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Tumor-infiltrating IL-17A⁺ cells determine favorable prognosis and adjuvant chemotherapeutic response in muscle-invasive bladder cancer

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ABSTRACT

The role of IL-17A⁺ cells remains controversial among various cancer types. This study aimed to investigate the effects of IL-17A⁺ cells on tumor immune contexture and clinical outcome in muscle-invasive bladder cancer (MIBC). In this study, we enrolled 141 patients from Zhongshan Hospital, 118 patients from Shanghai Cancer Center and 403 patients from TCGA cohort. In vitro studies were conducted in 32 freshly resected tumors. Survival analysis was conducted using Kaplan–Meier and Cox regression analysis. The results suggested that patients with high levels of IL-17A⁺ cells had prolonged overall survival and recurrence-free survival (HR = 0.268, P < .001; and HR = 0.433, P < .001). Moreover, these patients tended to be at lower risk of death and recurrence after adjuvant chemotherapy (P = .012 and P = .004). An increased number of IL-17A⁺ cells correlated with the infiltration of several anti-tumor immune cells into tumors. In addition, IL-17A⁺ cells had an influence on the recruitment, proliferation, and activation of CD8⁺ cells, and were positively associated with the expression of several anti-tumor immunity in MIBC. Besides, high infiltration of IL17A⁺ cells can predict benefit from ACT for MIBC patients.

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Introduction

Bladder cancer is one of the most common malignancies involving the urinary system.¹ Approximately a quarter of patients with bladder cancer present with muscle invasive disease and often have a much worse prognosis.² For these patients, radical cystectomy alone is associated with an overall cure rate ranging from 50% to 65%. However, the five-year survival rate drops to 40% or lower in patients with invasion beyond the bladder muscle.³ Besides surgical treatment, cisplatin-based chemotherapy has been proved to be effective for advanced and metastatic bladder cancer.⁴ There have been trials estimating the benefit of adjuvant chemotherapy for muscle-invasive bladder cancer patients, but they have been inadequately powered and prematurely closed for lack of sufficient data.⁴ As a result, clinicians are not yet able to identify patients who are most likely to benefit from adjuvant chemotherapy.

The efficacy of chemotherapy may not only be determined by its direct effects on cancer cells, but by off-target effect within the host immune system as well. Recent studies have been focused on the role of immune response in chemotherapy-induced cytotoxicity.⁵⁻⁷ Moreover, the intrinsic immune features can predict survival and potential benefit from chemotherapy in patients with bladder cancer.⁸ Interleukin-17A (also known as IL-17A) has attracted much attention in the past few years for its central role in immunity. It could enhance the expression of various chemokines including CXCL9 and CXCL10 and is believed to be actively involved in the recruitment of other immune cells.⁹⁻¹¹ The IL-17Aproducing cells, which are composed of CD4⁺ T helper (Th17) cells, cytotoxic CD8⁺ T (Tc17) cells and $\gamma\delta T$ ($\gamma\delta$ -17) cells, are a fascinating cell population for their dichotomous nature in cancer.¹² On the one hand, IL-17A⁺ cells have exhibited prominent anti-cancer ability in several types of cancer,¹³⁻¹⁵ but, on the other hand, this cell population have been reported to promote tumor growth and metastasis as well.¹¹

In this study, we evaluated the prognostic value of $IL17A^+$ cells in a large group of patients with muscle-invasive bladder cancer, validated its predictive significance for survival benefit from cisplatin-based chemotherapy. To our knowledge, this study is the first to reveal the function of $IL-17A^+$ cells in bladder cancer.

*These authors contributed equally to this work.

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Supplemental data for this article can be accessed on the publisher's website.

Materials and methods

Ethics approval and consent to participate

Informed consent was collected from patients in TMAs and this study was approved by the Clinical Research Ethics Committee of Fudan University and each hospital. All procedures of specimen collection, handling, and application from patients are acknowledged by patients with written informed consents and were approved by institutional review board and ethics committee of Fudan University and Shanghai Jiao Tong University.

Study population

This study was approved by the Clinical Research Ethics Committee of Fudan University and each hospital. In this study, we enrolled three independent patient cohorts, Zhongshan Hospital (ZS) cohort (N = 141), Shanghai Cancer Center (SCC) cohort (N = 118) and the Cancer Genome Atlas (TCGA) cohort (N = 403). The ZS cohort comprised of 215 patients receiving radical cystectomy between 2002 and 2014 in Zhongshan Hospital (Shanghai, China) and 73 patients were excluded for benign or nonmuscle invasive disease. The SCC cohort included 178 patients receiving surgery between 2008 and 2012 in Shanghai Cancer Center (Shanghai, China). However, 41 patients with benign or non-muscle invasive disease and 18 patients without clinicopathological information were excluded. After surgery, 119 patients from the two cohorts received cisplatin-based combination chemotherapy for at least one therapeutic cycle. All these patients did not receive neoadjuvant chemotherapy or radiation treatment. During immunohistochemistry staining, one patient from each cohort was lost due to detachment. Overall survival and recurrencefree survival were calculated as interval from the date of cystectomy to death or first recurrence. The follow-up protocol was instructed by EAU guidelines.¹⁶ The follow-up period ended in July 2016. The median follow-up periods in ZS cohort and SCC cohort were 56 months and 29 months, respectively. The TCGA cohort was composed of 413 patients, among them, 407 patients were histologically diagnosed with muscle-invasive bladder cancer. Four patients were excluded for lack of IL-17A mRNA information. Two patients without overall survival time and 87 patients without recurrence data were excluded when performing survival analysis. The information of these patients was retrieved from http://www.cbio portal.org in Feb. 2018. Detailed characteristics of all three cohorts are presented in supplementary table 1. The cutoff value of IL-17A⁺ cells was 2 cells/HPF (*200 magnification) and the cutoff point of IL-17A mRNA was 0.2 (RNAseq Version2), which is determined by X-tile 3.6.1 (Yale University).

Immunohistochemistry (IHC)

Tissue microarray (TMA) construction and the immunohistochemistry protocol were conducted as previously described.¹⁷ For each patient, three sections of tumor tissue were taken from paraffin blocks to construct TMA. Primary antibodies (listed in supplementary table 2) were applied to sections at 4°C overnight for immunohistochemical staining. For different T helper cells, TH1 was defined as CD4⁺ T-bet⁺ cells and TH2 was defined as CD4⁺ GATA3⁺ cells. Then, slides were treated with polyperoxidase-conjugated IgG (OriGene). Staining was performed with diaminobenzidine solution (Biocare Medical) under a microscope and counterstained with hematoxylin. The primary antibody was omitted for the negative controls. The positive cells were enumerated from the representative view of the three sections in highpower field (HPF) and an average number was adopted. Two pathologists who were unaware of patient information evaluated the staining of each specimen with the assistance of Image-Pro Plus 6.0 (Media Cybernetics Inc.). In case of disagreement, the slides were reviewed and the two observers achieved a consensus.

Fresh tumor specimen and flow cytometry

All 32 fresh human specimens were obtained from patients underwent radical cystectomy in Shanghai Cancer Center, Zhongshan Hospital, and Shanghai General Hospital, respectively. Single cells were isolated from freshly resected tumor tissues using collagenase IV. About 1,000,000 cells were collected and incubated in RBC lysis buffer at 4°C for 5 min. Surface markers (listed in supplementary table 2) were stained in Cell Staining Buffer for 30 min at 4°C in dark after blocking Fc-receptors with Human TruStain FcX (Biolegend). Then, cells were fixed by Fixation Buffer (Biolegend), permeabilized by Intracellular staining permeabilization wash buffer (Biolegend) and intracellular cytokine (listed in supplementary table 2) was stained for 40 min at 4°C in dark. Specially, we performed transcription factor staining using the True-Nuclear transcription Factor Buffer set (Biolegend) according to the manufacturer's instructions. Dead cells were excluded by Live/Dead Fixable Dead Cell StainingKit (Invitrogen). Cells were acquired on the FACSCelesta Flow Cytometer (BD Biosciences) and analyzed with FlowJo software (Treestar).

Statistical analysis

Statistical significance was calculated by the Student t-test for comparisons between two groups or one-way ANOVA for multi-group comparisons using GraphPad Prism and SPSS software. Patient characteristics and the association with IL-17A mRNA expression/IL-17A⁺ cells were described statistically and evaluated by Chi-squared test, respectively. Spearman's correlation test was used to compare differences between two continuous variables. Survival analysis was carried out by Log-rank test. P < .05 was considered statistically significant.

Results

Identification of tumor-infiltrating IL-17A⁺ cells in muscle-invasive bladder cancer

IL-17A⁺ cells were evaluated by immunohistochemistry on tissue microarrays (Figure 1a). Interestingly, the fraction of



Figure 1. Identification of tumor-infiltrating IL-17A⁺ cells in muscle-invasive bladder cancer. (a) Representative immunochemistry staining of IL-17A⁺ cells in muscle-invasive bladder. (b) A fraction of patients with high/low IL-17A⁺ cells in different TNM stages. (c) The number of tumor-infiltrating IL-17A⁺ cells in different TNM stages. (d) Representative flow cytometry analysis of the surface markers expressed by IL-17A⁺ cells from fresh human muscle-invasive bladder cancer samples. (e) Quantification analysis of surface markers expressed by tumor-infiltrating IL-17A⁺ cells.

patients with high levels of tumor-infiltrating IL-17A⁺ cells and the number of IL-17A⁺ cells seemed to negatively correlate with TNM stage (Figure 1b,c, Supplementary table 3), indicating that the infiltration of IL-17A⁺ cells into tumor might associate with a better prognosis for patients. We next assessed the surface markers expressed by tumor-infiltrating IL-17A⁺ cells in freshly isolated clinical specimens using flow cytometry. Among all 32 fresh tumor samples, there were 6 tissues without detectable IL-17A⁺ cells. The majority of IL-17A⁺ lymphocytes expressed CD3 or CD4 (Figure 1d,e). This finding is consistent with previous studies that TH17 cells are a predominant source of IL-17A.¹⁸⁻²⁰

Prognostic value of IL17A⁺ cells and IL-17A mRNA

To elucidate the prognostic value of IL-17A⁺ cells, we performed Kaplan-Meier analysis using the data from Zhongshan Hospital (ZS), Shanghai Cancer Center (SCC) and TCGA database. We observed that patients with elevated expression of IL-17A⁺ cells and IL-17A mRNA tended to have prolonged overall survival in all three cohorts (P < .001, P = .003 and P = .041, respectively; Figure 2a). Moreover, high infiltration of IL-17A⁺ cells into tumors indicated longer recurrence-free survival in ZS cohort and SCC cohort (P < .001 and P = .012, respectively; Figure 2b). However, the IL-17A mRNA failed to predict recurrence-free survival in TCGA cohort (P = .095; Figure 2b).

We then combined the three cohorts to evaluate whether IL-17A is an independent prognostic factor for muscleinvasive bladder cancer patients. In a multivariate analysis including age, gender, tumor stage, lymphovascular invasion, and adjuvant treatment as confounding variables, the hazard ratio (HR) of disease mortality and recurrence is 0.468 (95% CI, 0.329 to 0.665; P < .001) and 0.685 (95%CI, 0.480 to 0.979; P = .038) respectively (Figure 2c,d). IL-17A proved to be an independent protective factor for patients.

Tumor-infiltrating IL-17A⁺ cells predict response to adjuvant chemotherapy

In the combined cohort, there was no evidence suggesting that adjuvant chemotherapy could prolong overall survival or recurrence-free survival for muscle-invasive bladder cancer patients (P = .598 and P = .891, respectively; Figure 3a,b). However, after discriminating patients with tumor-infiltrating IL-17A⁺ cells, adjuvant chemotherapy significantly enhanced overall survival and recurrence-free survival in IL-17A⁺ cells high group of patients (P = .012 and P = .004, respectively) while it failed in IL-17A⁺ cells low group (Figure 3cf). A test for the interaction between IL-17A⁺ cells and adjuvant chemotherapy further confirmed the survival benefit in terms of overall survival (P = .003, Figure 3c,e) and recurrence-free survival (P = .002, Figure 3d,f) observed in IL-17A⁺ cells high group. Taken together, these results suggested that tumorinfiltrating IL-17A⁺ cells could successfully identify patients with muscle-invasive bladder cancer who would most likely to benefit from adjuvant chemotherapy.

The TCGA cohort was not included because over a third of patients lacked information of whether they received adjuvant pharmaceutical treatment or not. Moreover, the criteria utilized to treat patients, the regimens utilized and the duration



Figure 2. Tumor-infiltrating IL-17A⁺ cells and IL-17A mRNA indicate better survival for patients. (a and b) Kaplan–Meier curves comparing OS and RFS in muscleinvasive bladder cancer patients with high and low tumor-infiltrating IL-17A⁺ cells/IL-17A mRNA. (c and d) Multivariate cox regression analysis for IL17A expression and clinic-pathological variables.



Figure 3. Tumor-infiltrating IL-17A⁺ cells indicate an improved therapeutic response to adjuvant chemotherapy. (a and b) Survival curves for the use of adjuvant chemotherapy (ACT) in all patients, (c and d) in patients with low intra-tumoral IL-17A⁺ cells, (e and f) and in patients with high intra-tumoral IL-17A⁺ cells.

of therapy were not clear. These variables may possibly influence outcomes.

Tumor-infiltrating IL-17A⁺ cells associate with enhanced anti-tumor immunity

The immune contexture, which is determined by the composition, density, and functional state of the tumor-infiltrating leukocytes, can yield information relevant to prognosis, prediction of a treatment response and a variety of other pharmacodynamic parameters.²¹ According to previous reports, IL-17A happened to be a chef orchestrator of immunity.²² Thus, we decided to looked into the differences of immune cells infiltration between two groups of patients in order to understand the possible mechanism underlying the phenomena mentioned above. We performed semiquantitative immunohistochemistry and calculated the correlation of IL-17A⁺ cells with several types of immune cells infiltrating into the tumor (Figure 4a, Supplementary figure 1A). IL-17A⁺ cells were positively correlated with CD8⁺ T cells, dendritic cells, B cells, neutrophils, and macrophages (Figure 4b, Supplementary figure 1B). When regarding tumor-infiltrating IL-17A⁺ cells as a dichotomous data, the number of B cells, dendritic cells, and CD8⁺ T cell

was significantly higher in IL-17A⁺ cells high group than in IL-17A⁺ cells low group. This was in line with our previous finding that B cells were a protective factor for muscle-invasive bladder cancer patients.²³ With consecutive tissue sections, we observed the distribution of IL-17A⁺ cells was close to CD8⁺ cells, together with increased expression of IFN- γ and GZMB, the interaction between these cells deserves to be further investigated (Supplementary figure 1 C).

Recent studies have revealed that CD8⁺ T cells were necessary for therapeutic activity⁵ and have determined the importance of cytokines in promoting CD8/TH1 response to chemotherapy.^{6,24,25} Thus, we compared the different functional states of CD8⁺ T cells between two groups of patients. The percentage of CD8⁺ T cells expressing GZMB, IFN-y and PRF1 was significantly higher in IL-17A⁺ cells high tumors, as compared to IL-17A⁺ cells low tumors (Figure 5a, Supplementary figure 2A). The proportion of $GZMB^+$ $CD8^+$ T cells, IFN- γ^+ CD8⁺ T cells, and PRF1⁺ CD8⁺ T cells was also increased among CD45⁺ cells in IL-17A⁺ cells high tumors (Figure 5b). In addition, CD8⁺ T cells in IL-17A⁺ cells high group exhibited more proliferative ability than their counterparts in IL-17A⁺ cells low tumors (Figure 5a,b). This could be one of the reasons that there were more CD8⁺ T cells detected in IL-17A⁺ cells present group



Figure 4. Association of IL-17A⁺ cells and various tumor-infiltrating immune cells. (a) Representative immunochemistry staining of B cells, Dendritic cells and CD8⁺ T cells in IL-17A⁺ cells high tumors and in IL-17A⁺ cells low tumors. (b) Pearson correlation analysis of IL-17A⁺ cells with other tumor-infiltrating lymphocytes. (c) Differences in tumor-infiltrating lymphocytes between IL-17A⁺ cells high tumors and IL-17A⁺ cells low tumors.



Figure 5. Tumor-infiltrating IL-17A⁺ cells indicate enhanced anti-tumor immunity. (a) Difference in effector cytokines and proliferation marker expressed by CD8⁺ T cells in IL-17A⁺ cells low tumors and IL-17A⁺ cells high tumors. (b) Proportion of CD8⁺ T cells expressing effector cytokines and proliferation marker in CD45⁺ cells. (c) Difference in immunosuppressive molecules expressed by CD8⁺ T cells. (d) Proportion of CD8⁺ T cells expressing immunosuppressive molecules in CD45⁺ cells. (e and f) Effector cytokines and immunosuppressive molecules expressed by CD45⁺ cells in IL-17A⁺ cells low tumors and IL-17A⁺ cells high tumors.

of patients (Figure 4c). However, the checkpoint expressed by $CD8^+$ T cells exhibited no difference between IL-17A⁺ cells high and IL-17A⁺ cells low tumors, except for TIGIT (Figure 5c, Supplementary figure 2B). Among CD45⁺ cells, the CTLA-4⁺ CD8⁺ cells and LAG-3⁺ CD8⁺ cells also significantly increased (Figure 5d) in IL-17A⁺ cells high tumors.

In addition to cytotoxic T cells, the percentage of CD45⁺ T cells expressing immune effector molecules all increased in IL-17A⁺ cells high tumors (Figure 5e). However, we also observed an elevated expression of PD-1 by CD45⁺ cells (figure 5f). Results from TCGA database further confirmed our findings. GSEA analysis using the Hallmark gene sets²⁶ identified inflammatory response and IFN-y response as two of the most significantly up-regulated pathways in IL-17A mRNA high tumors (Supplementary figure 3A). In IL-17A mRNA high group, expression of chemokines that could recruit CD8⁺ T cells was significantly higher (Supplementary figure 3B). This could be another explanation for more CD8 + T cells observed in IL-17A⁺ cells high group of patients. Genes mediating cytotoxic functions^{27,28} were also positively correlated with IL-17A mRNA expression (Supplementary figure 3 C). Taken together, these results suggested that tumor-infiltrating IL-17A⁺ cells were associated with enhanced anti-tumor immunity.

Discussion

It has been well established that IL-17A and IL-17A⁺ cells are important in promoting inflammation and linked to many immune/autoimmune-related diseases.^{29,30} However, their role in cancer remained fascinating and required further investigation.^{20,31} Our study revealed that IL-17A⁺ cells were correlated with prolonged survival and response to adjuvant chemotherapy. The underlying mechanism was found associated with the recruitment of various anti-tumor immune cells and activation of $CD8^+$ T cells.

Patients are of great heterogeneity in clinical outcomes even with identical clinicopathological features. Thus, integrating multiple biomarkers, especially the intrinsic immune features, has aroused much interest and been proved to provide a more accurate prediction.^{32,33} As a chief orchestrator of immunity, IL-17A has the ability to induce the expression of important proinflammatory cytokines and recruit a variety of immune cells.³³ Thus, IL-17A⁺ cells may indicate the immune contexture of hosts. Our study demonstrated that tumor-infiltrating IL-17A⁺ cell was indeed an ideal biomarker to differentiate patients suitable for more active surveillance and adjuvant chemotherapy. Of interest, almost half of the tumor tissues were with low expression of IL-17A⁺ cell or IL-17A mRNA. This suggested that tumors, particularly with advanced stage, could prevent the infiltration of IL-17A⁺ cells through some sort of unknown mechanism. This deserves further investigation and may be a potential target to enhance anti-tumor immunity.

In current clinical practice, adjuvant chemotherapy is applied to patients who are at great risk of disease relapse (pT3 + pT4 or N+), but the debate is still ongoing for it has not exhibited obvious benefit for muscle-invasive bladder cancer patients.³⁴ In our study, patients with high tumor-infiltrating IL-17A⁺ cells have improved response to adjuvant chemotherapy. According to previous researches, immune cell/cytokine pathways involving CD8⁺ T cells were responsible for mediating response to chemotherapy.⁶ Moreover, there have been clinical trials proving that the usage of immune checkpoint inhibitor following treatment with platinum-based chemotherapy was effective in patients with locally advanced or metastatic urothelial carcinoma.^{27,28} The efficacy seemed to be underlying genomic,

molecular, and immunological factors. Based on these researches, we hypothesized that the efficacy of adjuvant chemotherapy in IL-17A⁺ cells high group of patients was due to the activation of cytotoxic immune cells. Actually, we have observed enhanced expression of anti-cancer cytokines by CD8⁺ T cells in IL-17A⁺ cells high tumors. A confusing point is that intratumoral IL-17A⁺ cells were actually removed by surgery, how these cells contribute to the therapeutic outcome remains unknown. Our previous study revealed that immunophenotype, which is an important feature of cancer, could predict benefit from adjuvant chemotherapy in bladder cancer.⁸ IL-17 is reported to be a chief orchestrator of immunity.²² Thus, high infiltration of IL-17A⁺ cells could represent an immune-activated phenotype of bladder cancer and serve as a predictive biomarker for adjuvant chemotherapy. In addition, it is reported that the immune composition of metastatic site is close to that of the primary tumor.³⁵ Patients with a high frequency of intratumoral IL-17A+ cells might also have high infiltration of this cell population in their metastatic sites if the disease relapsed. It could be a possible reason for how these cells contribute to the therapeutic outcome but need to be confirmed in further study.

Interestingly, we also observed the elevated expression of immunosuppressive markers by lymphocytes in IL-17A⁺ cells high tumors, including PD-1. IL-17A mRNA was also positively correlated with several immunosuppressive genes (supplementary Figure 4d). This intrinsic correlation needs further investigation to be explained. We assumed that it might be a counterbalance mechanism in response to hyper-activated anti-tumor immunity status in IL-17A⁺ high tumors. This finding suggested that patients with high intratumoral IL-17A⁺ cells might remarkably benefit from immune checkpoint blockade therapy. However, we lacked sufficient clinical data to carry out a further investigation at present.

There were several limitations to be addressed in this study. It was a retrospective study. Therefore, a prospective study with a larger cohort is needed in the future. In addition, the data we obtained from TCGA database were the mRNA information of IL-17A; its correlation with tumor-infiltrating IL-17A⁺ cells remained unclear. Also, we have not uncovered how IL-17A⁺ cells influenced the downstream cells. Interventional and functional studies are required to fully understand the underlying mechanisms.

In conclusion, this is the first study identified IL-17A⁺ cells as favorable factors for muscle-invasive bladder cancer patients in three independent cohorts. Furthermore, high infiltration of IL-17A⁺ cells into tumors is associated with an anti-cancer immune contexture and could predict therapeutic response to adjuvant chemotherapy.

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Disclosure of potential conflicts of interest

No potential conflicts of interest were disclosed.

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Author contributions

Z. Wang, Q. Zhou, H. Zeng, and H. Zhang for acquisition of data, analysis, and interpretation of data, statistical analysis, and drafting of the manuscript; Z. Liu, Q. Huang, Y. Xiong, J. Wang, Y. Chang, Q. Bai, Y. Xia, Y. Wang, Y. Zhu, L. Xu and B. Dai for technical and material support; L. Liu, J. Guo and J. Xu for study concept and design, analysis, and interpretation of data, drafting of the manuscript, obtained funding and study supervision. All authors read and approved the final manuscript.

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