

Original Article

TRAIL affects NF- κ B and Survivin expressions in gastric cancer cell

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Abstract: Gastric cancer is a common malignancy in digestive tract. Tumor necrosis factor related apoptosis-inducing ligand (TRAIL) can selectively induce tumor cell apoptosis. Survivin and NF- κ B participate in the cell apoptosis process induced by TRAIL. This study investigated the impact of TRAIL on NF- κ B and Survivin expressions in gastric cancer cell and related mechanism. Gastric cancer cell line SGC-7901 was treated by TRAIL at 100 μ g/L, 200 μ g/L, and 500 μ g/L for 12 h, 24 h, and 48 h, respectively. Cell proliferation was tested by MTT assay. Cell apoptosis was evaluated by flow cytometry. Cell invasion and migration were determined by Transwell assay. Survivin and NF- κ B expressions were detected by Western blot. TRAIL significantly inhibited cell proliferation at different concentrations and times compared with control ($P < 0.05$). TRAIL intervention at 500 μ g/L for 48 h exhibited the strongest suppressive effect on cell proliferation. It obviously promoted cell apoptosis and declined cell invasion and migration ($P < 0.05$). Survivin and NF- κ B protein expressions markedly down regulated in SGC-7901 cells treated by TRAIL with time and dose dependence ($P < 0.05$). TRAIL significantly inhibited SGC-7901 cell proliferation, facilitated cell apoptosis, and restrained cell invasion and migration through suppressing Survivin and NF- κ B.

Keywords: TRAIL, gastric cancer, NF- κ B, Survivin

Introduction

Gastric cancer is a common malignant tumor of the digestive system. Tumor cell apoptosis plays an important role in the occurrence and development processes of cancer. It is mainly presented as the inactivation of apoptosis-inducing gene and overexpression of apoptosis inhibiting gene. Blockage of cell apoptosis appears in precancerous lesions. Some molecules participate in the signaling transduction to affect apoptosis [1-4]. Tumor necrosis factor related apoptosis inducing ligand (TRAIL) is a member of the tumor necrosis factor family that can selectively induce tumor cell apoptosis without significant cytotoxicity [5]. It was showed that TRAIL is not sensitive to all the malignant cells. Survivin and NF- κ B are the critical factors in cell apoptosis pathway induced by TRAIL [6]. Survivin, a new member of the apoptosis inhibiting protein family, is the strongest apoptosis inhibiting factor so far. It was found that Survivin has dual effect on cell behavior, including apoptosis inhibition

and proliferation regulation. It plays inhibiting effect through directly suppressing Caspase-3 and Caspase-7, which is specifically expressed in a variety of tumors [7]. NF- κ B is a kind of transcription factor existed in eukaryocytes with multiple biological functions. It regulates various genes expressions, such as cytokines, adhesion molecules, and growth factors. Its relationship with tumor attracts attention because of its role as a pleiotropic regulatory protein [8]. P65 is the major component of the active NF- κ B. It plays a central role in tumor cell apoptosis and proliferation as a multifunction transcription factor. Its aberrant transcription is closely associated with the up regulation of anti-apoptotic genes. It was demonstrated that NF- κ B plays a critical role in multiple malignant tumors, such as leukemia, lymphoma, and solid tumors. It is still controversy about the role of Survivin and NF- κ B in the occurrence and development of gastric cancer. This study used TRAIL to treat gastric cancer cell line SGC-7901 to analyze its impact on Survivin and NF- κ B expressions and related mechanism.

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Table 1. TRAIL affected SGC-7901 cell proliferation (OD value)

Time	Experimental group			Control
	100 µg/L	200 µg/L	500 µg/L	
12 h	0.985±0.072 ¹	0.723±0.046 ^{1,2}	0.232±0.024 ^{1,2,3}	0.071±0.021
24 h	0.811±0.045 ^{1,4}	0.512±0.034 ^{1,2,4}	0.176±0.016 ^{1,2,3,4}	0.072±0.017
48 h	0.321±0.031 ^{1,4,5}	0.256±0.026 ^{1,2,4,5}	0.121±0.011 ^{1,2,3,4,5}	0.067±0.019

¹P < 0.05, compared with control; ²P < 0.05, compared with 100 µg/L; ³P < 0.05, compared with 200 µg/L; ⁴P < 0.05, compared with 12 h; ⁵P < 0.05, compared with 24 h.

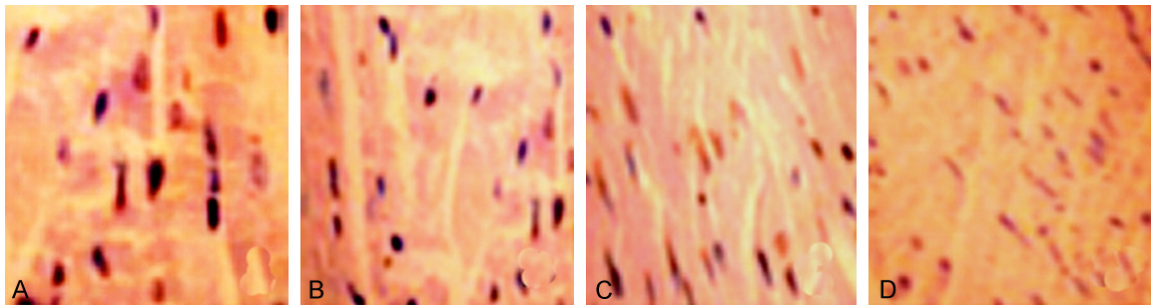


Figure 1. TRAIL affected SGC-7901 cell apoptosis (×400). A. 100 µg/L; B. 200 µg/L; C. 500 µg/L; D. Control.

Materials and methods

Experimental cells

Gastric cancer cell line SGC-7901 was provided by Soochow University (Suzhou, China).

Reagents

TRAIL (Sinopharm Chemical Reagent co., Ltd, lot no. 20050913). TUNEL assay kit (Roche, lot no. 11684817910). Rabbit anti human Survivin and NF-κB polyclonal antibodies, and goat anti rabbit secondary antibody (Santa Cruz). RPMI 1640 medium, penicillin-streptomycin, FBS (Gibco).

Instruments

Incubator (Thermo). Carbon dioxide incubator (SANYO). Inverted microscope (Nikon). Desktop low temperature centrifuge (Beckman). -80°C refrigerator (SANYO).

Methods

Routine cell culture

Gastric cancer cell line SGC-7901 was maintained in RPMI 1640 medium and cultured in incubator at 37°C and 5% CO₂.

TRAIL intervention

SGC-7901 cells in logarithmic phase were seeded in the plate overnight. After cultured in 2% FBS for 24 h, the cells were maintained in RPMI 1640 medium containing 10% FBS and treated by TRAIL at 100 µg/L, 200 µg/L, and 500 µg/L for 12 h, 24 h, and 48 h, respectively. Untreated SGC-7901 cells were selected as normal control.

MTT assay

The cells were treated by TRAIL at 100 µg/L, 200 µg/L, and 500 µg/L for 12 h, 24 h, and 48 h, respectively. The cells were added with 20 µl MTT solution at 5 mg/ml for 4 h. Then the cells were added with 150 µl DM-SO and vibrated for 10 min. At last, the plate was read at 570 nm to obtain the absorbance value.

TUNEL assay

The cells were treated by TRAIL at 100 µg/L, 200 µg/L, and 500 µg/L for 12 h, 24 h, and 48 h, respectively. The cells were washed by xylene and gradient ethanol. Next, the cells were treated by Proteinase K and added with 50 µl TUNEL solution. Then the cells were treated by 50 µl converter-POD together with 50-100 µl DAB

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Table 2. TRAIL affected SGC-7901 cell apoptosis

Time	Experimental group			Control
	100 µg/L	200 µg/L	500 µg/L	
12 h	11.18±1.02 ¹	17.27±1.44 ^{1,2}	21.24±2.45 ^{1,2,3}	3.01±1.26
24 h	15.21±1.33 ^{1,4}	18.48±1.67 ^{1,2,4}	24.37±2.63 ^{1,2,3,4}	3.21±1.43
48 h	18.26±1.54 ^{1,4,5}	20.85±2.12 ^{1,2,4,5}	28.92±2.77 ^{1,2,3,4,5}	3.32±1.39

¹P < 0.05, compared with control; ²P < 0.05, compared with 100 µg/L; ³P < 0.05, compared with 200 µg/L; ⁴P < 0.05, compared with 12 h; ⁵P < 0.05, compared with 24 h.

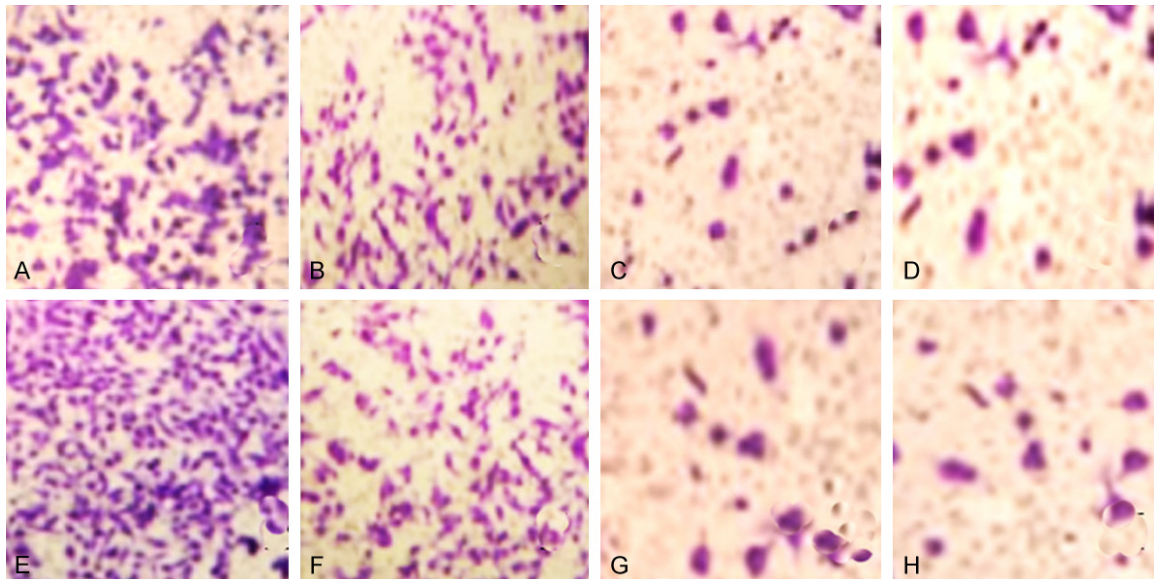


Figure 2. TRAIL affected SGC-7901 cell invasion and migration (×400). A. Control; B. 100 µg/L cell invasion; C. 200 µg/L cell invasion; D. 500 µg/L cell invasion; E. Control; F. 100 µg/L cell migration; G. 200 µg/L cell migration; H. 500 µg/L cell migration.

substrate. After redyed by hematoxylin or methyl green, the cells were counted and photographed.

Transwell assay

For cell invasion, the matrigel was mixed with serum free medium at 1:8 for 24 h and paved on the chamber. SGC-7901 cells were seeded at 10^6 /ml in 200 µl serum free medium. Another 1300 µl complete medium was added to the lower chamber. After 24 h, the upper chamber was fixed by ethanol and observed under the microscope after staining.

Western blot

Total protein was extracted and separated in 8% SDS-PAGE. After blocked at room temperature for 1 h, the membrane was incubated with primary antibodies (Survivin and NF-κB at 1:200, β-actin at 1:500, respectively) at 4°C

overnight. After washed by TBS-T, the membrane was incubated in secondary antibody (1:2000) for 1 h. Next, the membrane was treated by developer A and B and scanned. Quantity One software was used for band analysis.

Data analysis

SPSS 17.0 software was applied for data analysis. Measurement data were presented as mean ± standard deviation and analyzed by ANOVA. P < 0.05 was depicted as statistical significance.

Results

TRAIL affected SGC-7901 cell proliferation

Different concentrations of TRAIL were used to treat SGC-7901 cells for 12 h, 24 h, and 48 h. TRAIL significantly inhibited cell proliferation at

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Table 3. TRAIL affected SGC-7901 cell invasion and migration

Cell number	Experimental group			Control
	100 µg/L	200 µg/L	500 µg/L	
Invasion	127.23±15.33 ⁴	90.14±4.21 ^{1,2}	48.12±9.35 ^{1,2,3}	38.22±2.06
Migration	181.14±10.83 ⁴	99.28±3.61 ^{1,2}	152.55±7.03 ^{1,2,3}	37.62±2.86

¹P < 0.05, compared with control; ²P < 0.05, compared with 100 µg/L; ³P < 0.05, compared with 200 µg/L.



Figure 3. TRAIL affected Survivin and NF-κB protein expressions in SGC-7901 cells.

Table 4. TRAIL affected Survivin and NF-κB protein expressions in SGC-7901 cells

Item	Experimental group			Control
	100 µg/L	200 µg/L	500 µg/L	
Survivin				
12 h	0.71±0.21 ¹	0.66±0.18 ^{1,2}	0.16±0.11 ^{1,2,3}	0.74±0.37
24 h	0.54±0.15 ^{1,4}	0.44±0.13 ^{1,2,4}	0.10±0.03 ^{1,2,3,4}	0.72±0.38
48 h	0.33±0.13 ^{1,5}	0.28±0.09 ^{1,2,5}	0.14±0.01 ^{1,2,3,4,5}	0.69±0.41
NF-κB				
12 h	0.53±0.21 ¹	0.47±0.18 ^{1,2}	0.49±0.11 ^{1,2,3}	0.10±0.09
24 h	0.44±0.25 ^{1,4}	0.22±0.21 ^{1,2,4,5}	0.22±0.12 ^{1,2,3,4}	0.10±0.08
48 h	0.26±0.39 ^{1,4,5}	0.17±0.24 ^{1,2,4,5}	0.13±0.17 ^{1,2,3,4,5}	0.11±0.07

¹P < 0.05, compared with control; ²P < 0.05, compared with 100 µg/L; ³P < 0.05, compared with 200 µg/L; ⁴P < 0.05, compared with 12 h; ⁵P < 0.05, compared with 24 h.

different concentrations and times compared with control ($P < 0.05$). TRAIL intervention at 500 µg/L for 48 h exhibited the strongest suppressive effect on cell proliferation ($P < 0.05$) (Table 1).

TRAIL affected SGC-7901 cell apoptosis

TUNEL assay was applied to test the impact of TRAIL on SGC-7901 cell apoptosis. TRAIL obviously promoted cell apoptosis with dose and time dependence ($P < 0.05$) (Figure 1 and Table 2).

TRAIL affected SGC-7901 cell invasion and migration

Transwell assay was adopted to test the influence of TRAIL on SGC-7901 cell invasion

and migration. TRAIL markedly suppressed cell invasion and migration with dose and time dependence ($P < 0.05$) (Figure 2 and Table 3).

TRAIL affected Survivin and NF-κB protein expressions in SGC-7901 cells

Western blot was selected to determine Survivin and NF-κB protein expressions in SGC-7901 cells treated by TRAIL. Survivin and NF-κB protein expressions markedly down-regulated in SGC-7901 cells treated by TRAIL with time and dose dependence ($P < 0.05$) (Figure 3 and Table 4).

Discussion

Gastric cancer is a common digestive tract malignant tumor. Its occurrence pattern is a multi-stage process. Chronic gastritis may form atrophic gastritis, further leading to intestinal metaplasia and hyperplasia. It occurs canceration eventually, lead to the occurrence of gastric cancer [9]. TRAIL is a member of the tumor necrosis factor family shows high homology with FasL. It can induce most of

the tumor cell apoptosis [10, 11]. At present, several scholars proposed that Survivin may be related to the inhibition of apoptosis, the regulation of cell proliferation, and tumor angiogenesis [12]. NF-κB is a type of multifunction transcription factor that plays a central regulatory role in tumor cell proliferation and apoptosis. Its aberrant transcription activation is closely related to the upregulation of anti-apoptotic genes [13]. This study selected gastric cancer cell line SGC-7901 to analyze the impact of TRAIL on Survivin and NF-κB expressions and related mechanism.

Different concentrations of TRAIL were used to treat SGC-7901 cells for 12 h, 24 h, and 48 h. TRAIL significantly inhibited cell proliferation at different concentrations and times compared

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with control. TRAIL intervention at 500 µg/L for 48 h exhibited the strongest suppressive effect on cell proliferation. TUNEL assay was applied to test the impact of TRAIL on SGC-7901 cell apoptosis. TRAIL obviously promoted cell apoptosis with dose and time dependence. Transwell assay was adopted to test the influence of TRAIL on SGC-7901 cell invasion and migration. TRAIL markedly suppressed cell invasion and migration with dose and time dependence. TRAIL significantly inhibited SGC-77901 cell proliferation, facilitated cell apoptosis, and restrained cell invasion and migration. TRAIL belongs to TNF superfamily that can only induce apoptosis in transformed cells, tumor cells, and virus infected cells without normal cells. It was showed that TRAIL has certain apoptosis induction role on different types of gastric cancer cells with time dependence, such as AGS, MKN28, and MKN45 cell lines. However, the response of different cell lines to TRAIL was different, indicating the cell sensitivity to TRAIL was different [14, 15].

To explore the role of TRAIL on Survivin and NF-κB expressions in gastric cancer, this study applied different concentrations of TRAIL to intervene SGC-7901 cells. Survivin and NF-κB protein expressions markedly downregulated in SGC-7901 cells treated by TRAIL with time and dose dependence, suggesting that TRAIL can suppress Survivin and NF-κB expressions in gastric cancer. Survivin was found abnormally upregulated in gastric cancer, and its expression was higher in a tissue approaching the tumor infiltrated region [16]. Wakana collected 21 cases of intestinal type and diffuse type gastric cancer samples and found that Survivin mRNA and protein were negatively correlated with apoptotic index. It exhibited higher positive rate in diffuse type gastric cancer [17]. Over-expression of Survivin can inhibit cell apoptosis, elevate cell malignancy, and promoted cell metastasis in gastric cancer [18]. NF-κB demonstrated increasing trend in normal gastric tissue, adjacent tissue, and cancer tissue. Its excessive activation plays a role in the early and progressive stages of cancer [19, 20]. NF-κB suppresses gastric cancer apoptosis and facilitates cell proliferation and metastasis through regulating the downstream anti-apoptotic genes [21-23]. Our results suggested that TRAIL significantly inhibited SGC-77901 cell proliferation, facilitated cell apoptosis, and

restrained cell invasion and migration through suppressing Survivin and NF-κB.

TRAIL significantly inhibited SGC-77901 cell proliferation, facilitated cell apoptosis, and restrained cell invasion and migration through suppressing Survivin and NF-κB. It is worth of promoting Survivin and NF-κB detection in gastric cancer patients, especially in those received surgery, chemotherapy, or radiotherapy. Monitoring their expressions before and after treatment, together with clinicopathological characteristics may provide appropriate evaluation for the prognosis. More in-depth investigation on the aspects of animal model, gene, and protein may supply better therapeutic choice to improve the prognosis. The process of gastric cancer occurrence involves multiple cytokines with complex interactions. Precise detection of Survivin and NF-κB expressions may provide new strategy for the treatment of gastric cancer. Its specific mechanism still needs further exploration.

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Disclosure of conflict of interest

None.

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