TNF-α promotes colon cancer cell migration and invasion by upregulating TROP-2

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Abstract. High levels of tumor-associated calcium signal transduction protein (TROP)-2 have been demonstrated to be strongly associated with tumor necrosis factor (TNF)-α levels in colon cancer. In the present study, the effect of TNF-α on the regulation of TROP-2 expression and its effect in colon cancer cell migration and invasion were investigated in vitro. TROP-2 protein levels were evaluated in HCT-116 human colon cancer cells cultured with various concentrations of TNF-α using western blot analysis. Changes in the migratory and invasive potential of the cells were evaluated using a wound healing and transwell assay, respectively. Then, TROP-2 expression was downregulated in cells by use of siRNA, and TROP-2 knockdown was confirmed at the mRNA and protein level by reverse transcription-quantitative polymerase chain reaction and western blotting, respectively. The migration and invasion potential of cells transfected with TROP-2 siRNA was also evaluated. Levels of several mitogen-activated protein kinase proteins, namely p38, c-Jun N-terminal kinase (JNK) and extracellular signal-regulated kinase (ERK), were detected in cells treated with TNF-α using western blot analysis. The results demonstrated that TROP-2 protein levels increased in cells treated with lower concentrations of TNF-α, but decreased in cells treated with higher concentrations of TNF-α, compared with untreated control. Maximum TROP-2 levels were observed in cells treated with 20 µg/l TNF-α. Migration and invasion were enhanced in cells treated with 20 µg/l TNF-α. When TROP-2 was silenced in colon cancer cells by siRNA, migration and invasion were significantly decreased compared with control cells. TNF-α stimulation activated the ERK1/2 pathway, but did not significantly affect p38 and JNK phosphorylation levels. Treatment with a specific ERK1/2 inhibitor suppressed the TNF-α-induced upregulation of TROP-2 and the TNF-α-induced colon cancer cell migration and invasion. In conclusion, the present results demonstrated that low concentrations of TNF-α significantly enhanced colon cancer cell migration and invasion by upregulating TROP-2 via the ERK1/2 signaling pathway.

Introduction

Colon cancer is a common malignant tumor of the digestive tract, from which morbidity and mortality have been increasing in recent years. Tumor invasion and metastasis are the main causes of colon cancer-associated death, but the precise mechanism mediating these processes remains unclear. Previous studies have primarily focused on the characteristics of cancer cells; however, the effects of the tumor microenvironment on invasion and metastasis have increasingly been gaining attention in recent years (1).

The tumor microenvironment has an important role in tumorigenesis and disease progression. Tumor necrosis factor (TNF)-α is an inflammatory cytokine frequently present in the tumor microenvironment. TNF-α is primarily secreted by tumor-associated macrophages and it initiates chronic inflammation (2). TNF-α has biphasic effects, by which it induces tumor cell apoptosis at high doses while it accelerates tumor invasion and metastasis with long-term administration of low doses (3). Tumor cell invasion is potentiataed when tumor cells are co-cultured with macrophages, an effect primarily mediated by the secretion of TNF-α by macrophages (4). In addition, TNF-α induces the expression of angiogenic factors, promotes tumor angiogenesis, and accelerates tumour metastasis (5). However, the specific molecular mechanisms by which TNF-α facilitates tumor cells migration and invasion are unclear.

Tumor-associated calcium signal transducer protein (TROP)-2 is a transmembrane glycoprotein expressed at high levels in a variety of human epithelial tumors, including oral (6), pancreatic (7), gastric (8), ovarian (9), lung (10), and prostate cancer (11). By contrast, TROP-2 is barely detectable in non-cancerous tissues. Furthermore, TROP-2 is strongly associated with tumor invasion and metastasis, and with disease prognosis (12,13). A previous study from our group have demonstrated that both TROP-2 and TNF-α levels are

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Abbreviations: TNF-α, tumor necrosis factor-α; TROP-2, tumor-associated calcium signal transducer protein-2

Key words: colon cancer, tumor necrosis factor-α, tumor-associated calcium signal transduction protein-2, tumor invasion, metastasis
elevated in colon cancer cells and that TROP-2 and TNF-α levels are directly associated (12). The aim of the present study was to explore the effect of TNF-α on the regulation of TROP-2 levels in the context of colon cancer cell migration and invasion.

**Materials and methods**

**Cells and reagents.** The human colon cancer cell line HCT-116 was kindly provided by Professor Haitao Shen (Department of Pathology, Hebei Medical University, Hebei, China). The cells were frozen in liquid nitrogen. Recombinant human TNF-α was purchased from Sigma-Aldrich (Merck KGaA, Darmstadt, Germany). The Lipofectamine 2000 transfection kit was purchased from Invitrogen (Thermo Fisher Scientific, Inc., Waltham, MA, USA). TROP-2 small interfering (si)RNA (5'-CGT GGACAACGTGCCCTCTA-3') and the negative control siRNA (5'-UUCUCCGAACUGUGACAGUTT-3') were purchased from Biovol Biotechnology (Shanghai, China). Matrigel, a mixture of basement membrane matrix proteins, was purchased from BD Biosciences (Franklin Lakes, NJ, USA). Monoclonal mouse anti-human primary TROP-2 antibody (cat. no. ab65006; 1:1,000) and GAPDH antibody (cat. no. ab8245; 1:1,000) were purchased from Abcam (Cambridge, MA, USA). Monoclonal mouse antibodies against human phosphorylated (p)-c-Jun N-terminal kinase (JNK; cat. no. 9251; 1:1,000), total (t)-JNK (cat. no. 9252; 1:1,000), p-ERK1 (cat. no. 9101; 1:1,000), t-ERK1 (cat. no. 9212; 1:1,000), p-extracellular signal-regulated kinase (p-ERK) 1/2 (cat. no. 9107;1:1,000) were purchased from Cell Signaling Technology, Inc. (Danvers, MA, USA). The specific p-ERK1/2 inhibitor PD98059 was purchased from Cayman Chemical Company (Ann Arbor, MI, USA). Horseradish peroxidase-conjugated secondary antibody was purchased from Beijing Zhongshan Jinqiao Biotechnology Co., Ltd. (1:4,000; Beijing, China).

Cell culture. HCT-116 human colon cancer cells were cultured in McCoy's 5A medium (Gibco; Thermo Fisher Scientific, Inc., Waltham, MA, USA) supplemented with 10% fetal bovine serum (Invitrogen; Thermo Fisher Scientific, Inc.) and 1% antibiotics (penicillin and streptomycin). The cells were incubated at 37°C in a 5% CO₂ atmosphere.

Western blot analysis. HCT-116 cells were seeded in culture dishes (5x10⁵ cells/dish) and cultured until they reached 80% confluence. Then, the cells were cultured in serum-free medium for an additional 24 h. The cells were subsequently incubated with various concentrations of TNF-α for the indicated period of time. Once cells were harvested, total proteins (50 µg) were separated using SDS-PAGE (10% gel) and the separated proteins were transferred to polyvinylidene fluoride membranes. The membranes were blocked with 5% skim milk at room temperature for 1 h, and subsequently incubated at 4°C overnight with the primary antibody of interest and anti-GAPDH as a loading control. Then, the membranes were washed three times with TBS/0.1% Tween-20 and incubated with goat anti-rabbit secondary antibody at room temperature for 2 h. The protein signals were detected using enhanced chemiluminescence (Beyotime Institute of Biotechnology, Haimen, China). Western blot results were analyzed using ImageJ software (version 1.41; National Institutes of Health, Bethesda, MD, USA).

Wound healing assay. HCT-116 cells in the logarithmic phase were seeded in 6-well plates. Once the cells reached 80% confluence, wounds were generated by scratching the cells with a 100-µl pipette tip. Then, cells were washed with PBS and treated with TNF-α at a final concentration of 20 µg/l in serum-free medium for 24 h. Control cells were treated with PBS in serum-free medium. Images of the cells were captured at 0 and 24 h after the wound was created. The width of the wound at 5 randomly selected sites was measured using a Leica TCS-SP5 confocal microscope (magnification, x200; Leica Microsystems, Inc., Buffalo Grove, IL, USA), and the average wound width was calculated. The wound healing rate was calculated using ImageJ (version 2.1; National Institutes of Health) (14). The experiment was conducted in triplicate.

Transwell invasion assay. The invasive ability of cells was assessed using a Transwell assay. The invasive potential of HCT-116 cells was assessed by determining the number of cells that passed through a polycarbonate membrane (8 µm pore size) in 24-well transwell chambers. The upper Transwell filters (Corning Inc., Corning, NY, USA) were coated with Matrigel (BD Biosciences). Single cell suspensions of HCT-116 cells were cultured until they reached the logarithmic phase, and the cells (1x10⁵) were subsequently seeded into the upper chamber of Transwell filters in serum-free medium supplemented with 20 µg/l TNF-α. Complete medium was added to the lower chambers. The cells were incubated for 24 h, and non-migratory cells on the upper side of the filter were removed using cotton swabs. Cells that had migrated through the pores of the filter were fixed in methanol at 4°C for ~20-30 min, stained with 0.1% crystal violet for 30 min at 37°C, and washed with PBS. The number of migratory cells was determined using an inverted light microscope at a magnification of x200. The average number of cells in five randomly selected fields was calculated, and the experiment was conducted in triplicate.

siRNA-mediated silencing of TROP-2. In order to knock-down endogenous TROP-2 expression, TROP-2-specific siRNA was transfected into cells using Lipofectamine® 2000 transfection reagent (Invitrogen; Thermo Fisher Scientific, Inc.) according to the manufacturer's instructions. Sequences were as follows: TROP-2 siRNA, 5'-CGTGGACAACGTGCCCTCTA-3'; control siRNA, 5'-UUCUCCGAACUGUGACAGUTT-3'. Cells were transfected with TROP-2 siRNA (50 nM) or negative control siRNA (50 nM) using Lipofectamine® 2000. Non-transfected cells (blank control group) were included in the analysis. Two days after transfection, transfected cells were harvested and TROP-2 mRNA and protein expression was evaluated by reverse transcription-quantitative polymerase chain reaction and western blot analysis.

RT-qPCR. RT-qPCR was used to determine TROP-2 mRNA expression levels of the human colon cancer HCT-116 cell line.
Primers were as follows: TROP-2 forward, 5’-TCACCAACC GGAGAAAGTCG-3’ and reverse, 5’-AGGAAGCGGTGACT CACTTGG-3’; β-actin forward, 5’-CACGAAAATCCCTTC AACTCC-3’ and reverse, 5’-CATACTCCTGCTTGCTGTA TC-3’. Following transfection, HCT-116 cells were harvested for total RNA extract using TRIzol reagent (Invitrogen; Thermo Fisher Scientific, Inc.). For reverse transcription, RNase-free EP tubes containing Mix I (5 µl total RNA, 0.5 µl of 50 µmol/l oligo(dT), 0.5 µl random primer, 1 µl of 10 mmol/l dNTP solution and 5 µl diethylpyrocarbonate-treated water) were incubated in a 65°C water bath for 5 min and immediately transferred to ice water for 1 min. Then, 4 µl of 5X first-strand buffer, 2 µl of 0.1 mol/l DTT, 1 µl of 40 U/µl RNaseOUT (Invitrogen; Thermo Fisher Scientific, Inc.), and 1 µl SuperScript III reverse transcriptase (Invitrogen; Thermo Fisher Scientific, Inc.) were added to Mix 1 for a total reaction volume of 20 µl. The reactions were incubated as follows: 5°C for 5 min, 50°C for 60 min, and 70°C for 15 min. The resulting cDNA products were placed on ice immediately following the reaction. qPCR was performed using SYBR Premix DimerEraser (Takara Biotechnology Co., Ltd., Dalian, China). The conditions for the qPCR were as follows: 50°C for 30 min, a denaturation step at 95°C for 2 min, 40 cycles of 95°C for 10 sec and 60°C for 30 sec, and an elongation step at 70°C for 30 sec. All experiments were conducted in triplicate. Relative gene expression levels were calculated using the 2^ΔΔCq method (15).

Statistical analysis. SPSS software (version 17.0; SPSS, Inc., Chicago IL, USA) was used for statistical analyses. Data with a normal distribution are presented as the mean ± standard deviation. A t-test was used to compare differences in means between two groups. One-way ANOVA was used to compare differences in means among multiple groups, and the student Newman-Keuls-q method was used for multiple comparisons between groups. *P<0.05 was considered to indicate a statistically significant difference.

Results

Effect of TNF-α on TROP-2 levels in colon cancer cells. Previous studies from our group have demonstrated that TNF-α and TROP-2 protein levels are abundant in colon cancer, and higher levels of these proteins are associated with a poor prognosis (12). To investigate the effect of TNF-α on TROP-2 expression, HCT-116 colon cancer cells were incubated with various concentrations of TNF-α, and TROP-2 protein levels were subsequently examined using western blot analysis. As illustrated in Fig. 1, TROP-2 protein expression levels exhibited a biphasic response, increasing in cells incubated with decreased concentrations of TNF-α (10 and 20 µg/l), but decreasing in cells incubated with increased concentrations (30, 50, 100 and 200 µg/l). TROP-2 protein levels were the highest in cells incubated with 20 µg/l TNF-α. These results demonstrated that low concentrations of TNF-α upregulated TROP-2 expression in HCT-116 cells. Therefore, TNF-α at a concentration of 20 µg/l was used in the subsequent experiments.

Effect of TNF-α on TROP-2 regulation. Little is known about the signaling pathways that mediate TNF-α-induced upregulation of TROP-2. The mitogen-activated protein kinase (MAPK) signaling pathway is involved in multiple biological processes in tumors, including cell proliferation, cell cycle regulation, migration, and invasion. Since the MAPK pathway can be activated by a variety of inflammatory cytokines and TROP-2 is associated with aggressive tumor cell behaviors, the role of MAPK signaling in TNF-α-mediated regulation of TROP-2 was investigated. JNK, p38, and ERK1/2 are key members of the MAPK signaling pathway. As illustrated in Fig. 2, p-ERK1/2 levels increased in cells treated with TNF-α at 15 min post-stimulation, whereas there were no significant changes in p-JNK and p-p38 levels. Fig. 3 demonstrates that, compared with cells treated with TNF-α alone, p-ERK1/2 and TROP-2 protein levels significantly decreased in cells that were pretreated with the specific ERK1/2 inhibitor PD98059 (20 µM) for 30 min. Together, the present data suggested that TNF-α stimulation upregulated TROP-2 protein expression levels by activating the ERK1/2 signaling pathway.

TNF-α promotes colon cancer cell migration and invasion by upregulating TROP-2. Aggressive cancer cells are characterized by an increase in cell migration and invasion abilities. To investigate the role of TNF-α in metastasis, the effect of TNF-α stimulation on migration and invasion of HCT-116 cells was evaluated using wound healing and transwell assays, respectively. As presented in Fig. 4, the rate of
wound healing and the number of invasive cells at 24 h were significantly increased in the TNF-α-treated group compared with the control group. These results indicated that TNF-α significantly enhanced the migratory and invasive potential of HCT-116 colon cancer cells.

Although it was demonstrated that TNF-α upregulated TROP-2 protein levels and enhanced the invasive potential of colon cancer cells, it remained unclear if the latter effect was mediated by the upregulation of TROP-2. Therefore, the role of TROP-2 in the TNF-α-induced migration and invasion of colon cancer cells was examined by silencing TROP-2 expression with a specific siRNA. The negative control group was transfected with a non-specific scrambled siRNA, and non-transfected cells were used as a blank control. As presented in Table I and Fig. 5, TROP-2 expression levels in cells transfected with TROP-2 siRNA significantly decreased at both the mRNA and protein level compared with control groups. When migration and invasion were examined, the rate of TNF-α-induced wound healing and number of invasive cells at 24 h were significantly decreased in the TROP-2 siRNA group compared with the negative control siRNA group (Fig. 6). In addition, the rate of TNF-α-induced wound healing and invasion significantly decreased in cells pretreated with PD98059 (20 µM) for 30 min, compared with cells treated with TNF-α alone (Fig. 7). These data demonstrated that TROP-2 knockdown inhibited TNF-α-induced invasion and migration in colon cancer cells, and that TNF-α promoted invasion and migration in colon cancer cells by...
upregulating TROP-2 protein expression via the ERK1/2 signaling pathway.

Discussion

The tumor microenvironment is a popular topic in cancer research, and scholars consider tumor-associated inflammation as the seventh hallmark of cancer (16). The inflammatory microenvironment is strongly associated with tumorigenesis and cancer progression; however, the mechanism by which it promotes invasion and metastasis remains unclear. TNF-α is a key inflammatory cytokine present in the tumor microenvironment. Although TNF-α has been linked to tumor invasion and metastasis, the specific mechanisms underlying this association are unclear. Previous studies have demonstrated that TNF-α can activate the MAPK/ERK pathway, upregulate matrix metalloproteinase (MMP)-9, CD26 and fibroblast activation protein (FAP)-α levels, and enhance tumor cell invasion and metastasis in breast cancer (17). TNF-α may also upregulate MMPs in oral cancer cells, thereby promoting invasion and metastasis (18).
In addition, TNF-α promotes lymphangiogenesis and lymphatic metastasis in gallbladder carcinoma by activating the ERK1/2 or the activator protein-1/vascular endothelial growth factor (VEGF)-D pathway (19). However, there are no previous reports describing the role of TNF-α in colon cancer invasion and metastasis.

TROP-2 is a transmembrane glycoprotein expressed at high levels in a variety of human epithelial tumors. TROP-2 is strongly associated with tumor invasion and metastasis, and high levels of TROP-2 are associated with a poor prognosis in gastric carcinoma (8). However, the mechanism underlying TROP-2 overexpression in tumors remains unclear.

Figure 6. Effect of TNF-α on migration and invasion of TROP-2 silenced HCT-116 cells. Representative images and quantification of wound healing rate (magnification, x100) and Transwell invasion (magnification, x200). TNF-α, tumor necrosis factor-α; TROP-2, tumor-associated calcium signal transducer protein-2; siRNA, small interfering RNA; N, negative control.

Figure 7. Effect of the ERK1/2 inhibitor PD98059 on migration and invasion of TNF-α-stimulated HCT-116 cells. Representative images and quantification of wound healing rate (magnification, x100) and Transwell invasion (magnification, x200). ERK, extracellular signal-regulated kinase; TNF-α, tumor necrosis factor-α.
The mechanism by which TROP-2 is upregulated in tumors may provide insights that could facilitate the development of approaches for early diagnosis and treatment.

In previous studies, both TROP-2 and TNF-α were markedly expressed at the protein level in a large number of colon cancer tissue samples (12). Therefore, it was speculated that TNF-α may upregulate TROP-2 protein levels in colon cancer cells, thereby promoting tumor cell migration and invasion. The findings of the present study validate this hypothesis.

In the present study, the effect of various concentrations of TNF-α on TROP-2 protein levels was examined in colon cancer cells. TROP-2 protein levels increased in cells incubated with low concentrations of TNF-α and decreased in cells incubated with higher concentrations. Maximum TROP-2 levels were observed in cells treated with 20 μg/l TNF-α. It is hypothesized that TROP-2 levels are decreased in cells treated with high concentrations of TNF-α potentially due to increased cytotoxicity. Wound healing and cell invasion were enhanced in cells treated with 20 μg/l TNF-α compared with control untreated cells. Thus, the present data demonstrated that low concentrations of TNF-α upregulated TROP-2 protein expression and promoted colon cancer cell migration and invasion.

To further confirm the hypothesis that low concentrations of TNF-α promote invasion in colon cancer cells by upregulating TROP-2, expression of TROP-2 was silenced using siRNA technology. Cell migration and invasion were significantly inhibited in TROP-2 siRNA-transfected HCT-116 cells treated with TNF-α, compared with control siRNA-transfected cells. These data demonstrate that the inflammatory cytokine TNF-α promotes migration and invasion in colon cancer cells by upregulating TROP-2.

To date, there have been no reports describing the mechanism by which inflammatory cytokines, such as TNF-α, upregulate TROP-2. Choo et al (20) demonstrated that TNF-α may induce colon cancer cells to metastasize to the lungs by activating the ERK signaling pathway. Hagemann et al (21) reported that TNF-α secretion by tumor-associated macrophages promotes the invasion and metastasis of breast cancer cells by activating the JNK pathway and upregulating MMPs and VEGF. Mathukumaran et al reported that TNF-α enhances the metastatic potential of SKA and SKOV-3 ovarian cancer cells by activating the JNK signaling pathway and modulating CD44 levels (22). Therefore, it was hypothesized in the present study that TNF-α may upregulate TROP-2 in HCT-116 cells by activating the MAPK pathway. The results demonstrated that p-ERK1/2 levels significantly increased in cells treated with TNF-α and this effect was significantly inhibited in cells pretreated with PD98059, a specific ERK1/2 inhibitor. The effect of PD98059 on wound healing and invasion assays suggested that the TNF-α-induced tumor migration and invasion may be mediated by ERK1/2. By contrast, the phosphorylation levels of the MAPK proteins p38 and JNK did not significantly change in cells treated with TNF-α, suggesting that these proteins do not serve a role in the TNF-α-induced upregulation of TROP-2.

In conclusion, TNF-α is a key inflammatory cytokine in the tumor microenvironment, and the present results suggest that it may promote colon cancer invasion by upregulating TROP-2 expression via the ERK1/2 signaling pathway. Therefore, the ERK1/2 pathway may represent a potential therapeutic target for the treatment of colon cancer.

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