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Case Control Study

Oncogenic role of Tc17 cells in cervical cancer development

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Abstract

BACKGROUND

As one of the subsets of CD8⁺ T cells, Tc17 cells have recently been identified and are characterized by the secretion of interleukin (IL)-17, which is related to inflammatory diseases.

AIM

To assess the status of Tc17 cells in cervical cancer and investigate the biological function of Tc17 cells in cervical cancer development.

METHODS

Flow cytometry assay, immunohistochemistry, and immunofluorescence were performed to detect the levels and phenotype of Tc17 cells in blood and tumor samples from patients with cervical cancer. Prior to cell suspension culture, ELISA was carried out to measure the production of IL-6, IL-1 β , IL-23, CXCL12, and IL-17 in tumor tissue supernatant and co-cultured supernatant of patients with cervical cancer. In addition, multivariate analysis was performed to identify factors associated with overall survival using the Cox proportional hazards model.

RESULTS

Compared with normal tissues, Tc17 cells specifically accumulated in tumor tissues of cervical cancer patients. Cancer cells produced a greater amount of IL-6, IL-1 β , and IL-23, which in turn promoted Tc17 cell polarization. Unlike the traditional cytotoxic CD8⁺ T cells, Tc17 cells secreted IL-17, which subsequently promoted CXCL12 expression in tumor cells, eventually enhancing the proliferation and migration of tumor cells. Thus, the ratio of tumor-infiltrating Tc17 cells was highly correlated with poor clinical outcome in patients with cervical cancer.

CONCLUSION

Our data identified the oncogenic role of Tc17 cells in the development of cervical cancer. We propose that the ratio of Tc17 cells may be a useful index in the prognosis of patients with cervical cancer.

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Core tip: Inflammation contributes to cancer development. In this study, it was found that cervical cancer-elicited inflammation increased Tc17-polarizing cytokine production, which attenuated cytotoxic CD8⁺ T cell development. The high level of interleukin-17 production by Tc17 cells led to CXCL12 upregulation and cancer cell migration. Consistent with the oncogenic role of Tc17 cells in cancer development, the ratio of cancer-infiltrating Tc17 cells was highly associated with poor prognosis in patients with cervical cancer. Thus, our data demonstrate that Tc17 cells can be induced in cervical cancers and serve as a meaningful index in the prognosis of patients with cervical cancer.

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INTRODUCTION

As the fourth most common malignant tumor worldwide, the incidence of new cervical cancer is approximately 130000, accounting for 28% of the total number of cases globally. About 20000 women die of cervical cancer each year^[1,2]. Tumor progression has been recognized to be the product of evolving crosstalk between immune cells and tumor cells. By affecting immune cell activation or differentiation, cancer cells escape host immune attack and enhance tumor resistance to immunotherapies^[3,4].

As the primary component of tumor-infiltrating lymphocytes, T cells elicit a crucial effector function in cancer eradication. Recent studies showed that T cells that produce interleukin (IL)-17 are detected in human tumors, which have a certain pro-inflammatory effect^[5]. On the one hand, Th17 cell polarizing factor may induce differentiation and proliferation of Tc17 cells, despite exhibiting reduced cytotoxic activity in CD8⁺ T cells, thereby interrupting host immune surveillance^[5,6]. Moreover, in a monkey immunodeficiency virus infection model with mammals such as macaques and black-and-white marmosets, Tc17 cells play a pathological role in promoting disease progression^[7,8]. However, the role of Tc17 cells in human cervical cancer development remains unclear.

A large number of Tc17 cells were found in black and white monkey tumor tissues, which were induced by the cytokine IL-23^[8]. At the same time, other studies have shown that IL-6 may also be crucial for Tc17 cell differentiation^[9]. However, it is unclear whether other cytokines can induce Tc17 cell differentiation and how Tc17 cells affect cancer development is still poorly understood.

In this study, we showed that Tc17 cells are highly enriched within cervical cancer tissue. Cervical cancer cells produced a large amount of IL-6, IL-1 β , and IL-23, which induced Tc17 cell polarization. Increased levels of IL-17 induced by Tc17 cells led to CXCL12 upregulation in tumor cells, resulting in tumor cell proliferation and migration. Moreover, the percentage of Tc17 cells was associated with tumor progression and clinical outcome in patients with cervical cancer. Our data demonstrate the oncogenic role of Tc17 cells in cancer development and provide a theoretical basis for the clinical treatment of cervical cancer.

MATERIALS AND METHODS

Patients and tissue specimens

Fresh blood, tumor, peritumoral, or matched adjacent tissues (at least 5 cm distant from the tumor site) were obtained from patients with cervical cancer who underwent surgical resection at the Shanghai Seventh People's Hospital. None of these patients had received chemotherapy or radiotherapy before sampling. Individuals with autoimmune disease, infectious diseases, and multi-primary cancers were excluded.

Blood from healthy donors was used for control experiments. The clinical stages of tumors were determined according to the TNM classification system of the International Union Against Cancer.

Isolation and stimulation of tumor cells

Fresh tissues were washed 3 times with Hank's solution containing 1% fetal calf serum (FCS) before being cut into small pieces. The specimens were then placed in RPMI 1640 medium containing 1 mg/mL collagenase IV and 10 mg/mL deoxyribonuclease I and mechanically dissociated using a gentle MACS Dissociator (Miltenyi Biotec, Bergisch Gladbach, Germany). Dissociated cell suspensions were further incubated for 1 h at 37 °C under continuous rotation. The cell suspensions were then filtered through a 70 µm cell strainer (BD Labware, Bedford, MA, United States). Then, part of cells was used for flow cytometry to detect the number of Tc17 cells, and the remaining cells were cultured with IL-17 (0-10 ng/mL), IL-22 (0-10 ng/mL), interferon (IFN)-γ (0-10 ng/mL), IL-17 plus IL-22, or 100% Tc17 cell-polarizing culture for 48 h. The culture supernatants were harvested for ELISA.

In vitro monocyte-T-cell co-culture system

Peripheral blood mononuclear cells from cervical cancer patients were isolated by Ficoll density gradient centrifugation. Fresh peripheral blood CD8⁺ T cells were selected using positive isolation and negative isolation kits, respectively. In a 5-d incubation period, bead-puriWed peripheral CD8⁺ T cells were co-cultured with autologous blood monocytes at a 2:1 ratio in the presence or absence of recombinant human IL-6 (10 ng/mL), IL-1β (10 ng/mL) and IL-23 (10 ng/mL) in 200 µL RPMI 1640 medium supplemented with 10% FCS. After 5-d incubation, the supernatants were harvested for ELISA and the cells for intracellular cytokine staining.

Flow cytometry analysis

For detection of intracellular molecules, T lymphocytes were stimulated for 5 h with phorbol myristate acetate (50 ng/mL) plus ionomycin (1 µg/mL) in the presence of GolgiStop (BD Pharmingen, San Diego, CA, United States). Intracellular cytokine staining was performed after fixation and permeabilization using Perm/Wash solution (BD Pharmingen). The lymphocytes were analyzed by multicolor flow cytometry with FACSCanto II (BD Biosciences). Data were analyzed with FlowJo software (TreeStar, Ashland, OR, United States) or FACSDiva software (BD Biosciences).

Immunohistochemistry assay

Paraformaldehyde-fixed and paraffin-embedded samples were cut into 5 µm sections, which were incubated with rabbit anti-IL-17 antibody and stained by horseradish peroxidase anti-rabbit immunoglobulin G followed by diaminobenzidine. The sections were then incubated with mouse anti-CD8 antibody and stained using EnVision G2 System/AP Rabbit/Mouse (Permanent Red) (Dako, Glostrup, Denmark) and subsequently counterstained with hematoxylin. Slides were examined using an Olympus 71 inverted fluorescence microscope.

Immunofluorescence

Immunofluorescence was performed as previously described with minor modifications^[10]. In detail, Paraformaldehyde-Waxed cryostat sections of tumor tissues were washed in PBS and blocked for 30 min with 20% rabbit serum in PBS. Sections were incubated with goat anti-human IL-17 antibody (Ab) diluted in 5% rabbit serum. The bound Ab was detected with FITC-conjugated rabbit anti-goat Ab. After washing with PBS, the sections were blocked for 30 min with 20% goat serum in PBS and incubated with mouse anti-human CD8 Ab diluted in 5% goat serum. The bound Ab was detected with TRITC-conjugated goat anti-mouse Ab. After washing with PBS, the slides were examined with an Olympus 71 inverted fluorescence microscope.

ELISA

Cervical cancer tissues or their matched adjacent normal tissues were homogenized in 1 mL Protein Extraction Reagent (Rockford, IL, United States). Concentrations of IL-6, IL-1β, and IL-23 in the tissue supernatants; concentrations of CXCL12 in the co-culture supernatants or tissue supernatants; and concentrations of IL-17 in the co-culture supernatants were determined using ELISA kits according to the manufacturer's instructions.

Statistical analysis

Comparisons between data groups were performed as stated in each figure legend

with the use of GraphPad Prism ver.6. A value of $P < 0.05$ was considered statistically significant. The resulting data were presented as mean \pm SE.

RESULTS

Tc17 cells specifically accumulate in tumor tissues of cervical cancer patients

To assess the status of Tc17 cells in human cervical cancer tissues, we isolated immune cells from cancer tissues, matched adjacent normal tissues as well as peripheral blood. Compared with healthy donors, the percentage of Tc17 cells in the peripheral blood from patients with cervical cancers was identical (Figure 1A). Of note, we found that Tc17 cells were selectively induced in tumors compared to their matched adjacent normal tissues (Figure 1A). To further confirm this result, we analyzed the distribution of Tc17 cells in the paracancerous stroma, carcinoma nets, and intra-tumor sites. The results showed that Tc17 cells accumulated in all these sites, especially in carcinoma nets and intra-tumor sites (Figure 1B-D). Thus, these data show that Tc17 cells have an oncogenic role in cervical cancer development.

Cervical cancer cells produce a greater amount of Tc17-polarizing cytokines

Previous studies revealed that IL-6, IL-1 β , and IL-23 are essential for Tc17 cell differentiation^[11,12]. To investigate the mechanism of cervical cancer following modulation of Tc17 cell development, we assessed the levels of Tc17-polarizing cytokines in cancer-associated tissues. As expected, the concentrations of IL-6, IL-1 β , and IL-23 were significantly increased in peritumoral and intra-tumor tissue relative to their matched normal tissues or peripheral blood (Figure 2A-C). Thus, these data indicate that tumor-derived cytokines play a stimulatory role in the modulation of Tc17 polarization.

IL-6, IL-1 β , and IL-23 acts synergistically to enhance Tc17 cell differentiation

To evaluate the potential role of these cytokines in Tc17 cell differentiation, we isolated peripheral blood CD8⁺ T cells and autologous blood monocytes of cervical cancer patients. After 5-day co-culture incubation, the supernatants were harvested for ELISA and the cells for intracellular cytokine staining. The results showed that the addition of exogenous IL-6, IL-1 β , and IL-23 significantly increased the frequency of Tc17 cells either alone or in combination compared with co-culture without any cytokines added (Figure 3A and B). In addition, the ELISA results showed that IL-17 production was consistent with the percentage of Tc17 cells (Figure 3C). These findings showed that IL-6, IL-1 β , and IL-23 act synergistically to induce Tc17 cell polarization *in vitro* and suggest that a similar process might operate *in vivo*.

Tc17 cell-derived IL-17 promotes CXCL12 expression in tumor cells

To investigate the biological function of Tc17 cells in cervical cancer development, we isolated primary tumor cells and stimulated them with various concentrations of IL-17, IL-22, and IFN- γ , and the production of CXCL12 was detected in the culture supernatants by ELISA. As shown in Figure 4A, CXCL12 expression in cancer cells was increased upon IL-17 stimulation, while IL-22 or IFN- γ exhibited little effects on CXCL12 upregulation in cancer cells (Figure 4A). Moreover, unlike the co-culture supernatants from blood or non-tumor tissue monocytes, TAM-derived Tc17 cell culture supernatants selectively induced CXCL12 production in cancer cells (Figure 4B). The effects of CXCL12 upregulation were attenuated by anti-IL-17 neutralizing antibody, but not anti-IL-22 neutralizing antibody (Figure 4B). These findings suggested that Tc17 cell-derived IL-17 induced chemokine CXCL12 production by tumor cells.

Tumor-infiltrating Tc17 cells are related to the poor outcome of patients with cervical cancer

As CXCL12 upregulation drives cancer development, we assessed the relationship between the number of Tc17 cells and the survival rate of patients with cervical cancer. We first analyzed the number of Tc17 cells per million total cells in each tissue from patients in different TNM stages. The results showed that as cancer progressed, the number of Tc17 cells significantly increased in each of the tested tissues (Figure 5A). Conversely, patient survival was significantly reduced (Figure 5A and B). Simultaneously, we analyzed Tc17 cell percentage within the total CD8⁺ T cells in the same samples, and as cancer progressed, we found that the percentage of Tc17 cells significantly increased in the tested tissues, and was negatively associated with patient survival (Figure 5C and D). These findings suggested that the number of Tc17 cells was related to the survival rate of patients, thereby predicting the clinical

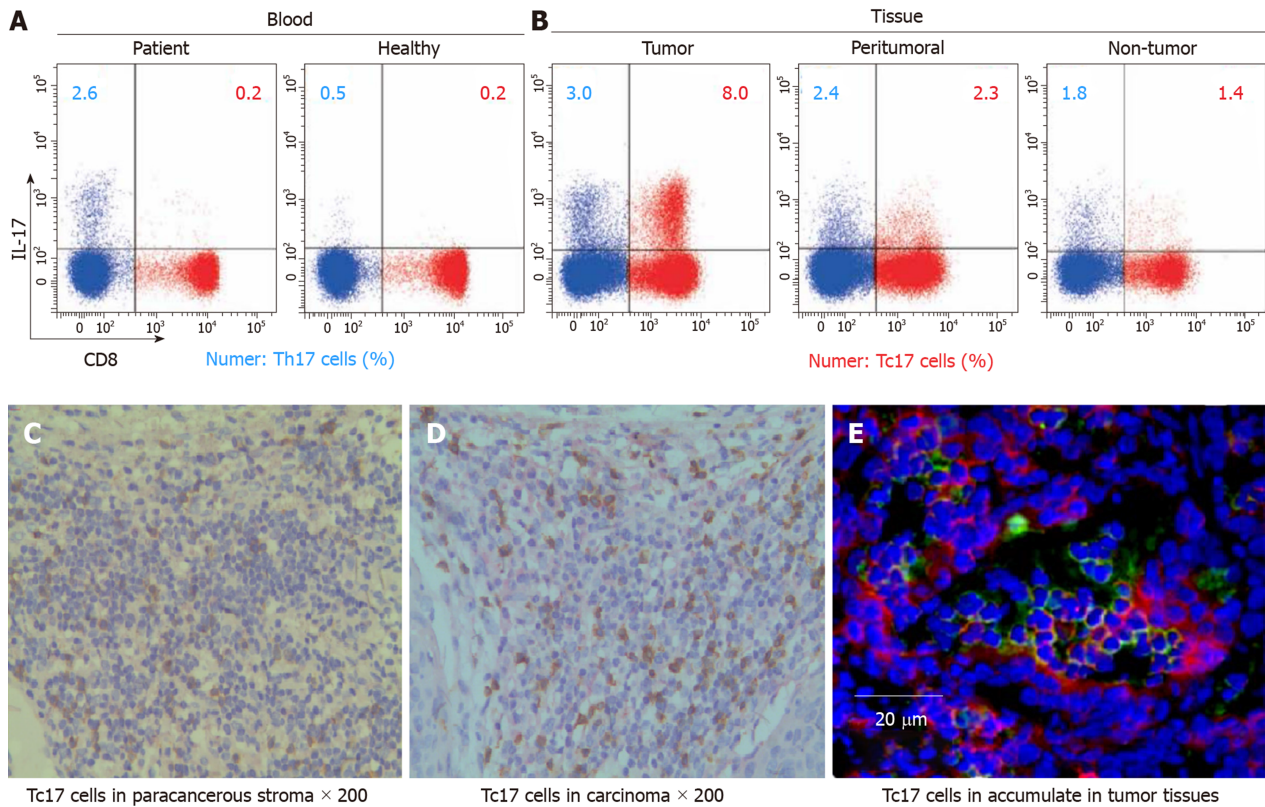


Figure 1 Tc17 cells accumulate in tumor tissues of cervical cancer patients. A, B: Dot plots of intracellular cytokine staining for Tc17 and Th17 in tumor cells; C: Immunohistochemistry staining for interleukin (IL)-17⁺ Tc17 cells in paracancerous stroma, the brown signal represents staining of IL-17, and the red signal represents staining of Tc17 (Envision, ×200); D: Immunohistochemistry staining for IL-17⁺ Tc17 cells in carcinoma nets, the brown signal represents staining of IL-17, and the red signal represents staining of Tc17 (Envision, ×200); E: Immunofluorescence staining for intra-tumoral IL-17⁺ Tc17 cells, the green signal represents staining of IL-17, the red signal represents staining of Tc17, and the blue signal represents DAPI-stained nuclei (scale bar, 20 μm). IL: Interleukin.

outcome of patients with cervical cancer.

DISCUSSION

Although Tc17 cell-mediated immune regulation has been studied in tumor progression in both mice and black-and-white monkeys^[13], the mechanism of Tc17 cells in human cervical cancer has been rarely reported. In this study, we found that Tc17 cells play a stimulatory role in the progression of cervical cancer. Compared with normal tissues, cervical cancer cells produced a greater amount of IL-6, IL-1β, and IL-23A, which are essential for Tc17 cell differentiation^[11,12]. Tc17 induction in turn augmented CXCL12 expression in tumor cells *via* activation of IL-17 signaling. Furthermore, the increased ratio of Tc17 cells in tumors predicted a poor prognosis in patients with cervical cancers. Thus, these data demonstrate that Tc17 cells have an oncogenic role in cervical cancer development.

Recent studies have shown that persistent antigenic stimulation leads to CD8⁺ T cell exhaustion^[14,15]. Blockade of the PD-1/PD-L1 strategy is promising for the treatment of cancer^[16]. However, an increasing number of studies have revealed that tumor-elicited inflammation enhances resistance to cancer immunotherapy, which suggests that cancer cells adopt other strategies to escape immune attack. In the present study, we found that Tc17 cells specifically accumulated in cervical cancer tissue. Unlike the necessary role of IL-6 and IFN-γ in stimulation of Tc17 cell expansion in colorectal or liver cancers^[14,15], cervical cancer cells secrete abundant Tc17-polarizing cytokines including IL-6, IL-1β, and IL-23, suggesting the enhanced effects on Tc17 cell development is attributed to the cancer microenvironment.

Murine tumor models have shown that Tc17 cells impair immune surveillance and promote *de novo* carcinogenesis and neovascularization of tumors^[16,17]. However, other groups have identified anti-tumor activity of Tc17 cells in mice^[18,19]. Therefore, the precise underlying mechanism is still unclear. Our data revealed that increased IL-17 derived from Tc17 cells drives CXCL12 expression in tumor cells, which has been reported to enhance cell proliferation and migration, thereby exacerbating malignant

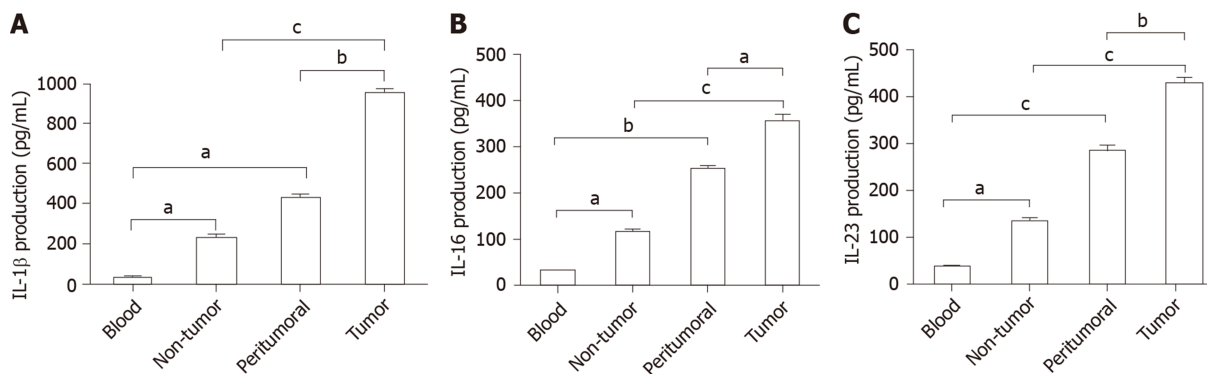


Figure 2 Interleukin-6, interleukin-1 β , and interleukin-23 accumulated in tumor tissues of cervical cancer patients. A-C: Before cell suspension culture, ELISA was carried out to show that interleukin (IL)-6, IL-1 β , and IL-23 were significantly upregulated in the supernatants of tumor and peritumoral tumor tissues compared with the supernatants isolated from autologous non-tumor tissues or peripheral blood. Comparisons were performed using the *t*-test. ^a*P* < 0.05; ^b*P* < 0.01. ^c*P* < 0.001. Error bars represent SE. IL: Interleukin.

formation. Finally, we assessed the relationship between the number of Tc17 cells and the survival rate of patients with cervical cancer. Similar studies have been conducted in mice and gastric cancer patients^[20]. The results showed that the increased ratio of Tc17 in tumors is highly correlated with poor outcome in patients with cervical cancer.

In conclusion, our data demonstrate that Tc17 cells are specifically induced by the cervical cancer microenvironment. Tc17 cells promote tumor development *via* activation of IL-17 signaling and are associated with the prognosis of patients with cervical cancer. Thus, we propose that Tc17 cells can be used as a useful clinical index in the prognosis of cervical cancer.

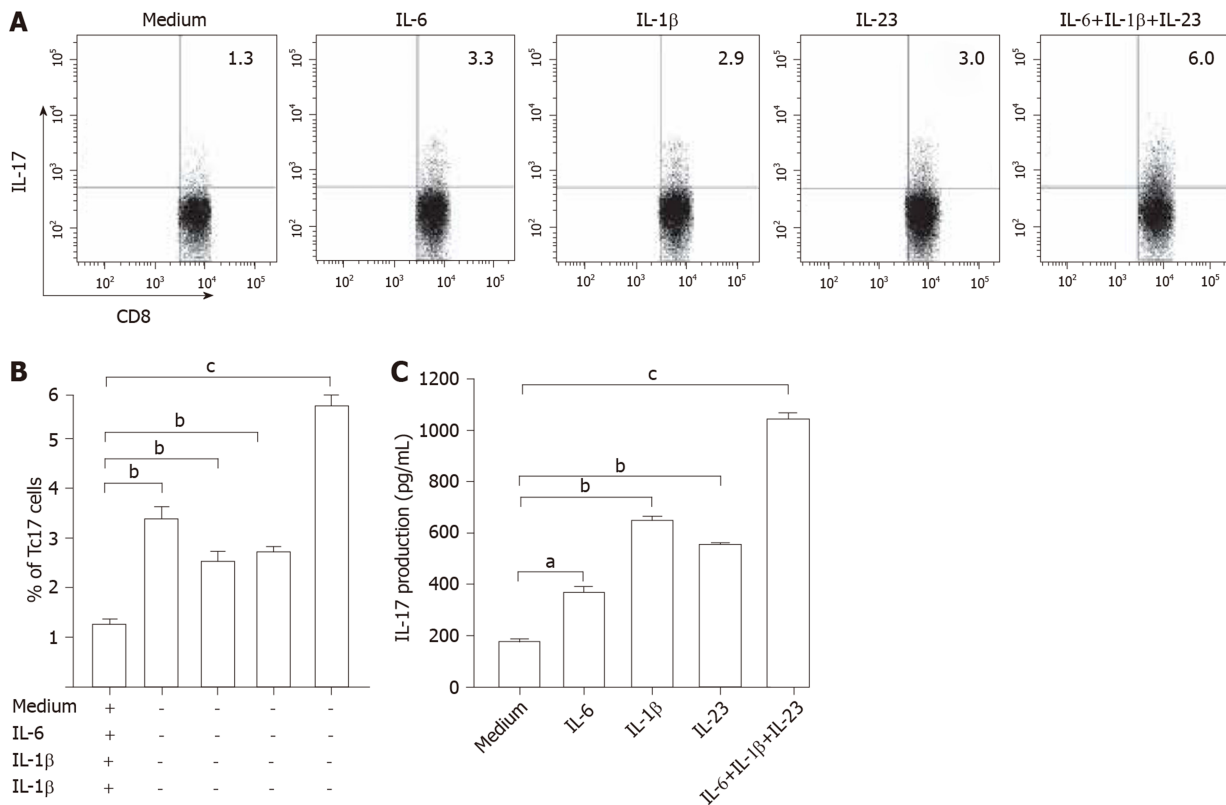


Figure 3 Interleukin-6, interleukin-1 β , and interleukin-23 induced differentiation of Tc17 cells. A-C: Peripheral CD8⁺ T cells and blood monocytes were co-cultured as described in Materials and Methods. Representative data and statistical analysis of Tc17 cell percentage in CD8⁺ T cells and interleukin-17 production in the culture supernatants from the co-culture systems. Comparisons were performed using the *t*-test. ^a*P* < 0.05, ^b*P* < 0.01, ^c*P* < 0.001. Error bars represent SE. IL: Interleukin.

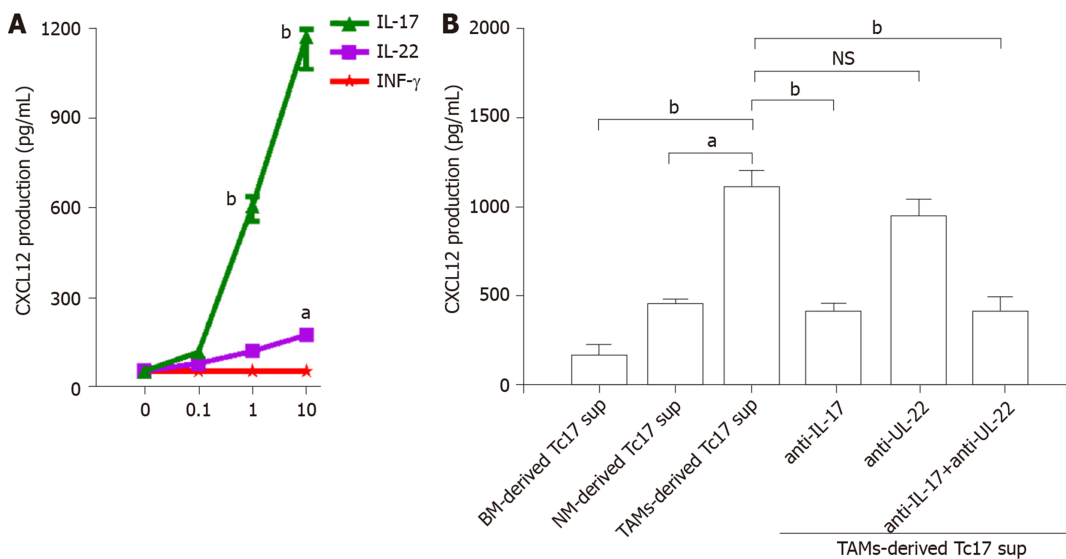


Figure 4 Tc17 cell-derived interleukin-17 induces tumor cells to produce CXCL12. A: Primary tumor cells were cultured with various concentrations of interleukin (IL)-17, IL-22, interferon- γ , and the production of CXCL12 was detected in the culture supernatants by ELISA and statistically analyzed; B: Primary tumor cells were cultured with various concentrations of cytokines or monocyte-derived Tc17 cells culture supernatants, and the production of CXCL12 was detected in the culture supernatants by ELISA and statistically analyzed. Comparisons were performed using the *t*-test. ^a*P* < 0.05; ^b*P* < 0.01. Error bars represent SE. NS: No significant difference; NM: Non-tumor tissue monocyte-derived; BM: Blood monocyte-derived; TAMs: Tumor tissue monocyte-derived.

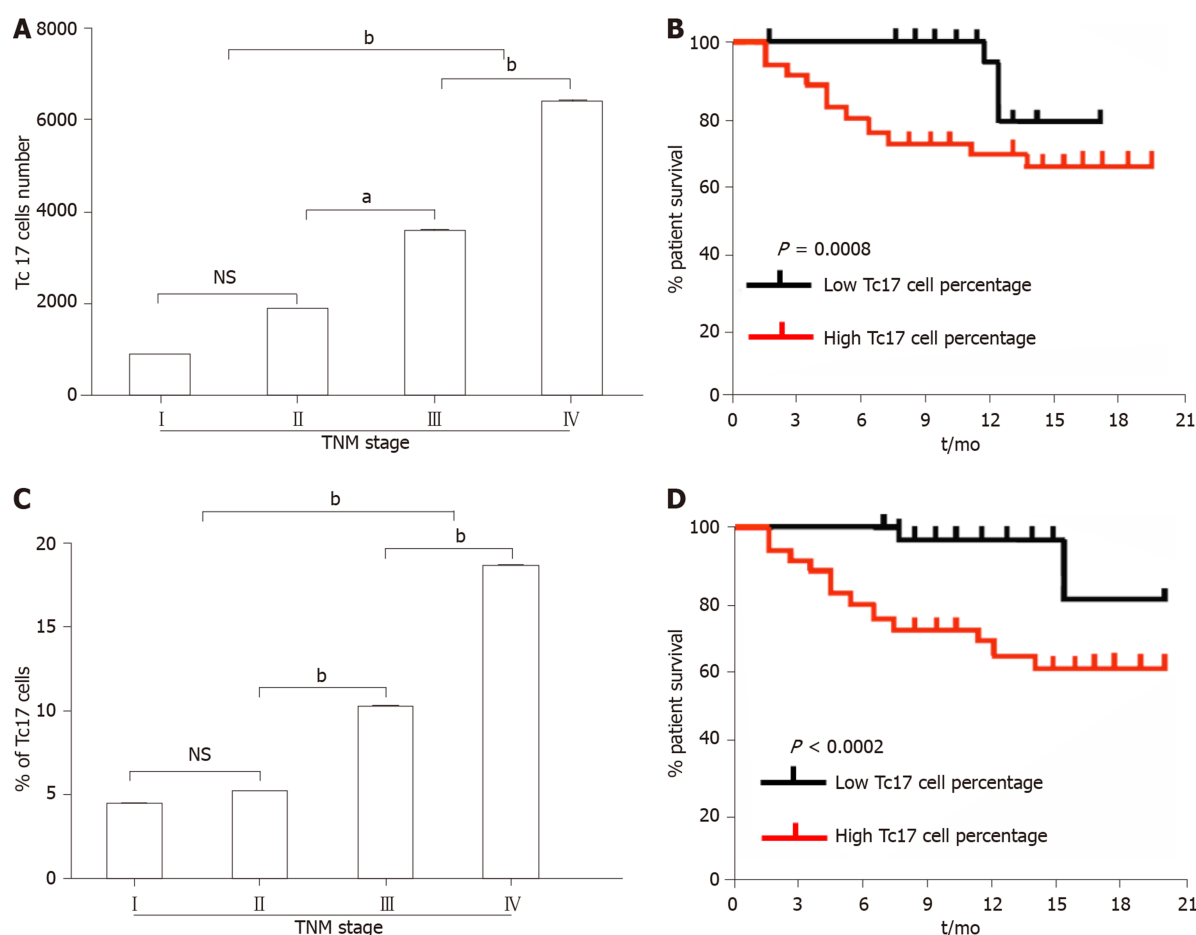


Figure 5 Analysis of the correlation between the percentage of Tc17 cells and the survival rate of cervical cancer patients. A: The total number of Tc17 cells per million total cells in each tissue in different TNM stage; B: The association between Tc17 cell number and patient survival times; C: Tc17 cell percentage in CD8⁺ T cells in each stage of cancer progression; D: Survival significantly decreased as a function of Tc17 cell percentage. Comparisons were performed using the *t*-test. NS indicates no significant difference; ^aP < 0.05; ^bP < 0.01. Error bars represent SE.

ARTICLE HIGHLIGHTS

Research background

The existence of Tc17 cells was recently shown in several types of inflammatory diseases.

Research motivation

The distribution and functions of Tc17 cells in cervical cancer have not been fully elucidated.

Research objectives

To investigate the role of Tc17 cells in the pathogenesis of cervical cancer.

Research methods

The frequency of Tc17 cells in blood and tumor samples from patients with cervical cancer was determined by flow cytometry. In addition, the levels and phenotype of Tc17 cells in tissue samples from cervical cancer patients were assessed by immunohistochemistry staining.

Research results

Tc17 cells specifically accumulate in the tumor tissues of cervical cancer patients. Cervical cancer-elicited inflammation increases Tc17-polarizing cytokine production, which attenuates cytotoxic CD8⁺ T cell development. High interleukin-17 production by Tc17 cells leads to CXCL12 upregulation and cancer cell migration.

Research conclusions

Consistent with the oncogenic role of Tc17 cells in cancer development, the ratio of cancer-infiltrating Tc17 cells is highly associated with poor prognosis in patients with cervical cancer.

Research perspectives

This study indicates that Tc17 cells in cervical cancer and their regulatory mechanisms are associated with cancer progression.

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