 and ITIH4 in AGS gastric cancer cells after H. pylori infection. (A) is ITIH4 and IL-6 expression in AGS cell by Western blot. (B) is the quantitative comparison of ITIH4 and IL-6 protein in AGS cells after H. Pylori infection by ELISA. ITIH4, inter-α-trypsin inhibitor heavy chain 4; IL-6, interleukin-6; *H. Pylori*, *Helicobacter pylori*; ELISA, enzyme-linked immunosorbent assay.



Figure 4 Expression of ITIH4 mRNA in AGS cells induced by rhIL-6. (A) is the ITIH4 protein expression in AGS cells with different amounts of IL-6 by Western blot. (B) is the expression level of ITIH4 in AGS cell lysate induced with rhIL-6 by ELISA. ITIH4, intera-trypsin inhibitor heavy chain 4; rhIL-6, recombinant human interleukin-6; H. Pylori, Helicobacter pylori; ELISA, enzyme-linked immunosorbent assay.

of 50 ng/mL, the expression of ITIH4 reached a plateau (Figure 4B). This suggests that the expression of ITIH4 in gastric cancer cells is closely related to the level of IL-6.

#### Discussion

The course of gastric cancer is a multifactor and multistep process. The occurrence and development of gastric cancer generally progresses through chronic superficial gastritis, atrophy, intestinalization, dysplasia, metamorphosis, and finally results in gastric mucosal cancer (9). More than 60% of gastric cancer cases originate from H. pylori (12), with smoking and eating particular foods, like pickled vegetables, also frequently being linked to this cancer (13). After its initial discovery, ample research has confirmed

H. pylori as a class I carcinogen of gastric cancer (14,15). Chronic inflammation, ulcers, atrophy, intestinal dysplasia, and dysplasia of gastric mucosa are caused by chronic H. pylori infection, with many cytokines and inflammatory factors being involved in this process (9,10). Recent studies have shown that peripheral serum levels of IL-6, IL-1, transforming growth factor  $\beta$  (TGF- $\beta$ ), and tumor necrosis factor  $\alpha$  (TNF- $\alpha$ ) are significantly increased in patients with gastric cancer patients (5,11,16). Some studies have suggested that the increase of serum IL-6 and IL-6R is closely related to the prognosis of tumors (16,17). Therefore, it is believed that serum IL-6 level can be used as a marker for the serological diagnosis and prognosis of gastric cancer.

Previous studies have shown that serum ITIH4 level is

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specifically elevated in patients with early gastric cancer (18,19). In this study, H. Pylori was first co-cultured with AGS cells. The results showed that the expression levels of ITIH4 and IL-6 genes in AGS cells increased simultaneously 3 h after H. pylori infection and reached a peak at 6 h. The expression level of IL-6 could be maintained at a high level after having increased; however, the expression of ITIH4 tended to decline at 12 h and returned to a lower level 24 h after infection. With changes of mRNA expression, the expression levels of ITIH4 and IL-6 proteins in AGS cells also presented a similar response after H. pylori infection. These results suggest that ITIH4 is an acute phase response protein for *H. pylori* infection. IL-6 expression level reached a peak after 6 h, and then decreased and maintained a moderate level, indicating that the elevated ITIH4 level may be mediated via IL-6. Therefore, the early rise of ITIH4 in AGS cell after H. pylori infection also suggests that it was a characteristic serum marker for the early stages of gastric cancer.

IL-6 is an inflammatory factor secreted by monocytes, endothelial cells, and various tumor cells (20). It not only plays an important role in immune response and inflammatory response, but also participates in the occurrence and development of tumors. IL-6 has two-way effects on the occurrence and development of tumors (21). For one, IL-6 can activate macrophages and prevent neutrophil apoptosis, thereby enhancing the killing effect of cancer cells; for another, IL-6 can induce vascular endothelial cell growth factor and thus contributes to tumor invasion. IL-6 can regulate the cell cycle of tumor cells through the JAK/ STAT signalling pathway and affect tumor angiogenesis (22). The results of this experiment showed that AGS cells can be induced to express IL-6 and ITIH4 at the same time after *H. pylori* infection, suggesting that IL-6 may participate in the occurrence and development of *H. pylori*-related gastric cancer. The increase in ITIH4 expression in AGS cells after H. pylori infection may be the result of increased IL-6 expression. In order to verify this finding, rhIL-6 was applied to induce AGS cells, and the expression of ITIH4 mRNA and protein were monitored at different IL-6 concentrations. The results showed that the expression of ITIH4 increased at a concentration of 6.25 ng/mL, and reached a plateau phase at a concentration of 50 ng/mL, suggesting that there may be some restriction or feedback relationship between the expression of ITIH4 and IL-6.

ITIH4 protein is often associated with the increased expression of ITIH4 in serum under various diseases, such as rheumatoid arthritis, fatty liver infection, chronic hepatitis, and cirrhosis (23,24). Persistent *H. pylori* infection can lead to sustained high levels of IL-6 expression, which in turn can positively increase the expression of ITIH4 via activating the STAT3 signalling pathway (25). Abnormal activation of STAT3 is associated with the occurrence and development of gastric cancer (26,27). This also implies that the elevated levels of ITIH4 and IL-6 in peripheral serum may be relevant to assessing the risk of canceration by being able to rule out infectious disease and chronic inflammation diagnoses, especially for patients with H. pylori infection. In short, they may function as early warning indicators of gastric cancer.

#### Conclusions

The detection of ITIH4 and IL-6 levels in gastric cancer cells were useful biomarkers for early gastric cancer diagnosis. The elevated ITIH4 level in gastric cancer cells is mediated by IL-6.

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#### Footnote

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*Conflicts of Interest:* All authors have completed the ICMJE uniform disclosure form (available at http://dx.doi. org/10.21037/tcr-20-1766). 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. This study protocol was conducted in accordance with the Declaration of Helsinki (as revised in 2013) and approved by ethical committee of Shanghai General Hospital (Approval #:2017KY210). Informed consent was not required because we only used human cancer cell line in the entire experiment.

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