



The Proportion of V δ 1T Cells in Peripheral Blood Is Correlated with Prognosis of Sepsis

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ABSTRACT

Background: Sepsis is a serious condition with a high mortality rate, and septic patients often have organ dysfunction, low tissue perfusion and hypoxia, lactic acidosis, oliguria, or functional brain changes.

Objective: To observe the number and the function of V δ 1T cells in peripheral blood of septic patients, to analyze the clinical significance of detecting V δ 1T cells, and to clarify the correlation of their presence with the prognosis of sepsis.

Methods: The basic data of the septic patients were recorded at admission. The immunosuppressive function-related molecules on the surface of V δ 1T cells were detected, and the immunosuppressive function of V δ 1T cells was also evaluated.

Results: Compared with the healthy controls, the proportion of V δ 1T cells in the blood of septic patients significantly decreased ($P < 0.01$). The proportion of V δ 1T cells in septic patients correlated with the patients' condition ($P < 0.05$). The expression of glucocorticoid-induced tumor necrosis factor receptor (GITR), cytotoxic T-lymphocyte-associated protein 4 (CTLA-4), and T-cell immunoglobulin and mucin domain-containing protein 3 (TIM-3) on the surface of V δ 1T cells in the blood of septic patients significantly increased ($P < 0.01$). The increase of V δ 1T cells in septic patients had inhibitory effects on T cell proliferation and interferon (IFN)- γ secretion. These findings implied that the immunosuppression of V δ 1T cells in the peripheral blood of septic patients was significantly higher than that of the healthy controls ($P < 0.01$).

Conclusion: Changes in V δ 1T cells in septic patients were closely related to the patient's condition and prognosis.

Keywords: Sepsis, Peripheral blood, V δ 1T cells, Immunosuppression, Proliferation

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INTRODUCTION

Sepsis refers to the systemic inflammatory

response caused by infection and diagnosed by the presence of bacteria or highly suspicious infections (1). Sepsis is a dangerous condition

with a high mortality rate, and septic patients often have organ dysfunction, low tissue perfusion, and hypoxia, lactic acidosis, oliguria, or functional brain changes (2, 3). Although a series of studies have been conducted on sepsis over the past 20 years, more than one-half of deaths in intensive care units (ICU) are still caused by septic shock and multiple organ dysfunction, while sepsis is still the main cause of death for non-cardiac patients in the ICU (4, 5). At present, there has been no fundamental breakthrough regarding the treatment of sepsis due to the unclear underlying mechanism. Therefore, studying the pathogenesis of sepsis can help us better understand this complex syndrome and discover a more effective treatment.

The immunosuppression that occurs with sepsis is also described as a compensatory anti-inflammatory response syndrome (CARS) (6). The immune response to sepsis is characterized by phagocytosis, antigen presentation, and apoptotic functions of dendritic cells, neutrophils, T cells and, B cells, and it is also accompanied by a large amount of secretion of anti-inflammatory cytokines, including interleukin (IL)-4, IL-10, etc. (7-9). Some researchers believe that the initial stage of inflammation after the onset of sepsis is associated with the presence of anti-inflammatory responses, so patients with sepsis may experience the inability to clear the pathogen or have a secondary infection (10). As the disease progresses, patients with sepsis experience an immunosuppressive state which may explain why most clinical trials involving anti-inflammatory strategies do not significantly improve the prognosis of patients with sepsis (11). It means that correcting the immunosuppressive state of patients with sepsis might be important for the treatment of this disease (12). In recent years, the functions of immunosuppressive cells including regulatory T cells have been proven to be involved in maintaining the immunosuppressive state in septic patients (12-16). According to the expression of T cell receptors (TCR), T cells are mainly divided

into 2 subsets: $\alpha\beta$ T cells and $\gamma\delta$ T cells (17). $\gamma\delta$ T cells are a kind of T lymphocytes expressing TCR consisting of γ chains and δ chains on their surfaces and are regarded as innate immune cells (18). $\gamma\delta$ T cells are divided into two cell subsets: V δ 1 T cells and V δ 2T cells. V δ 1T cells are mainly distributed in epithelial-related lymphoid tissue, and V δ 2T cells are mainly distributed in peripheral blood (19, 20). V δ 1T cells and V δ 2T cells are involved in different effects. V δ 2 T cells mainly play a role in the immune surveillance of tumors and defense responses (21, 22), while V δ 1 T cells mainly show immunoregulatory functions (23). $\gamma\delta$ T cells have become a hot topic of immunotherapy used for the treatment of cancers due to their unique characteristics (24). Of note, the immunosuppressive function of V δ 1T cells was confirmed more than 10 years ago. These cells have been studied to confirm their immunosuppressive function in patients with tumors and autoimmune diseases (25-27). However, the role of V δ 1T cells has been rarely reported in sepsis. Therefore, this study aimed to further clarify the immunosuppressive state of patients with sepsis experience, especially the role of V δ 1T cells that were closely related to the prognosis of septic patients.

MATERIAL AND METHODS

Case Collection

From October 2016 to December 2017, we collected 40 cases of sepsis (mean age 47.3 ± 10.6 years) from the ICU and 40 healthy controls (mean age 45.2 ± 11.6 years) for regular check-ups in our hospital. All patients were eligible based on the diagnostic criteria for sepsis (28) that were established by the International Conference on Sepsis in Washington, DC, in December 2001. Informed consent was obtained from all participants.

Peripheral Blood Mononuclear Cell (PBMCs) Separation

The 15 mL peripheral blood of the healthy controls and septic patients were collected in

a sodium citrate anticoagulant tube (Thermo Fisher Scientific Inc., Waltham, MA, USA) in the early morning. The appropriate amount of lymphocyte separation fluid (Biolegend, CA, USA) was added to the centrifuge tube; then, the diluted peripheral blood was slowly added to the centrifuge tube, placed on the upper level of the lymphocyte separation solution, and centrifuged as 200 g for 20 min. The white membrane layer of the middle white lymphocytes was gently absorbed with a straw and inserted using a centrifuge into a 10-mL sterile RPMI-1640 medium (Gibco, New York, USA) at 180 g for 15 min. The sterile RPMI-1640 medium was discarded, and the cell suspension was suspended in a 10-mL sterile RPMI-1640 medium and centrifuged at 150g for 8min. PBMCs were obtained and used for the following experiments.

Sorting of $\gamma\delta$ T Cells and V δ 1 T Cells

Sorting $\gamma\delta$ T cells and V δ 1 T cells was performed according to the published study in the literature (29, 30). Briefly, PBMCs were cultured with 0.2 mL of RPMI-1640 medium containing 0.125 μ g of anti-TCR V δ 1 monoclonal antibody (each well) on a 48-well plastic culture plate in an incubator at 37°C with 5% CO₂ for 2 h. After PBMCs were resuspended with RPI-1640 medium adding 10% fetal bovine serum (FBS; Gibco, New York, USA) cells were added to a 48-well plate (1.0 mL per well) coated with anti-TCRV δ 1 monoclonal antibody (Cell Signaling Technology, Inc., Danvers, MA, USA) and cultured. Following 2 weeks of culture, the V δ 1T cells with purity higher than 90% were sorted out by flow cytometry.

Detection of Surface Immunosuppressive Molecules in V δ 1 T Cells

In an Eppendorf tube, 1 \times 10⁶ PBMCs were obtained and mixed with 1 mL of the RPMI-1640 medium with 10% FBS (Gibco, New York, USA) for washing, and then centrifuged at 250 g for 8 min. The supernatant was abandoned, and the abovementioned operation was repeated. Subsequently, the cells were suspended in a 0.1

mL cell culture medium, and incubated with 3 μ L of PE-anti-CD3antibody and 3 μ L of FITC-anti-TCR V δ 1 antibody or3 μ L of PE-anti-CD3 antibody and 3 μ L of APC-anti-human GITR antibody/APC-anti-human CTLA-4 antibody/ Pcy5-anti-human TIM-3 antibody at 4°C for 30 min. After being washed twice with the RPMI-1640 medium containing 10% FBS, the cells were suspended in 0.1 mL of a 1% poly formaldehyde-fixed solution (Biolegend, CA, USA) for flow cytometry. All antibodies were purchased from Cell Signaling Technology, Inc., Danvers, MA, USA.

CFSE Cell Proliferation Detection

Naïve CD4 T cells were washed once with a serum-free 10-mL RPMI 1640 culture medium, and cultured with a CFSE staining solution (Invitrogen, Carlsbad, CA, USA) with a final concentration of5 mmol/L for 10 min in a 5% CO₂ incubator at 37°C. And then 5 mL of pre-chilled CFSE staining stop solution containing 5% cultures of 1640 medium was added to the cell and placed on ice for dyeing. After centrifuging at 400 g for 8 min and washing it once with a 10 mL RPMI 1640 culture medium, the cells were suspended in an RPMI-1640 complete medium. The T cells and naïveCD4 T cells (1:5 ratio) were added to a 48-hole plate and incubated for 5 days, and the cells were detected using flow cytometry.

Detection of Cytokine Secretion by V δ 1T Cells in CD4 T Cells

PBMCs were obtained using an aseptic sorting method, and V δ 1T cells and CD4 T cells with a purity greater than 90%were obtained using flow cytometry. Two cell types were added to the 48-hole plate using a 1:1 ratio of 1 μ g/mL of CD3antibody and 2 g/mL of CD28antibody (BD, New Jersey, USA). After incubating for 72 h at 37°C in a 5% CO₂ incubator, a 10- μ L cell activation cocktail was incubated for 6h in 24-plates. Cells were collected, stained, and mixed with an immobilized cell membrane. After washing the cells with a permeable liquid twice, 5 μ L of IFN- γ antibody (BD, New Jersey, USA) was

added at room temperature for 30 min. The cells were suspended twice in 0.2 mL poly formaldehyde-fixed solution at a concentration of 1% (Invitrogen, Carlsbad, CA, USA), and the cells were detected using flow cytometry.

Statistical Analysis

Statistical Product and Service Solutions (SPSS) version 18.0 (SPSS Inc., Chicago, IL, USA) was used to analyze the data. Counting data were expressed as percentages, and the measurement data were expressed with mean±standard deviation (SD). A t-test was used to compare the measurement data between the two groups. Comparison of experimental groups was evaluated using a one-way Analysis of Variance (ANOVA) analysis, followed by a Bonferroni analysis. A P value<0.05 was considered significant.

RESULTS

Characteristics of Patients

As shown in Table 1, patients with sepsis were matched with the healthy controls in age and sex, and there were no significant differences between the two groups (P>0.05). Clinical data for patients with sepsis indicated

that the sources of infection included lung infection (52.5%), abdomen infection (40.0%), and urinary tract infection (5.0%). Pathogens were gram-positive bacilli (47.5%), gram-negative bacilli (20.0%), and Fungus (32.5%). 20 patients died in ICU within 30 days after admission. One patient was transferred from ICU to the general ward and died of malignant arrhythmia on the 45th day after admission. One patient died of multiple organ failure on the 73rd day after admission. A total of 22 patients with sepsis died during their hospitalization, 20 of them died within 30 days, and the remaining 18 patients were discharged after improvement. During the 1-year follow-up, 5 of the 18 patients re-developed pulmonary infection, 2 died of respiratory failure, and the remaining 3 were discharged from the hospital. 3 of the 18 patients had chronic renal insufficiency treated with hemodialysis. 1 patient of the 18 patients developed chronic heart failure and pulmonary hypertension.

Detection of Vδ1T Cell Ratio in the Blood of Septic Patients

The proportion of Vδ1T cells in the peripheral blood of septic patients was measured (Figure 1). The proportion of

Table 1. Clinical characteristics of the patients

Items	Health controls	Patients with sepsis
Number	40	40
Age (year)	45.2±11.6	47.3±10.6
Sex (male/female)	20/20	21/19
SOFA score	-	11.3±4.4
Source of infection		
Lung	-	21(52.5%)
Abdomen	-	16(40.0%)
Urinary tract infection	-	2(5.0%)
Pathogen		
Gram-positive bacilli	-	19(47.5%)
Gram-negative bacilli	-	8(20%)
Fungus	-	13(32.5%)
White blood cell (×10 ⁹ /L)	-	15.9±6.2
PCT, ng/mL	-	5.3±4.8
Mechanical ventilation	-	13(32.5%)
Renal transplantation	-	9(22.5%)
ICU hospitalization (day)	-	23.7±8.1
Mortality (survival/death)	-	55.0%(18/22)

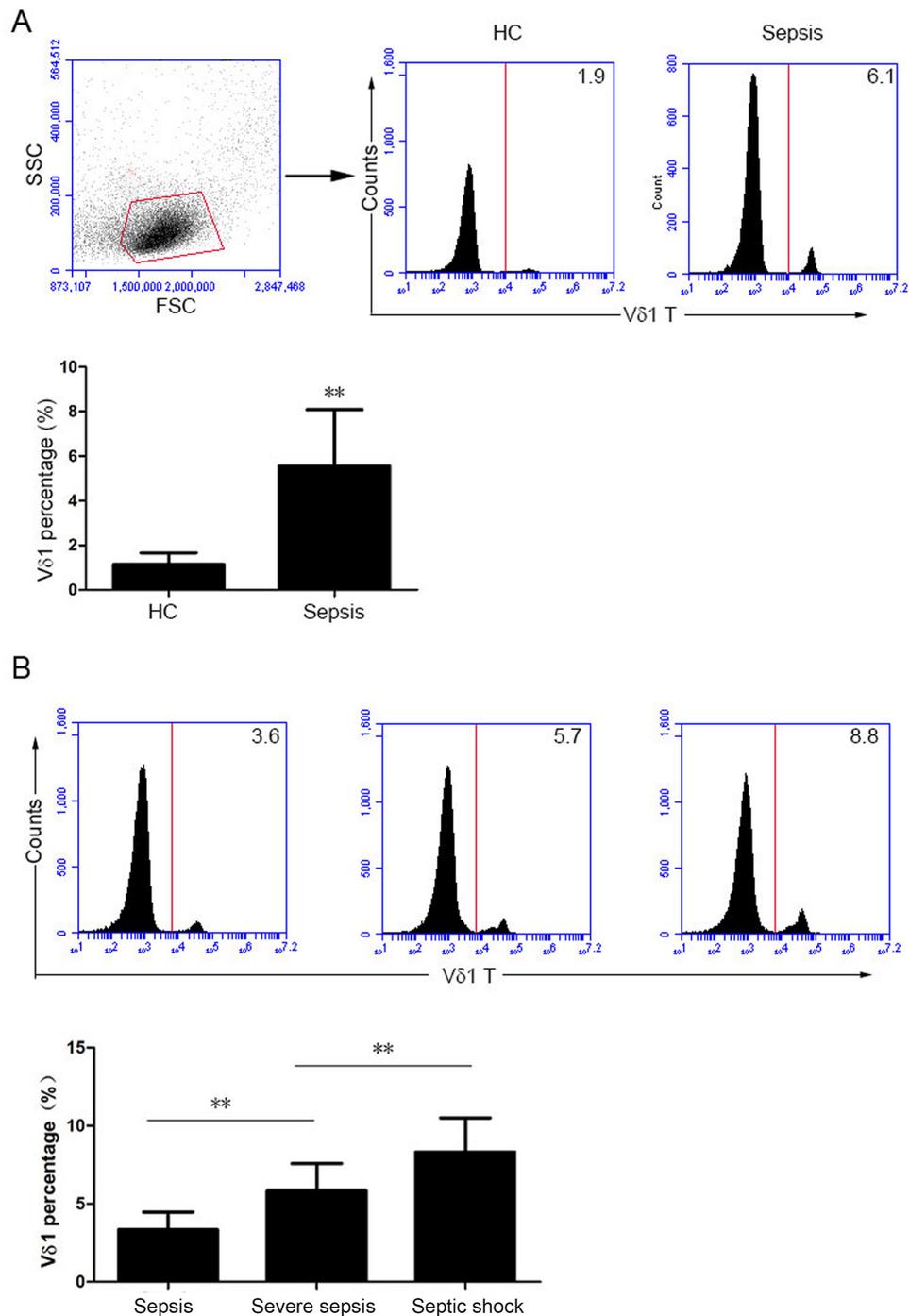


Figure 1. Detection of the proportion of Vδ1T cells in the peripheral blood. Flow cytometry was used to detect the proportion of Vδ1T cells in peripheral blood of the healthy controls and septic patients (A), and the ratio of Vδ1T cells in peripheral blood of the patients with sepsis in different conditions (B). **P<0.01 vs. the health control.

Vδ1T cells in the peripheral blood of the healthy controls (HC) and septic patients was $1.15 \pm 0.51\%$, and $5.56 \pm 2.53\%$, respectively, and the proportion of Vδ1T cells in the peripheral blood of patients with sepsis was significantly higher than that of the HC ($P < 0.01$; Figure 1A). The proportion of

Vδ1T cells in the peripheral blood of septic patients was observed in different conditions. The results in Figure 1B showed that the proportion of Vδ1T cells was $3.34 \pm 1.13\%$ in the peripheral blood of patients with moderate sepsis, $5.84 \pm 1.74\%$ in patients with severe sepsis, and $8.32 \pm 2.18\%$ in patients with septic

shock. With the exacerbation of sepsis, the proportion of Vδ1T cells in the peripheral blood gradually increased ($P<0.01$).

Correlation between the Ratio of Vδ1T Cells in the Peripheral Blood of Septic Patients and Their Condition

Correlation between the ratio of Vδ1T cells in the peripheral blood of septic patients and their condition was observed and the results are shown in Figure 2. The proportion of Vδ1T cells in septic patients was positively correlated with the SOFA score ($r=0.4204$, $P<0.01$; Figure 2A), APACHE II score ($r=0.4744$, $P<0.01$; Figure 2B), and lactic acid level ($r=0.3570$, $P<0.05$; Figure 2C). The proportion of Vδ1T cells in survival patients with sepsis was lower than in non-survival septic patients ($P<0.01$; Figure 2D).

Treatment Changed the Proportion of Vδ1T Cells in the Peripheral Blood of Septic Patients

The proportion of Vδ1T cells in the blood of septic patients was compared before

and after treatment. Compared with before treatment, the proportion of Vδ1T cells in the peripheral blood of septic patients after treatment reduced ($5.56\% \pm 2.53\%$ vs $2.98 \pm 1.43\%$) ($P<0.01$; Figure 3B).

Expression of Immunosuppressive Molecules on Vδ1T Cells in the Peripheral Blood in Septic Patients

The expression of immunosuppressive molecules on Vδ1T cells in the peripheral blood in patients with sepsis and the healthy controls was measured. The percentage of GITR-, CTLA-4-, and TIM-3- positive Vδ1T cells in the peripheral blood of the healthy controls was $7.34 \pm 3.52\%$, $3.82 \pm 1.62\%$, and $12.8 \pm 3.31\%$, respectively. The percentage of GITR-, CTLA-4-, and TIM-3- positive Vδ1T cells in the peripheral blood of the septic patients was $30.2 \pm 6.36\%$, $13.1 \pm 4.27\%$, and $28.1 \pm 6.36\%$, respectively. Compared with the healthy controls, the percentage of GITR-, CTLA-4-, and TIM-3- positive Vδ1T cells in the peripheral blood of septic patients significantly increased ($P<0.01$; Figure 4).

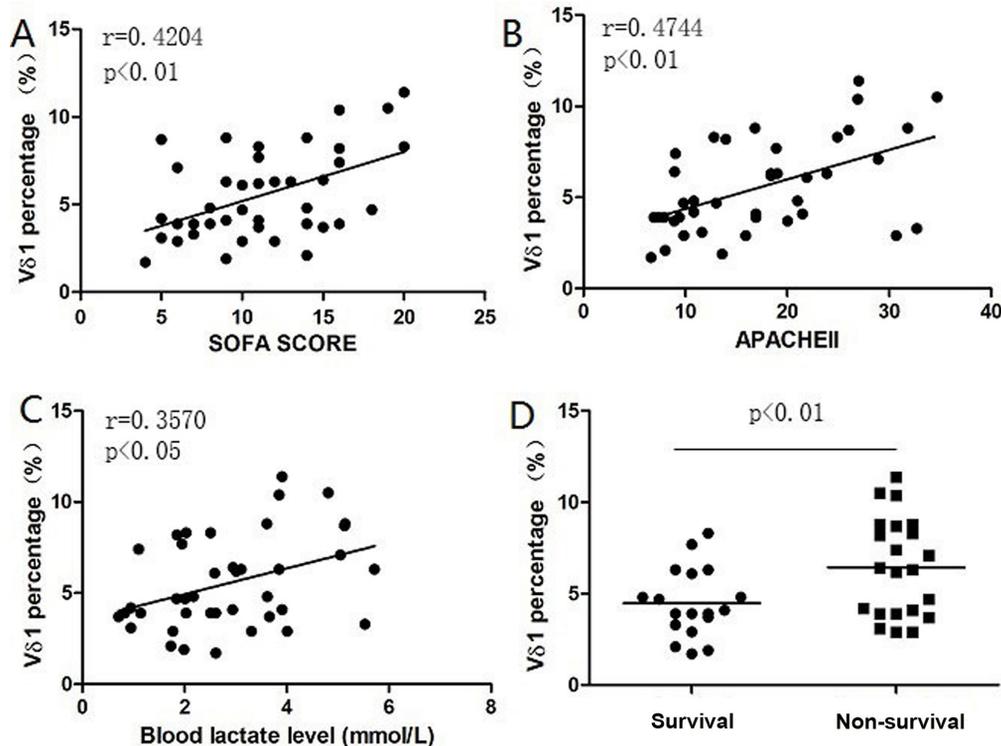


Figure 2. The proportion of Vδ1T cells in septic patients positively correlated with their condition. Correlation between Vδ1T cells in the peripheral blood is the SOFA score (A), APACHE II score (B), the lactate level (C), and the patient survival rate (D). ** $P<0.01$ vs. the survival group.

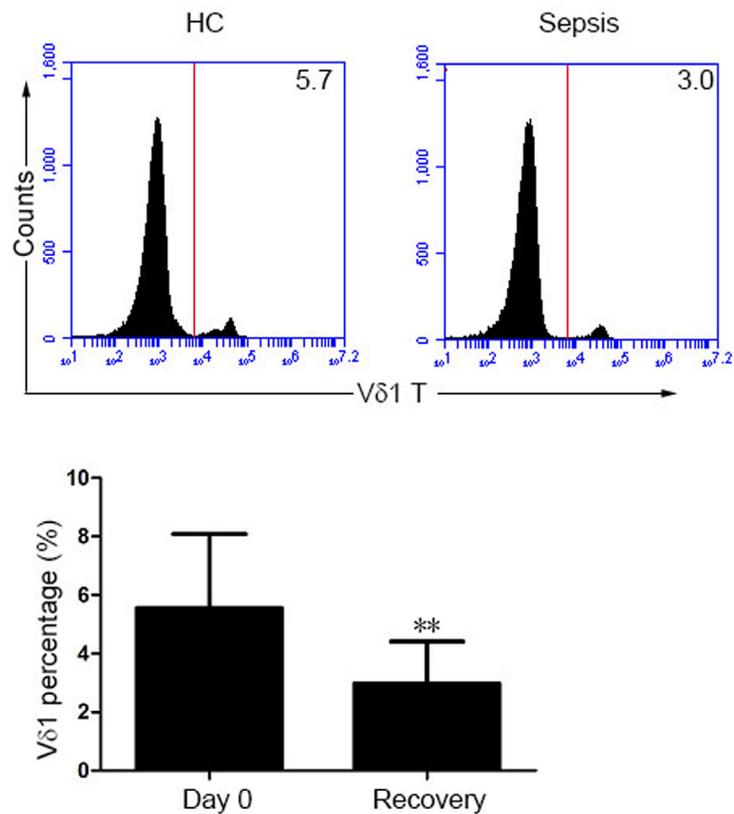


Figure 3. Proportion of Vδ1T cells in the peripheral blood of patients with sepsis after treatment. Flow cytometry was used to detect the proportion of Vδ1T cells in peripheral blood of septic patients before and after treatment (A). The proportion of Vδ1T cells significantly decreased after treatment (B). **P<0.01 vs. the Day 0 group.

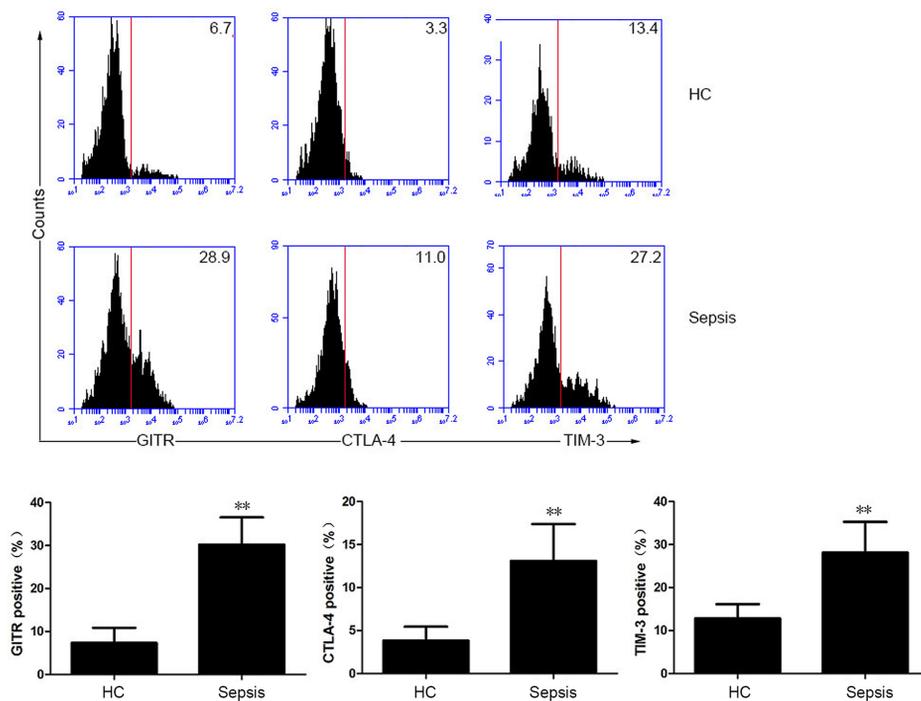


Figure 4. The expression of immunosuppressive molecules on Vδ1T cells in the peripheral blood of patients with sepsis. Flow cytometry was used to detect the expression of GITR, CTLA-4, and TIM-3 on Vδ1T cells in sepsis patients increased (Figure 4A-B). **P<0.01 vs. the Health control group. GITR, glucocorticoid-induced tumor necrosis factor receptor; CTLA-4, cytotoxic T lymphocyte-associated antigen-4; TIM-3, T cell immunoglobulin and mucin domain 3.

V δ 1T Cells Inhibited the Proliferation of and IFN- γ Secretion by CD4 T Cells

The proliferation level of CD4 T cells in the peripheral blood of HC was $95.2 \pm 6.83\%$, and it reduced to $70.7 \pm 7.92\%$ after the V δ 1T and CD4 T cells were incubated. The proliferation level of CD4 T cells in the peripheral blood of patients with sepsis was $99.5 \pm 8.14\%$, and it was changed to $51.1 \pm 8.01\%$ after the V δ 1T cells were added (Figure 5A). It indicated that the increase of V δ 1T cells in the sepsis patients had inhibitory effects on T cell proliferation. The IFN- γ secreted by CD4 T cells in the peripheral blood of the HC was $34.1 \pm 5.66\%$, and it was $21.4 \pm 6.21\%$ after the V δ 1T and CD4 T cells were incubated. The IFN- γ secreted by CD4 T cells in the peripheral blood of septic

patients was $35.4 \pm 8.82\%$, but it changed to $15.8 \pm 4.95\%$ in co-incubated V δ 1T and CD4 T cells from septic patients (Figure 5B). These results suggest that the immunosuppressive function of V δ 1T cells in the peripheral blood of septic patients is significantly higher than that of the healthy controls ($P < 0.01$; Figure 5).

DISCUSSION

$\gamma\delta$ T cells represent a minor population of T lymphocytes in human peripheral blood, which can directly recognize and bind antigens and play a crucial role in immune surveillance and regulation (31). In the healthy controls, V δ 1T cells are found predominately

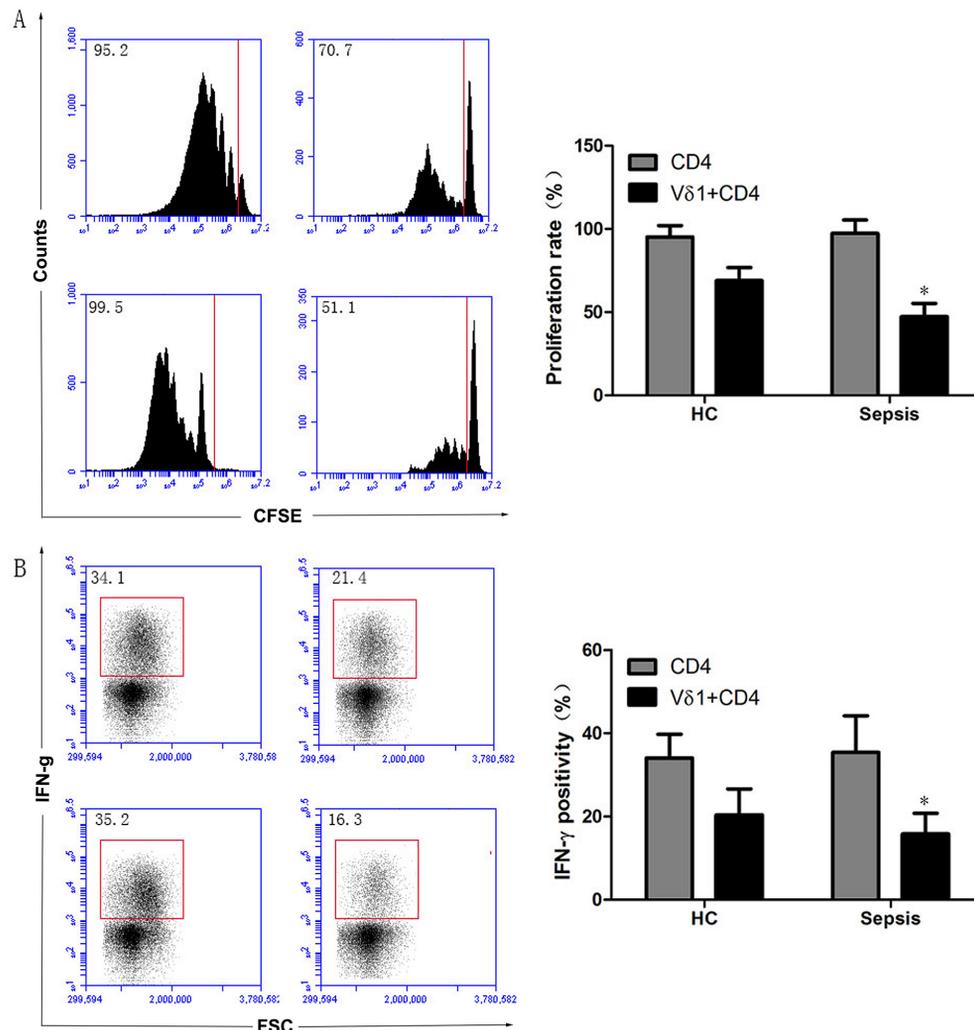


Figure 5. V δ 1T cells inhibited the proliferation of CD4 T cells. Flow cytometry showed that CD4 T and V δ 1T cells were incubated in peripheral blood of septic patients, CD4 T cell proliferation (A) and IFN- γ secretion ability (B) significantly decreased compared with CD4T cell culture alone. ** $P < 0.01$ vs. the healthy control group. IFN- γ , interferon- γ . The histograms are just representative examples of data.

at mucosal sites and can respond to non-classical MHC molecules expressed on stressed cells (32). Conversely, V δ 2T cells represent among 70% of circulating $\gamma\delta$ T cell subsets, and respond to phosphor antigens without MHC restriction (33). In this study, we found that the proportion of V δ 1T cells in the blood of septic patients is higher than that of HC, which positively correlated with the severity of the disease. After some effective treatment, the percentage of V δ 1T cells in the peripheral blood of patients with sepsis significantly reduced. In addition, the percentage of GITR-, CTLA-4-C, and TIM-3- positive V δ 1T cells in the blood of septic patients is significantly higher than that of the healthy controls, and the immunosuppressive function of V δ 1T cells in the peripheral blood of septic patients is significantly higher than that of the controls.

A large amount of experimental and clinical studies have shown that sepsis can cause remission of immunosuppression in the body, which in turn leads to priming infections, mainly opportunistic infections (28). Consistent with this evidence, the proportion of peripheral blood regulatory T (Treg) cells in the peripheral blood of patients with septic shock significantly increased (34). Wan et al. (2010) showed that during the early stages of sepsis, a significant proportion of abnormal Treg cells exists which is mainly manifested by an increase in the proportion and enhancement of immunosuppression (13). In addition, many studies have confirmed the abnormal proportion and function of Treg in septic patients. For instance, Huang et al. (2015) have found a significant increase in the proportion of CD39⁺Treg in the peripheral blood of septic patients, and the increase in the proportion of CD39⁺Treg in the peripheral blood of septic patients is closely related to the prognosis of patients. Shao et al. (2011) provided evidence that the proportion of CD4⁺CD25⁺Treg cells in the peripheral blood of septic patients significantly increased (14). The increase in the proportion of CD4⁺CD25⁺Treg cells can lead to a decrease

in CD3⁺, CD4⁺, and CD4⁺/CD8⁺ levels, which can inhibit immune function and participate in the pathogenesis of sepsis (15). In addition, the study by Pagel et al. (2016) confirmed that the proportion of CD4⁺CD25⁺Treg cells significantly increased in patients with clinical premature sepsis in preterm infants (16). In recent years, V δ 1T cells have been proven to play an immunosuppressive function in autoimmune diseases and tumor escape (23-26). Referring to the previous results, the role of Treg cells in the pathogenesis of sepsis, and the state of immunosuppression in patients with sepsis, we suspect that V δ 1T cells are involved in the development of sepsis and are associated with patient prognosis. Our results confirmed that the proportion of V δ 1T cells in the peripheral blood increased in septic patients, and shows a positive correlation with the severity of the disease. At the same time, the proportion of V δ 1T cells in septic patients was closely related to the patients' condition. These results are consistent with the changes observed in the immunosuppressive cells in most studies involving patients with sepsis. Based on the above-mentioned findings, we can preliminarily conclude that V δ 1T cells are closely related to the development and prognosis of sepsis. According to the conclusion of the comprehensive literature, the incidence of sepsis is not a result of a change in the proportion or function of immunosuppressive cells but related to a variety of immunosuppressive cells including Treg and V δ 1T cells, which are jointly related to the development of sepsis. Future regulatory DC cells and regulatory B cells that are not much concerned in sepsis patients will also be the focus of research in sepsis mechanisms.

Some studies have shown that in patients with tumors, the function of V δ 1T cells, in addition to the changes in proportion, mainly includes enhanced proliferation capacity and enhanced immune suppressive functions (35-37). The immunosuppressive state of septic patients is similar to that of those who have

tumors, and the changes in the function of V δ 1T cells in septic patients are also being researched. A recent study evaluated the characterization of V δ 1 and V δ 2 T cells in the peripheral blood of septic patients, indicating the involvement of imbalance and functional changes of V δ 1 and V δ 2 T in sepsis progression (38). This study indicated that the percentage of GITR-, CTLA-4-, and TIM-3-positive V δ 1T cells in the peripheral blood of septic patients is significantly higher than that of HC, and the inhibition of naïve CD4 T-cell proliferation and ability of V δ 1T cells in the peripheral blood to secrete cytokines in septic patients are significantly higher than that of the controls.

CONCLUSION

In summary, our study suggested that the proportion of V δ 1T cells in the blood of patients with sepsis significantly increased, and their function significantly enhanced. The immune function of septic patients is inhibited. Changes in the immune function of V δ 1T cells in septic patients may be closely related to the preconditioning of patients with sepsis.

ETHICAL APPROVAL

All procedures performed in studies involving human participants were in accordance with the ethical standards of the institutional and/or national research committee and with the 1964 Helsinki declaration and its later amendments or comparable ethical standards. This study is approved by the Ethics Committee of Huangshi Central Hospital of Edong Healthcare Group. Written informed consent was obtained.

AUTHOR'S CONTRIBUTION

Fawei Yuan carried out study concepts and design, experimental studies and data

acquisition; Huan Yin helped to manuscript preparation and editing; Juan Tan contributed to literature research and data analysis; Kun Zheng was the guarantor of integrity of the entire study and helped to clinical studies; Xiping Mei carried out definition of intellectual content and statistical analysis; Lixue Yuan helped to manuscript review.

Conflict of Interest: None declared.

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