



Interleukin-17 inhibits the growth of oral squamous cell carcinoma by promoting the differentiation of T helper 17 cells

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Background: The infiltration of T helper 17 (Th17) cells has been reported in multiple solid tumors. Recent studies implicated that Th17 cells played an inhibitory role in the tumor growth. However, if Th17 cells infiltrated and played a role in oral squamous cell carcinoma (OSCC) remained unknown.

Methods: Fluorescence activated cell sorting (FACS) was performed to confirm the increase of Th17 cells in the peripheral blood and the infiltration in tumor samples from OSCC patients. To mimicking the effect of Th17 cells on OSCC growth, an OSCC line stably expressing human interleukin-17A (IL-17A) was established. Human-PBL-SCID-beige mice were employed as the hosts for the implantation of OSCC/IL-17A cells. Quantitative PCR, ELISA and Western blotting were performed to analyze Th17 differentiation and function in the implanted tumors.

Results: The percentage of Th17 cells increased significantly in both the peripheral blood and the tumor tissues from OSCC patients. Expression of IL-17A promoted the differentiation of Th17 cells and suppressed the growth of implanted OSCC tumors. The transcription as well as the protein levels of IL-6 and granzyme B in the implanted OSCC cells was remarkably elevated by IL-17A expression.

Conclusions: The increased infiltration of Th17 cells was suggested to result in a therapeutic effect on OSCC as Th17 cells did in other tumors. IL-17 could promote the differentiation of Th17 cells by increasing IL-6 expression. The inhibitory effect of Th17 cells on OSCC growth was associated with their increased cytotoxic activity.

Keywords: T helper 17 (Th17); oral squamous cell carcinoma (OSCC); interleukin-17; tumor implantation; cytokines

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Introduction

The tumors in head and neck are usually characterized by the progressive malignancy and occupy 6% of the newly diagnosed neoplasms (1). Approximately, 95% cases of the head and neck tumors are squamous cell carcinoma (2,3). Even taking the advancements in surgery, radiology and

chemotherapy into account, the prognosis of the squamous cell carcinoma is still pessimistic because of the terribly high ratio of relapse (4). The infiltration of lymphocytes in head and neck tumors has been reported in many studies (5,6), however, the exact roles of the tumor infiltrated lymphocytes (TIL) were still unclear. Moreover, since previous studies mainly focused on nasopharyngeal cancers,

if the infiltration of TIL exists in oral squamous cell carcinoma (OSCC) is also in debated.

T helper 17 (Th17) cell is a IL-17⁺CD4⁺ subpopulation of T helper cells discovered recently (7). Th17 cell plays critical roles in mucosal immunity and autoimmunity, such as multiple sclerosis, immune enteritis and psoriasis (7). Besides interleukin-17A (IL-17A), Th17 cells also secrete IL-17F, IL-21, IL-22 and IL-26, and are marked by CD-161, CCR6, IL-23R and ROR2C (8,9). IL-17 induces adjacent cells to produce chemokines, matrix metalloproteinases and anti-microbial peptides, which trigger inflammation and recruit neutrophils and macrophages (10). Previous studies confirmed the infiltration of Th17 cells in the tumors of human and mouse (11). Under the combined stimulus from TGF- β and IL-6, the primitive CD4⁺ T cells preferentially differentiate into Th17 cells, and IL-23 plays a key role in the maintenance and proliferation of Th17 cells (12).

Although the mechanism remains unknown, the inhibitory effects of Th17 cells on tumor have been reported (13-15). Since the complicated immune-escape mechanism is involved in the tumor inhibition, if Th17 cells serve as a kind of TIL and exert an inhibitory effects on OSCC as they do in other tumors remain elusive. In this study, we detect an aberrant increase and infiltration of Th17 cells in OSCC patients, and then, assess the influence of Th17 cells on OSCC by enhancing their differentiation.

Methods

The diagram of the experimental procedures could be found in *Figure S1*.

Tissue specimens

The blood and tissue specimens were taken from the OSCC patients who got OSCC excision in Chinese PLA General Hospital from May 2016 to May 2017. All OSCC patients were diagnosed as OSCC according to the pathological biopsy. There were 8 OSCC patients donating their blood samples and OSCC tissues. The normal control tissues were collected from the neighbouring tissues at least 0.5 cm away from the OSCC focus. The control blood specimens were contributed by the health donors in the Infusion Center of Chinese PLA General Hospital. All the patients and healthy donors had no history of radiotherapy, chemotherapy, biotherapy and traditional Chinese medicine treatments before specimen collection (*Table 1*).

Fluorescence activated cell sorting (FACS)

FACS was conducted to detect the percentage of the Th17 cells in all T lymphocytes in the circulating blood and OSCC tissues. The antibodies for cell sorting were mouse anti-human IL-17A Alexa Fluor[®] 647 (Becton, Dickinson and Company, USA) and PerCP mouse anti-human CD4 (Becton, Dickinson and Company, USA). ALEXA 647 mouse IgG1 was used as the isotype control (Becton, Dickinson and Company, USA). The cell sorting was performed and analyzed by flow cytometer (BD FACS Calibur[™]) as the manufacturer instructed (Becton, Dickinson and Company, USA).

Cell lines

TCA8113, an OSCC line from human tongue squamous cell carcinoma was preserved by our institute. The vector expressing human IL-17A was purchased from KainuoBio Co., Ltd. (Beijing, China) by inserting the coding sequence of human IL-17A (480bp) into the backbone of pcDNA3.1. The transfection of TCA8113 with the plasmid pcDNA3.1/IL-17A, and the subsequent screen by G418 were performed as the manufacturer's protocol (Invitrogen, Carlsbad, CA). The surviving cells (TCA8113/pcDNA3.1/IL-17A) were assessed by ELISA Kit for IL-17A production (U-Cytech, Dutch) and CellTiter 96[®] Aqueous MTS Kit for cell proliferation (Promega, Madison, WI, USA).

Host mice

The human lymphocytes were extracted from the circulating blood of health donors by Ficoll-Paque density gradient centrifugation as previous described (16). The concentrated lymphocytes were injected at the 3×10^7 cells/mouse into the abdominal cavity of the 4-week-old SCID-beige mouse (average weight is 20–25 g; purchased from Beijing Vital River Laboratory Animal Technology Co., Ltd., Beijing, China) to generate human-PBL-SCID-beige mice. After 4 weeks of injection, the serum was taken from tail vein for human IgG calibration with the human IgG calibration Kit (MabTech).

OSCC xenograft tumors

TCA8113/pcDNA3.1-IL-17A, TCA8113/pcDNA3.1 and TCA8113 cells (2.5×10^6 cells/mouse) were, respectively, subcutaneously injected into the left flank of the bodies of the 5-week-old human-PBL-SCID-beige mice to form

Table 1 General information of patients (P) and donors (D)

No.	Age (years)	Gender	Smoking	Drink	Cancer	Other disease
P1	36	Male	Yes	No	No	No
P2	61	Male	Yes	No	No	No
P3	57	Female	No	No	No	No
P4	53	Male	No	Yes	No	No
P5	35	Female	No	No	No	No
P6	43	Male	No	No	No	No
P7	46	Male	No	No	No	No
P8	37	Female	No	No	No	No
D1	50	Female	No	No	No	No
D2	41	Male	Yes	No	No	No
D3	30	Female	No	No	No	No
D4	34	Female	No	No	No	No
D5	38	Male	No	No	No	No
D6	39	Male	Yes	No	No	No
D7	42	Male	No	No	No	No
D8	44	Male	No	No	No	No

xenograft tumors. 27 host mice were randomly assigned into 3 groups. After mice were sacrificed by cervical dislocation, tumors were harvested for following investigations.

Quantitative PCR (Q-PCR)

Total RNA was extracted from implanted OSCC samples by using trizol total RNA reagent (Invitrogen, Carlsbad, CA). Primers were synthesized by Ao Ke Inc. (Beijing, China), and their sequences are shown as follow: human granzyme B (forward: 5'-GGTGGCTTCCTGATACGA-3' and reverse: 5'-GGATGGGTCTT TTCACAG-3'); human IL-1 β (forward: 5'-TACAGTGGCAATGAGGAT-3' and reverse: 5'-AACCAGCATC TTCCTCAG-3'); human IL-6 (forward: 5'-CCACACAGACAGCCACTC-3' and reverse: 5'-TTTGGAAGG TTCAGGTTG-3'); human IL-17A (forward: 5'-TACTACAACCGATCCACCT-3' and reverse: 5'-GGACCA GGATCTCTTGCT-3'); human IL-23A (forward: 5'-ATGATGTTCCCCATATCC-3' and reverse: 5'-TATCA GGGAGCAGAGAGG-3'); human ROR γ t (forward: 5'-CTTGGGGAAGTGGGACAG-3' and reverse: 5'-GGAGAAGATGTTGGAGCG-3'); human GAPDH (forward: 5'-ATCATCAGCAATGCCTCC-3'

and reverse: 5'-AGTCCTTCCACGATACCAA-3'). All the amplifications were conducted in MiniOpticon™ Real-Time PCR system (Bio-Rad Laboratories, USA).

Western blotting

The procedures for the lysis of the implanted OSCC sample and the manipulation of Western blotting followed those procedures described in previous study (17). The antibodies against human granzyme B (Santa Cruz Biotechnology, Inc. USA), and ROR γ t (Merck Millipore, Darmstadt, Germany) were mouse anti-human IgG, while the antibody against human Actin was rabbit IgG (Santa Cruz Biotechnology, Inc. USA). The secondary antibodies conjugated with HRP were goat anti-mouse and goat anti-rabbit IgGs (Santa Cruz Biotechnology, Inc. USA). The ECL chemiluminescent (Thermo Fisher Scientific, USA) was applied for color development. The image analysis was performed with Gel-Pro analyzer4.5 (Media Cybernetics).

ELISA

The ELISA Kits for human IL-17A (U-CyTech), IL-1 β

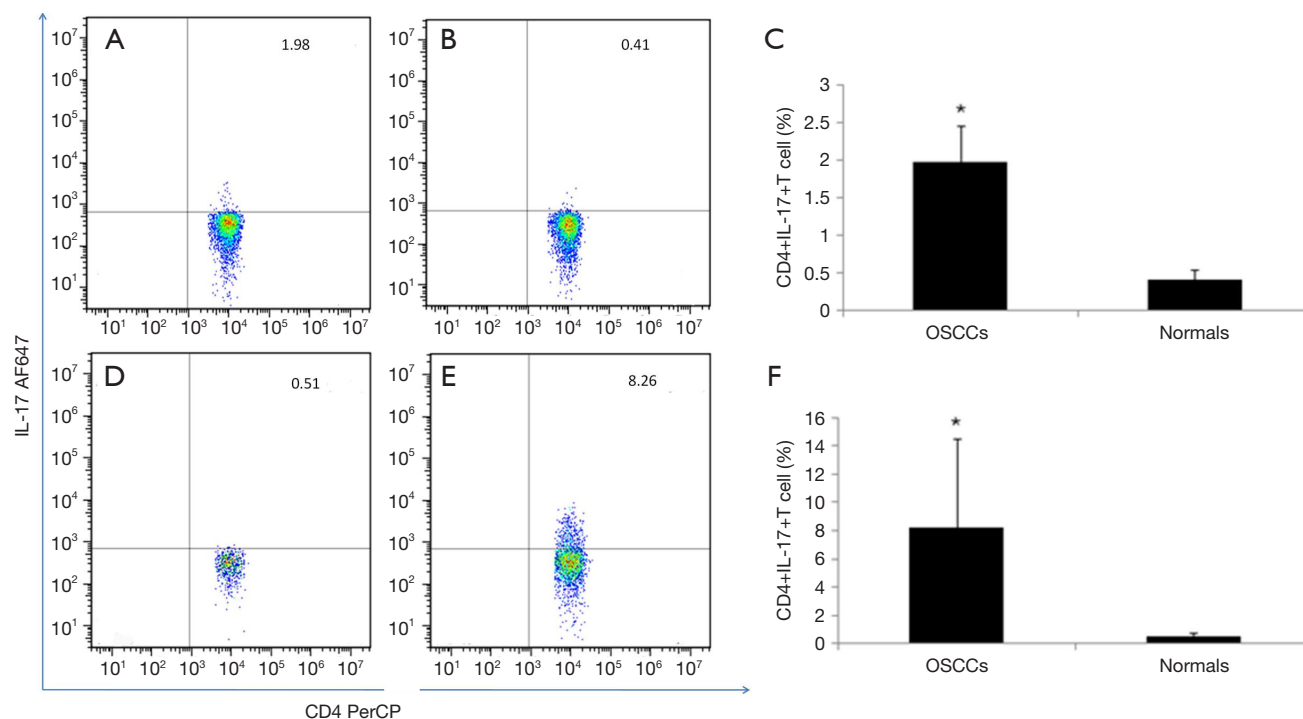


Figure 1 The content of Th17 cells in the OSCC blood and tissues by flow cytometry analysis. (A) Peripheral blood from OSCC patients; (B) peripheral blood from normal donors; (C) statistical analyses for the Th17 percentage in blood samples; (D) normal adjacent tissues from the OSCC focus; (E) tumor tissues from OSCC patients; (F) statistical analyses for the Th17 percentage in tissue specimens. *, $P < 0.05$.

(MabTech), IL-6 (GenWay) and IL-23 (CUSABIO) were purchased from Dakewe Biotech Co., Ltd. (Shenzhen, China). The ELISA manipulation followed the procedures provided by the manufacturers.

Statistical analysis

The data were analyzed using SPSS 17.0 software (IBM, NY, USA). The SNK test and the Student's *t*-test were used in this study. $P < 0.05$ was regarded as a statistically significant difference.

Results

The differentiation and infiltration of Th17 cells in OSCC patients were promoted

To address if there was an infiltration of Th17 cells in OSCC samples, FACS assay was employed to enrich Th17 lymphocytes by the antibodies against CD4 and IL-17. The percentage of CD4⁺IL-17⁺ cells in the total CD4⁺ cells, which represent differentiated Th17 lymphocytes,

was $1.98\% \pm 0.472\%$ in the peripheral blood of the OSCC patients (Figure 1A), while the counterpart population was only $0.41\% \pm 0.14\%$ in the control blood ($P < 0.01$; Figure 1B,C). It suggested that the differentiation of Th17 cells in the OSCC patients was significantly promoted. On the other hand, $8.259\% \pm 6.25\%$ cells of the CD4⁺ population from OSCC tissues was IL-17 positive (Figure 1D). In contrast, IL-17⁺ cells was only $0.51\% \pm 0.22\%$ in all the CD4⁺ cells in the normal tissues ($P < 0.05$; Figure 1E,F). This result suggested that the infiltration of Th17 cells took place in the OSCC tissues.

The construction of IL-17A-expressing cell line and tumor-implantation hosts

The eukaryotic expression vector (pcDNA3.1-IL-17A) was constructed by inserting the full length cDNA of human IL-17A (480 bp) into the backbone of pcDNA3.1 (Figure 2A,B). To get an OSCC line stably expressing IL-17A, the pcDNA3.1-IL-17A was transfected into TCA8113 cells (human tongue squamous cell carcinoma). After the screen by G418, the TCA8113 cells stably expressing

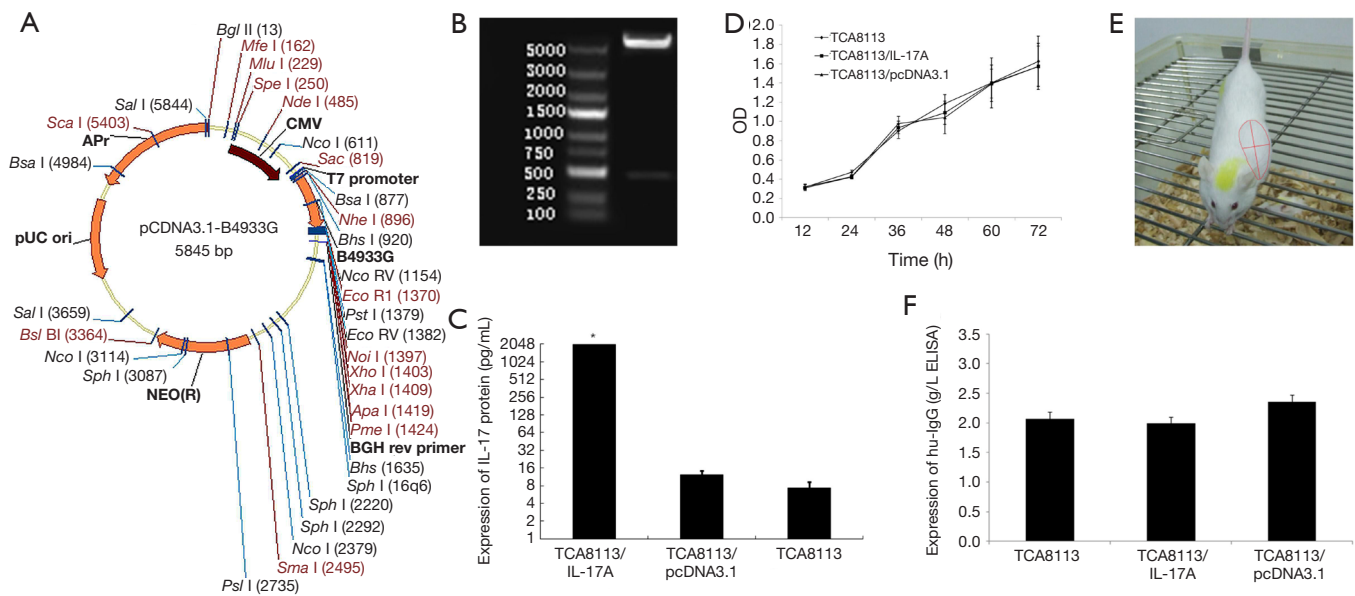


Figure 2 The validity of human-PBL-SCID-beige mice and IL-17A-expressing TCA8113 cell line. (A) The map of pcDNA3.1 vector; (B) identification of pcDNA3.1-IL-17 by NheI/EcoRI cutting; (C) IL-17A production in TCA8113, TCA8113/pcDNA3.1-IL-17A and TCA8113/pcDNA3.1 cells by ELISA assay; (D) cell proliferation of TCA8113, TCA8113/pcDNA3.1-IL-17A and TCA8113/pcDNA3.1 cells by MTS assay; (E) the human- PBL-SCID-beige mouse carrying OSCC xenograft (outlined by red circle); (F) ELISA assay confirmed the human IgG production in human- PBL-SCID-beige mice showed no significant difference. *, $P < 0.05$.

IL-17A was named after TCA8113/pcDNA3.1-IL-17A. ELISA assay confirmed that the average production of IL-17A in the TCA8113/pcDNA3.1-IL-17A cell line reached 2,016.256 pg/mL, which was remarkably higher than those on control groups (Figure 2C). Meanwhile, MST test gave rise to the similar re-absorptions among TCA8113/pcDNA3.1-IL-17A, TCA8113/pcDNA3.1 and TCA8113 groups ($P > 0.05$; Figure 2D), indicating that the secretion of IL-17A exerted little effect on the proliferation of TCA8113/pcDNA3.1-IL-17A cells.

To mimicking the *in vivo* immune environment of human, human-PBL-SCID-beige mice, in which human immune system was reconstructed, were employed as the hosts for tumor implantation (Figure 2E). All the 27 human-PBL-SCID-beige mice were randomly separated into 3 groups. The average concentration of human IgG in the peripheral blood of the host mice was approximately 2.105 g/L and no significant difference among groups (Figure 2F).

IL-17A inhibited OSCC growth by promoting the differentiation of Th17

To assess the *in vivo* effects of IL-17A on OSCC growth,

we constructed OSCC implantation model by transplanting the TCA8113/pcDNA3.1-IL-17A cell mass containing 2.5×10^6 cells into human-PBL-SCID-beige mice. During the first 5–9 days after implantation, the tumor size in the TCA8113/pcDNA3.1-IL-17A group differed little from those in TCA8113/pcDNA3.1 and TCA8113 groups (Figure 3A). From 14 days after implantation on, the tumor sizes in the TCA8113/pcDNA3.1-IL-17A group were noticeably smaller than those in TCA8113/pcDNA3.1 and TCA8113 groups (Figure 3A). In contrast, the tumor sizes in TCA8113/pcDNA3.1 and TCA8113 groups showed few discrepancy from each other in all the period (Figure 3A). After 29 days of implantation, the final volume of the tumor in TCA8113/pcDNA3.1-IL-17A group reached $1.219 \pm 0.626 \text{ cm}^3$, which was remarkably smaller than the $2.858 \pm 1.188 \text{ cm}^3$ in TCA8113/pcDNA3.1 group or the $2.593 \pm 1.199 \text{ cm}^3$ in TCA8113 group (Figure 3B), indicating that the *in vivo* growth of OSCC was suppressed by the production of IL-17A.

To verify if IL-17A could enhance the differentiation of Th17, the expression of IL-17A and ROR γ t in the implanted tumors was measured. Q-PCR showed that the transcription of both IL-17A and ROR γ t was significantly

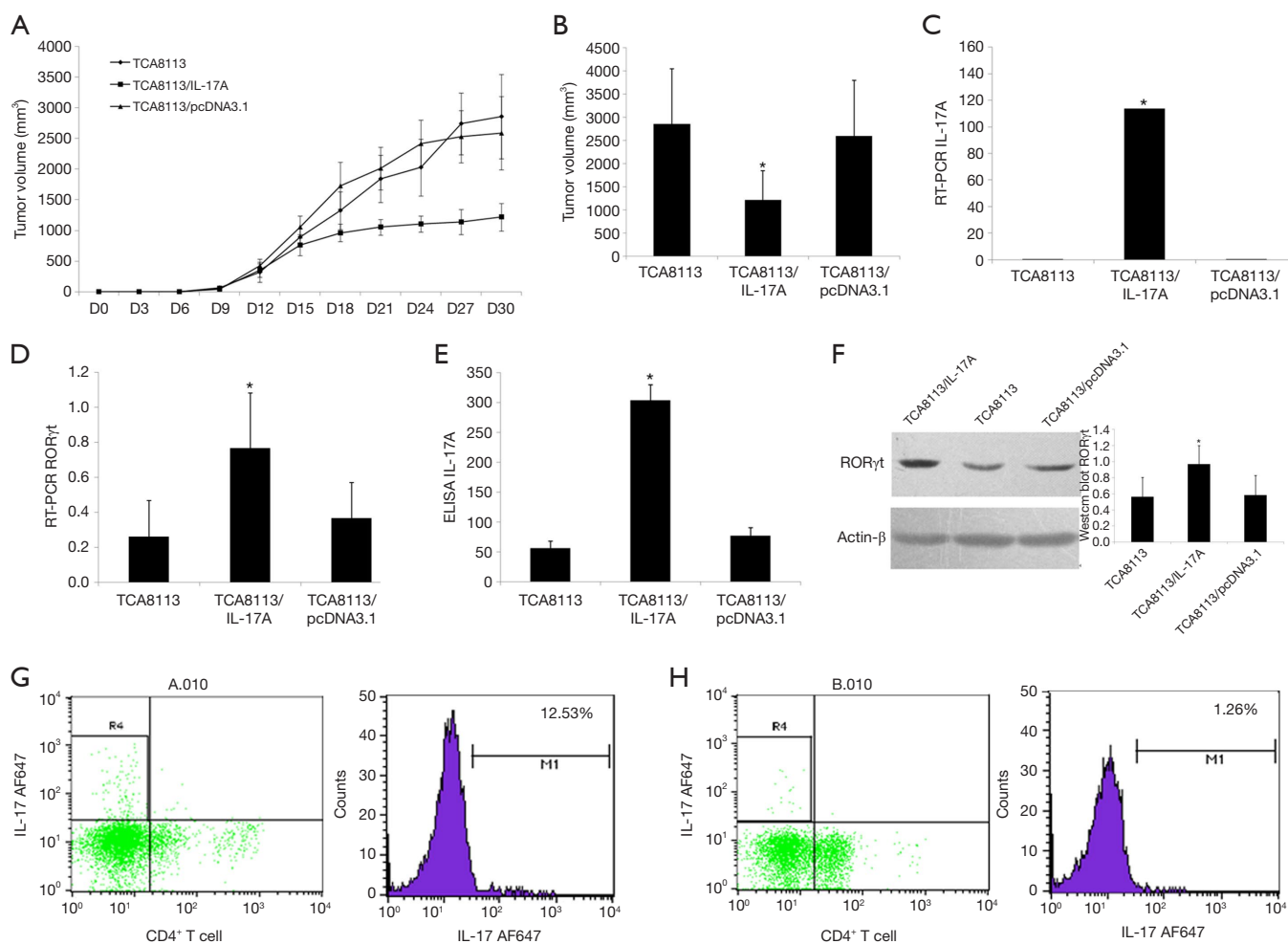


Figure 3 IL-17A impaired OSCC growth and promoted the differentiation of Th17 cells. (A) The growth curves of OSCC xenografts in 30 days after implantation; (B) the average tumor size in TCA8113/pcDNA3.1-IL-17A group was evidently smaller than those in control groups; (C) the IL-17A transcripts in TCA8113/pcDNA3.1-IL-17A group was much higher than those in control groups; (D) the mRNA of RORγt in TCA8113/pcDNA3.1-IL-17A group was increased compared with control groups; (E) the protein level of IL-17A in TCA8113/pcDNA3.1-IL-17A group was much higher than those in control groups; (F) the RORγt band of Western blotting in TCA8113/pcDNA3.1-IL-17A group was significantly denser than those of controls; (G) in a TCA8113/pcDNA3.1-IL-17A samples, the percentage of the Th17 positive cells in all CD4⁺ T lymphocytes was 12.53%; (H) the percentage of the Th17 positive cells in all CD4⁺ T lymphocytes was only 1.26% in a TCA8113/pcDNA3.1 sample. *, P<0.05.

increased in the tumors of TCA8113/pcDNA3.1-IL-17A group compared to TCA8113/pcDNA3.1 or TCA8113 group (Figure 3C,D). Consistently, ELISA assay indicated that the IL-17A in TCA8113/pcDNA3.1-IL-17A group was 5–6 folds higher than those in TCA8113/pcDNA3.1 and TCA8113 groups (Figure 3E). Western blotting also exhibited a much stronger band of RORγt protein in TCA8113/pcDNA3.1-IL-17A group compared to TCA8113/pcDNA3.1 and TCA8113 groups (Figure 3F). To

further confirm the enhanced differentiation of Th17 cells in the TCA8113/pcDNA3.1-IL-17A group, the percentage of Th17 cells in all CD4⁺ T lymphocytes was measured by FACS. In all the T lymphocytes screened by anti-CD4 antibody, the percentage of Th17 cells was 12.53% in the TCA8113/pcDNA3.1-IL-17A sample (Figure 3G), while 1.26% in the TCA8113/pcDNA3.1 sample (Figure 3H). These findings suggested that IL-17A could suppress OSCC growth by promoting Th17 differentiation.

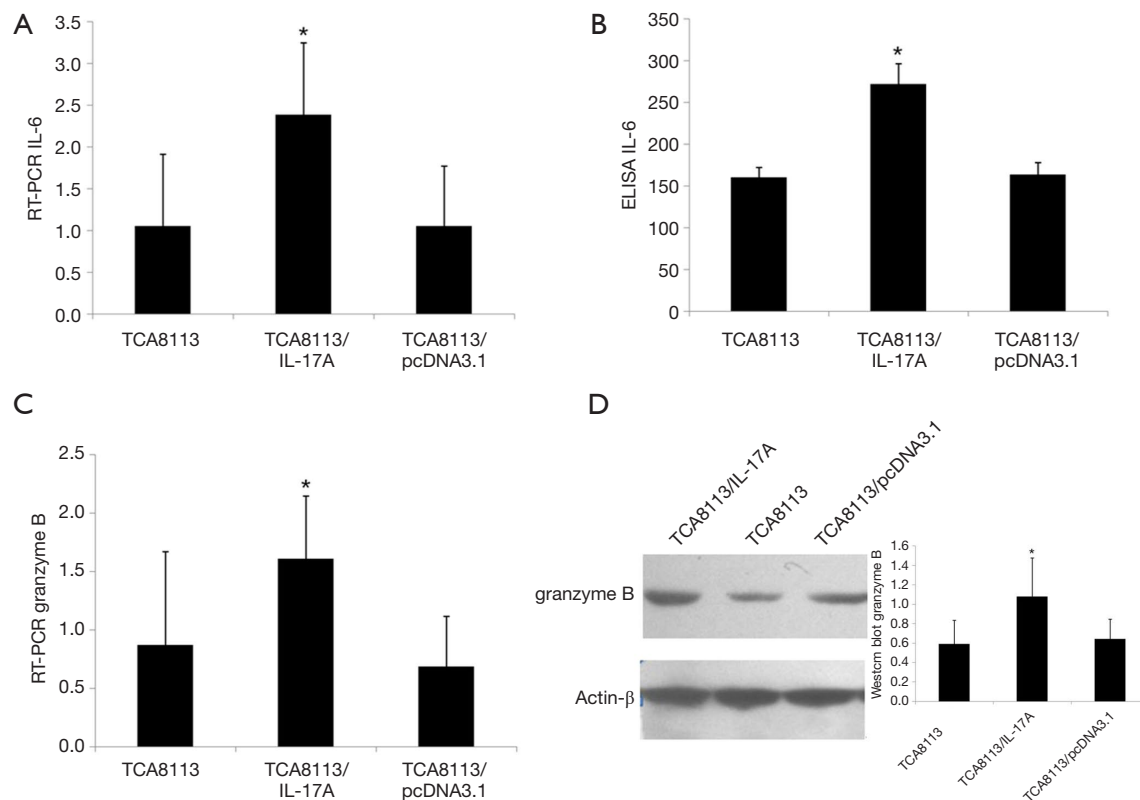


Figure 4 The increased expression of IL-6 and granzyme B in OSCC implants. (A,B) Both the mRNA and the protein level of IL-1 in TCA8113/pCDNA3.1-IL-17A group were significantly higher than those in control groups; (C,D) compared with controls, the transcripts and protein of granzyme B were also elevated in TCA8113/pCDNA3.1-IL-17A group. *, $P < 0.05$.

The expression of IL-6 and granzyme B was increased in OSCC implants

To explore the mechanism how IL-17A induces Th17 differentiation, the expression of IL-1 β , IL-6 and IL-23A in the OSCC implants was investigated. Q-PCR revealed that only the mRNA of IL-6 was significantly elevated in the TCA8113/pCDNA3.1-IL-17A group (Figure 4A), while the transcription of IL-1 β and IL-23A showed no difference (data not shown). ELISA assay confirmed the elevated IL-6 expression in the TCA8113/pCDNA3.1-IL-17A group in contrast to the TCA8113/pCDNA3.1 and TCA8113 groups (Figure 4B). Therefore, it was suggested that the differentiation of Th17 lymphocytes in the OSCC micro-environment was promoted by the elevated IL-6. Moreover, the expression of granzyme B was also examined to explore how Th17 cells repressed OSCC growth. In the TCA8113/pCDNA3.1-IL-17A group, the transcription of granzyme B was higher than those in TCA8113/pCDNA3.1 and TCA8113 groups (Figure 4C). Consistently, the Western

blotting band of granzyme B in TCA8113/pCDNA3.1-IL-17A group was also much denser (1.081 ± 0.394) than the bands in TCA8113/pCDNA3.1 (0.646 ± 0.201) and TCA8113 (0.591 ± 0.245) groups (Figure 4D). These results suggested that differentiated Th17 lymphocytes retarded OSCC growth by recruiting more cytotoxic T lymphocytes (CTLs) in OSCC micro-environment.

Discussion

The micro-environment of tumor cells not only decides their survival and growth, but also affects their pharmacological responses to anti-tumor drugs. Tumor micro-environment is constituted by non-tumor cells, extracellular matrix, blood vessels and signaling molecules (18). As one component of the non-tumor cells in tumor micro-environment, the population of the immune cells was a mixture of T lymphocytes, B lymphocytes, natural killer cells and macrophages. However, the

infiltrated immune cells were found to play dual roles in tumor progression. The immune responses were putatively regarded to inhibit tumorigenesis, but in the context resembling chronic inflammation, the immune responses promoted malignancy (19). For example, the CD4⁺ Th1 cells function as the major anti-tumor cells by secreting IFN- γ , while the CD4⁺ Th2 cells were reported to prevent or enhance tumor progression in different context (20). Similar to CD4⁺Th2 cells, the functions of CD4⁺ Th17 cells in tumor progression were also controversial based on the different kinds of tumors (21,22).

As the newly discovered subpopulation of T helper cells, Th17 lymphocytes have been detected in several tumor tissues and circulating blood (6). Previous studies demonstrated that OSCC has a close correlation to immune responses, such as the pre-tumor lesion, lichen planus or vitiligo, in which there is noticeable lymphocyte infiltration (23). Based on above findings, we hypothesized that the immune micro-environment of OSCC was also aberrant as other carcinomas. The increased concentration of Th17 cells detected by FACS in the peripheral blood and tumor tissues from OSCC patients verified the infiltration of Th17 cells in OSCC micro-environment. However, it requires to be elucidated the function of the infiltration of Th17 cells.

The tumors in immune-compromised mice were diminished after inoculated with the *ex vivo* expanded Th17 cells (24). This finding implicated the anti-tumor activity of Th17 lymphocytes. However, only 10–30% T lymphocytes differentiated into Th17 cells in *ex vivo* expansion (25). Hence, such a low ratio limited the application of transfusing the *ex vivo* expanded Th17 cells into patients because of the scarce sources and expensive price. Fortunately, the proceeding study discovered that Th17 lymphocytes could be induced in the tumor micro-environment and amplified *in vivo* (25). A variety of cytokines secreted by tumor cells, such as IL-1 β , IL-23, IL-6 and TGF- β , provides a micro-environment facilitating the differentiation of Th17 cells (26). Further investigations found that these cytokines also originated from tumor associated macrophage (TAM), bone marrow-derived suppressing cells (MDSC), antigen present cells (APCs) and fibroblasts (27). IL-17 was reported to stimulate above cells to secrete the cytokines (28). Therefore, we hypothesize that the increase of IL-17 in tumor micro-environment can inhibit tumor growth by promoting Th17 differentiation.

In our study, we generate the human OSCC cell line stably expressing human IL-17A by transfecting TCA8113

with full length cDNA of human IL-17A. By inoculating the IL-17A-expressing TCA8113 cells into human-PBL-SCID-beige mice, we get an artificial environment mimicking human immune response to address if the elevation of IL-17A could enhance the Th17 infiltration in local OSCC micro-environment and to explore the mechanism how Th17 cells suppress OSCC growth. In the TCA8113/pcDNA3.1-IL-17A group, the expression of IL-17A was remarkably higher than those in TCA8113/pcDNA3.1 and TCA8113 groups. As the marker of Th17 differentiation, the expression of transcription factor ROR γ t was also robustly activated in TCA8113/pcDNA3.1-IL-17A group compared to control groups, suggesting that an increased Th17 differentiation resulted from the IL-17A stimulus. Finally, the evidently reduced tumor volume in TCA8113/pcDNA3.1-IL-17A group confirmed the inhibitory role of Th17 cells on OSCC growth. On the other hand, we noticed that the growth curve of TCA8113/pcDNA3.1-IL-17A group was only distinct from those of TCA8113/pcDNA3.1 and TCA8113 groups after 9 days of tumor implantation. This phenomenon may result from the lower production of IL-17A by TCA8113/pcDNA3.1-IL-17A cells in the initial stage or the window period required for Th17 differentiation. Since there are few concerns on the differential roles of Th17 cells in the early and later stages of tumors, and/or the dosage-dependent manner of Th17 cells, we plan to clarify those concerns in our further study.

Previous studies disclosed that IL-1 β , IL-6, IL-21 or IL-23 was necessary for the differentiation of native T cells into Th17 cells (26–28). To clarify the mechanism contributing to the increase of Th17 cells, the expression of above mentioned cytokines was examined in the implanted OSCC samples. The elevated IL-6 expression was detected in the TCA8113/pcDNA3.1-IL-17A group, while the transcription of IL-1 β and IL-23A showed no difference among groups. This finding suggested that IL-6 played a critical role in the promoted differentiation of Th17 cells, which was coincided to the previous report (28). Although IL-6 was also reported to enhance the angiogenesis through IL-6/Stat3 pathway in tumors (28), which conflicted to the tumor-suppressing effect in our study, it could be interpreted that the inhibitory effect of Th-17 cells promoted by IL-6 overwhelmed the angiogenesis stimulated by IL-6. Of course, further study is required to verify this assumption.

Although there were a lot of studies confirmed the inhibition of Th17 lymphocytes on tumor growth (29,30),

the mechanism still needs to be elucidated. Putatively, Th17 cells were regarded to induce more CTLs against tumor specific-antigen in the local micro-environment (31). So the granzyme B, the hallmark of CTLs, was tested in the implanted OSCC samples. Both Q-PCR and Western blotting results indicated an increase of granzyme B in the TCA8113/pcDNA3.1-IL-17A cells. Because the repression of the increased activity of CTLs on tumor growth has been established (31), it implicated that the same mechanism was also responsible for the inhibition of Th-17 cells on OSCC growth.

In summary, our study suggested that IL-17 was able to induce the differentiation of Th17 cells by increasing IL-6 production, and Th17 cells inhibited OSCC growth by exerting the cytotoxic effect. Application of differentiated Th17 cells may greatly retard the growth and reduce the mass and size of OSCC, which would gain more time for patients or decrease the dosage of anti-tumor drugs. Additionally, the original concentration of the differentiated Th17 cells in the OSCC focus could be exploited as an index for prognosis, in which the more Th17 cells means the better prognosis. Even though, we have to acknowledge that the repression of Th17 cells in OSCC could work through more than the cytotoxic activity, which needs more explorations. Moreover, if Th17 cells have the differential roles in OSCC at different stages also requires further investigation to decide the timing of its clinical administration. Another limitation in the application of Th17 cells is the potential risk of enhancing the progression of tumors other than OSCC, because its function in other tumors has not been clarified yet. Taking above shortcomings into account, the local induction of the differentiation of Th17 cells, instead of the intravenous injection of differentiated Th17 cells would be the preferential approach for OSCC treatment, which has been performed in our study.

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Footnote

Conflicts of Interest: All authors have completed the ICMJE uniform disclosure form (available at <http://dx.doi.org/10.21037/tcr.2018.06.05>). 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. The study was conducted in accordance with the Declaration of Helsinki (as revised in 2013). This study was approved by the Ethical Committee of Chinese PLA (People's Liberation Army) General Hospital. All of the patients participated in this study have signed the informed consent for using their medical records and tissue specimens for research purposes. The animal experiments were approved by the Animal Research Committee at Beijing Laboratory Animal Research Center and carried out in accordance with the authorized Protocol (P2015015).

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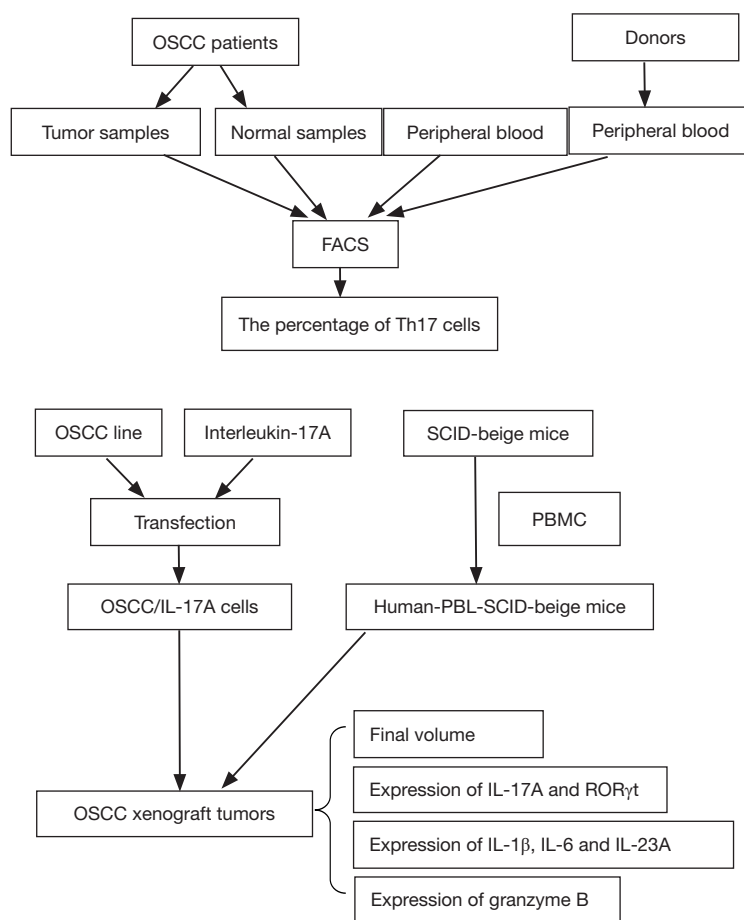


Figure S1 Research process of this study.