Original Article TIGIT marks exhausted T cells and serves as a target for immune restoration in patients with chronic HBV infection

Yan-Yan Wei^{1,2*}, Jing Fan^{2*}, Meng-Xuan Shan², Dan-Dan Yin², Li-Li Wang², Wei Ye², Wei Zhao²

¹Department of Infectious Disease, The First Affiliated Hospital of Anhui Medical University, Hefei 230022, Anhui, China; ²Department of Clinical Research Center, The Second Hospital of Nanjing, Nanjing University of Chinese Medicine, Nanjing 210003, Jiangsu, China. ^{*}Equal contributors.

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Abstract: Background: Chronic HBV infection is a serious worldwide health problem that mainly causes liver cirrhosis and hepatocellular carcinoma (HCC). Few studies have explored how T cell exhaustion helps HBV avoid immune system attack and how to reverse that exhaustion. Recently, T cell immunoglobulin and immune receptor tyrosine-based inhibitory motif (ITIM) domain (TIGIT) have been identified as coinhibitory receptors, similar to PD-1. This study explores the expression of TIGIT and the T cell function changes in patients with chronic HBV infection. Results: In this study, we found that the expression of TIGIT on T cells increased significantly in patients with chronic HBV infection. High expression of TIGIT on T cells is associated with functional exhaustion. Importantly, this study demonstrates that blocking TIGIT can reverse T cell exhaustion and restore function in patients with chronic HBV infection. Conclusions: HBV induces T cell exhaustion by up-regulating the expression of TIGIT. Blocking the TIGIT/ PVR signaling pathway can reverse T cell exhaustion, so this discovery provides an immunotherapy target to battle chronic HBV infection.

Keywords: T cell immunoglobulin and ITIM domain, TIGIT, programmed death receptor-1, PD-1, T cell exhaustion

Introduction

Hepatitis B virus (HBV) is the most common hepatotropic virus that causes liver inflammation and injury. In the stage of acute viral infection, the response of T cells to HBV infection is characterized by the activation of large numbers of epitope-specific CD4⁺ and CD8⁺T cells [1, 2]. However, in the stage of chronic viral infection, the increase of antigen level and HBV viral load continue to stimulate T cells, leading to gradual loss of T cell function, which is defined as "T cell exhaustion" [3]. The "exhaustion" state of patients with chronic HBV infection is characterized by weak cytotoxic activity of effector cells, impaired production of cytokines, and continuous expression of multiple inhibitory receptors, such as cytotoxic T lymphocyte-associated antigen-4 (CTLA-4), lymphocyte activation gene-3 (LAG-3), T cell immunoglobulin-3 (TIM-3), and programmed cell death-1 (PD-1) [4-7].

When the immune system cannot eradicate an infection quickly, it will start a process of various actions to inhibit the infection and prevent immune-mediated tissue damage. This process is defined as exhaustion [8]. In patients with chronic HBV infection, the elevated expressions of PD-1, TIM-3, LAG-3, and CTLA-4 are associated with T cell exhaustion and persistent viral infection [8-12]. In addition, previous studies have demonstrated that blocking PD-1, Tim-3, LAG-3, and CTLA-4, whether alone or in combination, can improve the function of HBV-specific CD8⁺T cells in vitro [10-14]. It is worth noting that compared to other inhibitory receptors, the PD-1 pathway blockade leads to the strongest enhancement of T cell function in chronic HBV infection [11]. However, blocking the inhibitory receptors in patients with chronic HBV infection has only a limited effect on the recovery of HBVspecific CD8⁺T cell function [15], which indicates the necessity of exploring new inhibitory receptors.

TIGIT (T cell immunoglobulin and ITIM domain) was first identified as a new type of inhibitory receptor of CD28 family by bioinformatics algorithm in 2009 [16, 17]. TIGIT is a coinhibitory receptor of the Ig superfamily, and is specifically expressed on T cells and natural killer (NK) cells [17-19]. In recent years, studies have reported that blocking both TIGIT and PD-1 pathways can reverse T cell exhaustion in HIV-infected patients [20, 21]. However, there are few reports about the expression and function of TIGIT in patients with chronic HBV infection.

In this study, we analyzed the expression profile and function of TIGIT in peripheral T cells of patients with chronic HBV infection. We confirmed the effect of TIGIT blockade on the recovery of T cell function in patients with chronic HBV infection. Our study demonstrates that T cell exhaustion is the key to HBV immune escape, and blocking this checkpoint may reverse the functional impairment of T cells to HBV infection.

This study obtained approval from the Medical Ethics Committee of the Second Hospital of Nanjing in Jiangsu province of China (2017-LYkt021), and informed consent was obtained from each enrolled subject. This study had no influence on the subsequent management of patients.

Materials and methods

Patient sample collection and isolation

Heparinized peripheral blood was collected from patients with chronic HBV infection and healthy donors. Patients with chronic HBV infection were defined as HBV surface antigenpositive at least six months before enrollment without evidence of cirrhosis or hepatocellular carcinoma (HCC) according to the criteria established by the Chinese Medical Association (Chinese Society of Hepatology and Chinese Society of Infectious Diseases) [22]. These donors were considered healthy, had had no infection or immunity in the past month, had no known immunodeficiency or any history of chemotherapy or radiotherapy, and had not received systemic steroids or any other immunosuppressive drugs in the past 6 months. All samples were obtained in accordance with the requirements of the ethics committee of the Second Hospital of Nanjing and processed in

accordance with the principles of the International Conference on Harmonization Guidelines and the 1962 Declaration of Helsinki. All participants or their legal representatives voluntarily provided written informed consent before being included in the study. Venous blood was collected from each participant by venipuncture. Within 3 hours after collection, peripheral blood mononuclear cells (PBMCs) were extracted from blood samples with Lymphocyte separating solution [23]. Cell viability was detected by 0.4% trypan blue vital staining. All cells were cryopreserved in fetal bovine serum (FBS) containing 8% dimethyl sulfoxide (DMSO) and stored in liquid nitrogen until use.

T cell immunostaining and flow cytometric analysis

Cryopreserved PBMCs were thawed rapidly in warm 10% cRPMI (RPMI-1640) medium (GIBCO) supplemented with 10% FBS (GIBCO), 1% penicillin-streptomycin (GIBCO), 10 mM HEPES (GIBCO), 2 mM L-glutamine (GIBCO), and 10 µg/ mL DNase I (Sigma-Aldrich, Dorset, United Kingdom), washed with PBS+2% FBS (GIBCO) (complete RPMI). The cells were stained with aqua fluorescent reactive dye Dead Cell Stain Kit (AARD: Invitrogen, Carlsbad, California), and then incubated with panels of conjugated antihuman monoclonal antibodies (mAbs) in the dark at 4°C for 30 min. The following directly conjugated mAbs used in this study were obtained from eBioScience (Inc., San Diego, CA): CD3 eFluor450 clone (OKT3), CD8a APCeFluor® 780(OKT-8), TIGIT PE-Cyanine7 (MBS-A43), Mouse IgG1 K Iso control PE-Cyanine7, Mouse IgG1 APC, and PE-conjugated anti-CD28 (Cat. No. 2071). CD45RA FITC (LeuTM-18, Cat. No. 347723) was obtained from IMMUNOTECH. PMA/Ionomycin mixture (250×) was obtained from MultiSciences. All cells were washed with PBS+2% FBS and then placed in 1% paraformaldehyde (PFA, Electron Microscopy Sciences, Hatfield, Pennsylvania), and data were obtained on a CANTO II flow cytometer (BD Biosciences) within 18 hours. For each sample, 100,000 to 500,000 lymphocyte events were gathered. The isotype control or fluorescence minus one (FMO) samples were prepared for gating. Flowjo software version 10.0 (Treestar, Ashland, Oregon, USA) was used to analyze the data.

Isolation of PBMCs and stimulation of HLA-A2restricted HBV peptides in vitro

PBMCs were separated from fresh heparinized blood by Ficoll-Hypaque density gradient centrifugation, and then resuspended in RPMI 1640 with 25 mM HEPES, 50 µg/mL of gentamicin, 2 mM L-glutamine, and 8% human serum (complete medium). For the expansion of CD8⁺T cells, PBMC were resuspended in complete medium supplemented with concentration 2×10⁶/ml, sown at 200 µL/well into 96-well plates, and stimulated with HLA-A2-restricted HBV peptides (FLPSDFFPSV) in 10 µg/mL final concentration. Recombinant IL-2 (50 IU/mL), IL-7 (10 ng/mL), and IL-12 (10 ng/mL) were added on the 3rd and 6th days of culture, respectively. The immunological test was performed on the 10th day.

Intracellular staining and flow cytometry

PBMCs were cultured in RPMI-1640 medium (GIBCO) containing 10% FBS, stimulated with PMA/ionomycin (50 ng/mL and 1 µg/mL) or HBV peptides for 6 hours, and Golgi^{plug} (BD Biosciences) for 5 hours. The cell surface was stained with CD3 eFluor450 clone (OKT3), CD8a APC-eFluor[®] 780(OKT-8), and TIGIT PE-Cyanine7 (MBSA43); intracellular staining was done with IL-2 PerCP-eFluor[®] 710(MQ1-17H12), IFN-γ-PE (4S.B3), and TNF-α-FITC (MAb11). The cell viability was measured with Fixable Viability Dye eFluor 450 (eBioScience). Finally, the cells were analyzed by 8-color BD flow cytometry.

Antibody blockade of T cell function in vitro

Cryopreserved PBMCs (6×10⁵ cells) were blocked with anti-TIGIT mAb or IgG in wells of concave-bottomed 96-well plates (BD Falcon, Stockholm, Sweden) and cultured in a 5% CO incubator at 37°C for 40 min. Next, the peptides (10 µg/mL) were extracted from HBV epitope core-18 FLPSDFFPSV, env-183 FLLTRILTI, env-335 WLSLLVPFV, pol-455 GLSRYVARL peptides, and then put into the previous 96-well plates with concave bottom and incubated at 37°C for 96 hours in a 5% CO, incubator. Then, similar lyophilized peptides were added into the 96-well plates with concave bottom at 37°C for 10 hours in a 5% CO₂ incubator, and the Golgi^{plug} was added into the concave-bottomed 96-well plates at 37°C for 6 hours in 5% CO₂ incubator. Finally, the cells were washed and stained with LIVE/DEAD[®] Fixable Aqua Dead Cell Stain Kit and conjugated antibodies against CD3 and CD8 before intracellular staining of IFN- γ and IL-2 was carried out. As mentioned above, the cells were obtained by flow cytometry.

Statistical analysis

Prism Graphpad release 5.0 (Graphpad Software, San Diego, California) and SPSS 22.0 software (SPSS, Chicago, IL, USA) were used for statistical analysis. Under the assumption that the sample size and sample population obey the normal distribution, the independent-samples t test was selected to compare the two independent groups. In the event that the sample's size and population do not obey the normal distribution, the Mann-Whitney U test was selected to compare the two independent groups. Repeated measures were analyzed byone-way ANOVA attended by Tukey's multiple comparisons. Wilcoxon matched-pairs signed ranked test was used for pairing tests, and Spearman's rho test was used for correlation analyses. The significance measures are denoted in our tables and figures as *P<0.05, **P<0.01, or ***P<0.001.

Results

Expression of TIGIT on T cells in patients with chronic HBV infection

To investigate whether TIGIT is involved in the pathogenesis of chronic HBV infection, we detected the expression of TIGIT on T cells of peripheral blood mononuclear cells (PBMCs) in patients with chronic HBV infection and healthy donors (HD). 50 patients with chronic HBV infection (42 males and 8 females, median age 42.34 years, range 25-76 years) were selected as the study group, and 20 healthy donors (13 males and 7 females, median age 39.94 years, range 21-50 years) formed the control group. The mean percentage of CD4⁺T cells was 65.08±11.29% in the chronic HBV infection group and 67.87±11.14% in the HD group (P=0.249). The mean percentage of CD8⁺T cells was 33.02%±11.18% in the chronic HBV infection group and 30.63%±10.66% in the HD group (P=0.302). Details of participants' characteristics are shown in Table 1.

The percentage of TIGIT⁺CD4⁺T cells in the chronic HBV infection group was significantly

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Value	HBV-Uninfected Donors (HD; n=20)	Chronic hepatitis B (CHB; n=50)
Age (Years)	29.94±2.45	42.34±1.43
Gender (Male/Female)	13/7	42/8
ALT (IU/L)	37.23±5.72	32.18±3.05
AST (IU/L)	31.44±3.31	33.84±4.08
GGT (IU/L)	66.16±19.04	39.79±5.35
HBsAg	NA	1045.16±251.95
HBeAg	NA	23.40±17.21
HBcAb	NA	9.44±2.93
HBV viral load (copies/ml) Log10	NA	3.06±1.12
CD4 ⁺ T (%)	67.87±11.14	64.26±11.26
CD8+T (%)	30.63±10.66	33.81±11.17

 Table 1. Participant characteristics

CD8⁺T cells in the chronic HBV infection group. It should be noted that the expression of TIGIT in effector CD4⁺ and CD8⁺T cell subsets of patients with chronic HBV infection was the highest, followed by expression in memory CD4⁺T cell subsets (Effector, Memory, Naïve, and Intermediate CD4⁺T cell subset results: 18.92%±1.60%, 16.18% ±6.25%, 13.81%±1.88%, and 8.64%±0.80%, P<0.001) and CD8⁺T cell subsets (Effector, Memory, Naïve, and Intermediate CD8⁺T cell subset

higher than in the HD group ($12.28\pm0.93\%$ vs.7.98 $\pm0.86\%$, *P*=0.0083, **Figure 1A**, **1B**). In addition, the percentage of TIGIT*CD8*T cells in the chronic HBV infection group was significantly higher than in the HD group ($30.77\pm2.00\%$ vs. $16.61\pm2.17\%$, *P*=0.0001, **Figure 1A**, **1C**). These results suggest that the expression of TIGIT on CD4*T cells and CD8*T cells might be related to the pathogenesis of chronic HBV infection.

Analysis of TIGIT phenotype of T cell subsets in patients with chronic HBV infection

Previous studies have demonstrated that chronic HBV infection can lead to the expansion of immature effector CD4⁺ or CD8⁺T cells [24, 25]. Our study found that the expression of TIGIT on T cells in patients with chronic HBV infected was elevated. In order to analyze the expression of TIGIT in different T cell subsets, T cells were further divided into the Naive group, Memory group, Intermediate group, and Effector group according to the expression of CD28 and CD45RA [24, 25].

We analyzed the expression pattern of TIGIT in the above heterogeneous CD4⁺T cells and CD8⁺T cell subsets. **Figure 2A** shows a representative flow cytometry flow chart of coexpression of CD28 and CD45RA on CD4⁺ and CD8⁺T cells in chronic HBV infection. Compared to the HD group, TIGIT was highly expressed in four differentiation states of CD4⁺ and CD8⁺T cells in the chronic HBV infection group (**Figure 2B**, **2C**). In addition, we compared the expression of TIGIT in four subsets of CD4⁺ and results: 49.61%±14.66%, 37.62%±15.29%, 34.45%±9.65%, and 30.42%±14.13%, *P*< 0.001, **Figure 2B**, **2C**).

Furthermore, the expression of TIGIT on all of the Naive, Memory, Intermediate, and Effector subsets of T cells was significantly higher in the chronic HBV infection group than in the HD group. Additionally, the highest expression of TIGIT was found in the effector T cell subsets during chronic HBV infection, suggesting that the high expression of TIGIT may be related to the dysfunction of effector T cells in patients with chronic HBV infection.

Cytokine secretion by TIGIT⁺ and TIGIT⁻CD8⁺T cells after stimulation of HLA-A2-restricted HBV peptides

HBV-specific CTL can secrete antiviral cytokines such as IFN- γ , IL-2, and TNF- α , and it plays a role in virus clearance [8]. The dysfunction of cytokine secretion by exhausted HBV-specific CTLs in patients with chronic HBV infection could lead to the abnormal secretion of proinflammatory cytokines including decreases in IFN-y, IL-2, and TNF- α [8]. The PBMCs from chronic HBV infection patients were stimulated by HLA-A2-restricted HBV peptides, and the cytokine secretion by TIGIT⁺ and TIGIT⁻CD8⁺T cells were compared to study the effect of TIGIT on the cytokine secretion of CD8⁺T cells. The percentage of IFN-y secreting cells in the TIGIT⁺CD8⁺T cell group (1.16±0.14%) was significantly lower than in TIGIT-CD8+T cells stimulated by the antigenic peptides group (3.13±0.79%) after stimulation by HBV peptides. There was a significant difference bet-

TIGIT and exhausted T cells in HBV

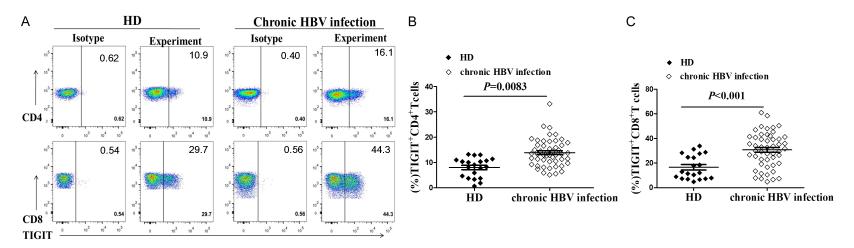


Figure 1. Expression of TIGIT in T cells elevated in chronic HBV infection. Cryopreserved PBMCs from 50 patients diagnosed with chronic HBV infection and 20 healthy donors were thawed and surface phenotypes were tested for TIGIT expression. (A) Representative flow cytometry flow plots showing TIGIT expression on CD4⁺ and CD8⁺T cells. Graphs show compiled data of TIGIT expression on (B) CD4⁺ and (C) CD8⁺T cells in the HBV-uninfected healthy donors group (HD, black diamond; n=20) and chronic HBV infection group (open diamond; n=50).

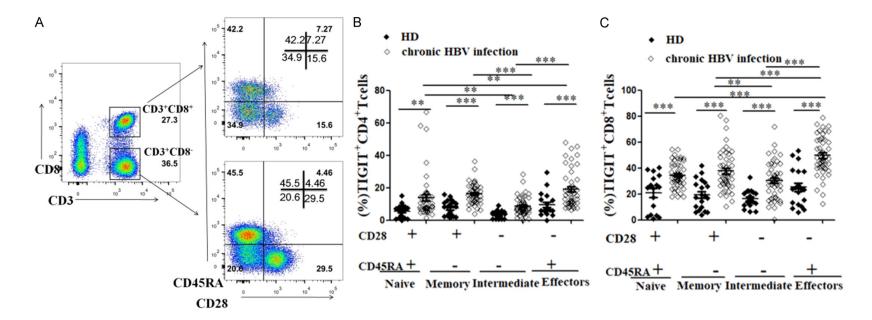


Figure 2. Expression of TIGIT is elevated at different stages of T cell differentiation in patients with chronic HBV infection. After thawing of frozen PBMCs, TIGIT expression on CD4⁺ and CD8⁺T cell subsets with different CD28 and CD45RA expression was detected. A. Representative flow cytometry flow plots show CD28 and CD45RA coexpression on CD4⁺ and CD8⁺T cells in chronic HBV infected patients. B. Graph shows compiled percentage of TIGIT⁺T cells in different CD4⁺T cell subsets in HD and chronic HBV infection groups. HBV-uninfected healthy donors (HD, black diamond; n=20), chronic HBV infection (open diamond; n=50). C. Graph shows compiled percentage of TIGIT⁺T cells in four CD8⁺T cell subsets in HD and chronic HBV infection groups. HBV-uninfected healthy donors (HD, black diamond; n=20), chronic HBV infection (open diamond; n=50). P values were calculated using independent-samples t test.

ween the two groups (P=0.0407, Figure 3A, **3D**). The percentage of TNF- α secreting cells in the TIGIT⁺CD8⁺T cells group (1.72±0.37%) was significantly lower than that in TIGIT the CD8⁺T cells stimulated by the antigenic peptides group (3.32±0.59%). There was a significant difference between the two groups (P=0.0087, Figure 3B, 3E). The percentage of IL-2 secreting cells in TIGIT⁺CD8⁺T cells stimulated by antigenic peptides group (5.73±0.86%) was significantly lower than that in TIGIT CD8⁺T cells stimulated by antigenic peptides group (8.91±1.48%) after stimulation of HBV peptides. There was a significant difference between the two groups (P=0.0446, Figure 3C, 3F). These results suggest that the expression of TIGIT on CD8⁺T cells stimulated by antigenic peptides might be related to the inhibition of cytokines secreted by CD8⁺T cells stimulated by antigenic peptides.

TIGIT blockade on T cells reversed the cytokine secretion of HBV-specific CD8⁺T cells

Boni et al. [2] indicated that the blockade of the PD-1/PD-L1 pathway can restore exhausted T cell function in patients with chronic HBV infection. Through previous studies, the authors found that the expression of TIGIT on the surface of CD8⁺T cells may lead to dysfunction of CD8⁺T cells in vitro. It is unclear whether blocking TIGIT signaling pathways can affect the cvtokine secretion of HBV-specific CTL in patients with chronic HBV infection. Therefore, we added anti-TIGIT mAb alone to PBMCs from 10 patients with chronic HBV infection to block TIGIT receptors in vitro. Then, the HBV peptide pool was added to the system and co-cultured with PBMCs for 96 hours. Finally, the expression of cytokines was detected by flow cytometry.

The percentage of IFN- $\gamma^{+}CD8^{+}T$ cells after stimulation by the HBV peptide pool in blocking the TIGIT signaling pathway group (1.40±0.12%) was significantly higher than of the Ig group (0.65±0.08%). There was a significant differ-

ence between the two groups (P<0.001, Figure **4A**, **4C**). The percentage of CD8⁺T cells secreting IL-2 in the group blocking TIGIT signaling pathway (2.03±0.16%) was significantly higher than that in Ig group (0.73±0.08%). There was a significant difference between the two groups (P<0.001, Figure **4B**, **4D**). Therefore, blocking the TIGIT signaling pathway in patients with chronic HBV infection partially reverses the cytokine secretion function of HBV-specific CD8⁺T cells.

Relationship between increased TIGIT expression on T cells and clinical values in patients with chronic HBV infection

To investigate the correlation between the expression level of TIGIT on T cells and clinical indexes in patients with chronic HBV infection, the expression level of TIGIT on T cells, liver function, and HBV viral load were detected in 50 patients with chronic HBV infection. The characteristics of these 50 patients are listed in **Table 1**.

The results show that the percentage of TIGIT⁺CD4⁺T cells was positively correlated with the level of alanine-transaminase (ALT), gamma-glutamyl transpeptidase (GGT) and HBV viral load, but not with the level of aspartateaminotransferase (AST) and HBsAg (Figure 5A-C and Supplementary Figure 1A, 1B). Similarly, the percentage of TIGIT⁺CD8⁺T cells was also positively correlated with the level of ALT and HBV viral load (Figure 5D, 5E), but not with the level of AST, GGT, and HBsAg (Supplementary Figure 1C-E). These results suggest that the expression of TIGIT on CD4⁺ and CD8⁺T cells in patients with chronic HBV infection positively correlates with liver inflammation and HBV viral load.

Discussion

The persistent expression of multiple inhibitory receptors in patients with chronic HBV infection

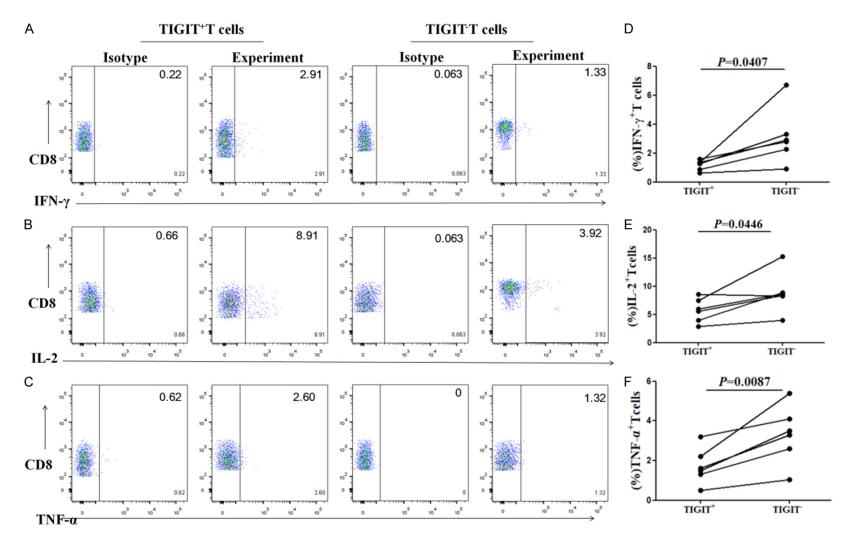


Figure 3. Secretion of IFN- γ , TNF- α , and IL-2 by TIGIT⁺CD8⁺T cells is lower than that of TIGIT⁻CD8⁺T cells after stimulation of HLA-A2-restricted HBV peptides. Representative flow cytometry flow plots show percentage of (A) IFN- γ , (B) IL-2, and (C) TNF- α expression on TIGIT⁺ and TIGIT⁻CD8⁺T cells after stimulation of HLA-A2-restricted HBV peptides. Graphs show the compiled percentage of (D) IFN- γ , (E) IL-2, and (F) TNF- α expression in TIGIT⁺ or TIGIT⁻CD8⁺T cells after stimulation of HLA-A2-restricted HBV peptides in chronic HBV infection group (n=6). *P* values were calculated using independent-samples t test.

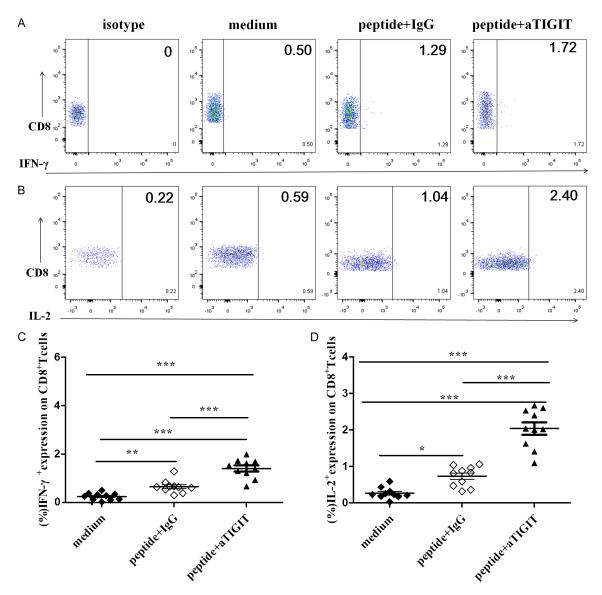


Figure 4. Effect of blocking with anti-TIGIT mAbs on HBV-specific CTL responses *in vitro*. In the presence or absence of mAb blocking antibodies, the HBV peptide pool was added to stimulate PBMCs from patients with chronic HBV infection *in vitro*. Representative flow cytometry flow plots gated on CD8⁺T cells, showing (A) IFN- γ and (B) IL-2 responses for medium, HBV peptide pool stimulation with IgG, HBV peptide pool stimulation with anti-TIGIT mAb from a chronic HBV-infected individual. Compiled data showing variation in the percentage of (C) IFN- γ^+ and (D) IL-2⁺ CD8⁺T cells in response to medium, HBV peptide pool stimulation with IgG, and HBV peptide pool stimulation with anti-TIGIT mAb (n=10). *P* values were calculated with repeated-measures one-way ANOVA test, followed by Tukey's multiple comparisons test.

is one of the characteristics of T cell exhaustion [1, 3]. There have been many studies on the expression of PD-1 and other inhibitory receptors on T cells of patients with chronic HBV infection [26-28]. TIGIT is a member of the immunoglobulin superfamily, especially expressed in immune cells, as a coinhibitory receptor [17-19]. Previous studies have demonstrated that TIGIT is expressed on activated T cells,

memory T cells, regulatory T (Treg) cell subsets, NK cells, and follicular helper T (Tfh) cells [16, 29, 30]. In addition to protecting against autoimmune diseases [31, 32], TIGIT also plays an important role in cancer and chronic viral infections [21, 33, 34]. In a recent study, TIGIT blockade or deficiency was found to lead to chronic hepatitis, fibrosis, and even HCC in HBsAgtransgenic mice in a CD8⁺T cell-dependent

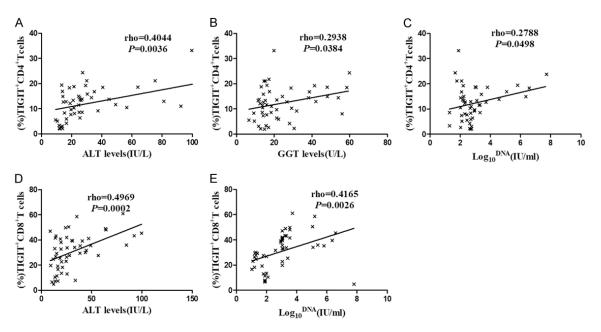


Figure 5. Relationship between the expression of elevated TIGIT on T cells and clinical parameters in patients with chronic HBV infection. Graphs show correlation between the percentage of TIGIT⁺CD4⁺T cells and (A) ALT (IU/L) levels, (B) GGT (U/L) levels, and (C) HBV viral load (\log_{10} IU/mI) levels in chronic HBV infection group (n=50). Graphs show correlations between the percentage of TIGIT⁺CD8⁺T cells and (D) ALT (IU/L) levels, and (E) HBV viral load (\log_{10} IU/mI) levels in chronic HBV infection group (n=50). Graphs show correlations between the percentage of TIGIT⁺CD8⁺T cells and (D) ALT (IU/L) levels, and (E) HBV viral load (\log_{10} IU/mI) levels in chronic HBV infection group (n=50). Spearman's rho tests were performed to detect correlations.

manner. The expression of the TIGIT gene in intratumoral regions of HCC patients is also higher than in paracancerous regions [35].

In this study, we studied the phenotype and function of T cells expressing TIGIT in patients with chronic HBV infection. The important findings of this study are as follows. First, the expression of TIGIT on T cells is elevated. Second, the expression of TIGIT on the Effector T cell subset was higher than that on the Naïve, Memory, and Intermediate T cell subsets. Third, the high expression of TIGIT on T cells was related to the decrease of cytokines secreted by CD8⁺T cells after stimulation by HBV peptides. Moreover, blocking the TIGIT signaling pathway partially restores T cell function. Finally, in patients with chronic HBV infection, increased TIGIT expression on T cells is positively correlated with liver inflammation and HBV viral load.

In patients with adult T cell leukemia [36], rheumatoid arthritis [37], HIV [38], and primary and metastatic melanoma [39], the expansion of T cell immune checkpoints may affect the skewness of T cell subsets. Speculating that TIGIT could play a dominant role in the upregulation of effector T cell subsets in patients with chronic HBV infection, we analyzed the expression pattern of TIGIT in T cell subsets (naïve T cells, Memory T cells, Intermediate T cells, and Effector T cells). We found that TIGIT was highly expressed in four differentiation states of CD4⁺ and CD8⁺T cells in the chronic HBV infection group compared with the HD group, respectively. In addition, the expression of TIGIT on the effector of CD4⁺ and CD8⁺T cell subsets was the highest in patients with chronic HBV infection.

Strikingly, this study found that TIGIT expression was relatively stable in the naïve CD8⁺T cell population, but there was no significant difference in TIGIT expression between the CD8⁺T cells naive subsets and intermediate subsets, and their expression values were lower than those of the effector subsets and memory subsets. Similarly, Chew et al. showed that TIGIT expression was relatively stable in the naïve population (CD28⁺CD45RA⁺CD8⁺T cells) [21].

In addition, TIGIT expression significantly increased on the CD8⁺T cell effector subsets, with the highest expression of TIGIT on the effector CD8⁺T cell subset, followed by interme-

diate and transitional subset values during chronic HIV infection. These results are consistent with a role for TIGIT as a potential regulator of intermediate, transitional, and effector T cell responses. However, we found a significant difference in TIGIT expression between the CD4⁺T cells naive subsets and effector subsets, with the naïve subset's value being lower than the effector subset. The reason might be that CD4⁺T cells include many cell types, including Tregs, Th1, Th2, Th17, and many others.

Previous studies have shown that the level of TIGIT expression can regulate CD4⁺T cell subsets. TIGIT is a newly identified coinhibitory receptor with up-regulated expression in effector Treg cells, and the marking Tregs specifically control Th1 and Th17 responses [40]. Additionally, gene expression analysis had shown that the frequency of TIGIT⁺CD4⁺T cells associated with a skewed CD4⁺T cell molecular profile indicates an exhausted Th1 profile [41]. TIGIT is critical for Th2 immunity. In vitro research has shown that the up-regulation of TIGIT expression in Th2 effector cells and its interaction with CD155 expressed in dendritic cells are important during the development of Th2 responses, blockade of TIGIT inhibited Th2 cells responses [42]. TIGIT is up-regulated in effector Treg cells, and this subset allows selectivity in inhibiting the response of Th1 and Th17 cells rather than Th2 cell response [30].

However, our study found a significant difference in TIGIT expression between CD4⁺T cell Naive subsets and Effector subsets, such that their values were lower than those of the Effector subset. Therefore, these results suggested that increasing TIGIT affects theskewing of T cell subsets, especially the Effector T cell subset. In other words, the high expression of TIGIT may be related to the dysfunction of Effector T cells in patients with chronic HBV infection. In addition to analyzing the general regulatory role of TIGIT as a coinhibitory receptor of CD8⁺T cells, the increased expression of TIGIT on CD4⁺T cells indicates a special regulatory role of TIGIT in shifting the cytokine balance, which has been reported in other studies and needs further study in patients with chronic HBV infection [6].

PD-1 is a receptor of the Ig superfamily. It has been demonstrated to play a major negative regulatory role in the T cell-mediated immune response [28]. Previous studies have found that PD-1 and other inhibitory receptors are coexpressed on T cells, leading to T cell exhaustion in chronic HBV infection [29, 33, 34]. In this study, the expression of TIGIT on CD4⁺ and CD8⁺T cells in the chronic HBV infection group was significantly higher than that of the HD group. Therefore, TIGIT may also be a marker of T cell exhaustion in patients with chronic HBV infection.

The synergistic effects of TIGIT expression on T cell function in AML patients [33], melanoma patients [43, 44], and HIV patients [21, 45] have been elucidated in the literature. Based on that, we speculated that the synergistic effect of TIGIT on T cells in patients with chronic HBV infection could inhibit the cytokines secreted by CD8⁺T cells. Therefore, we studied the effect of TIGIT expression on cytokine secretion of CD8⁺T cells in patients with chronic HBV infection in vitro, finding that the cytokine secretion of the TIGIT+CD8+T cell group was significantly reduced after stimulation of HBV peptides, suggesting that TIGIT could lead to the down-regulation of cytokine secretion of HBVspecific CD8⁺T cells. Moreover, blocking the TIGIT signaling pathway significantly increases the production of IL-2 in CD8⁺T cell groups stimulated by antigenic peptides-after stimulation by HBV peptides. It was deduced that anti-TIGIT mAb could partially restore the HBV-specific CD8⁺T cell response. This result is similar to previous reports [15, 21, 26, 46-48], but not exactly the same, so it needs to be confirmed with enlarged samples. Other similar studies have found that TIGIT is associated with T cell exhaustion in AML patients [33] or melanoma patients [43]. In this study, our data clearly demonstrate the key role of TIGIT in regulating T cell exhaustion in chronic HBV infection.

Chew et al. [21] and Tanaka et al. [36] illustrated that the viral load was associated with high expression of TIGIT in HIV patients. Therefore, considering the persistence of the TIGIT pathway under precise conditions of virus inhibition, blocking TIGIT and other inhibitor receptor signaling pathways can be used to enhance the response of CD8⁺T cells [21, 36]. Similar results were found in our study. The percentages of TIGIT⁺CD4⁺T cells and TIGIT⁺CD8⁺T cells were positively correlated with HBV viral load. Moreover, the percentage of TIGIT⁺T cells was positively correlated with ALT and GGT levels. Our results also indicate that the expression of TIGIT on CD4⁺ and CD8⁺T cells in patients with chronic HBV infection is positively correlated with liver inflammation and HBV viral load. However, the exact role of TIGIT in HBV clear-ance remains unclear.

Conclusions

This research elucidated the role of TIGIT in the pathogenesis of chronic HBV infection and dysfunction of T cells during chronic HBV infection. It is noted that blocking the TIGIT pathway can reverse T cell exhaustion and restore T cell function, which provides a basis for the treatment of chronic HBV infection.

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Disclosure of conflict of interest

None.

Address correspondence to: Drs. Wei Ye and Wei Zhao, Department of Clinical Research Center, Second Hospital of Nanjing, Nanjing University of Chinese Medicine, No. 1-1 Zhongfu Road, Nanjing 210003, Jiangsu, China. Tel: +86-13951729085; E-mail: yewei@njucm.edu.cn (WY); Tel: +86-13951-901772; E-mail: zhaoweiseu@126.com (WZ)

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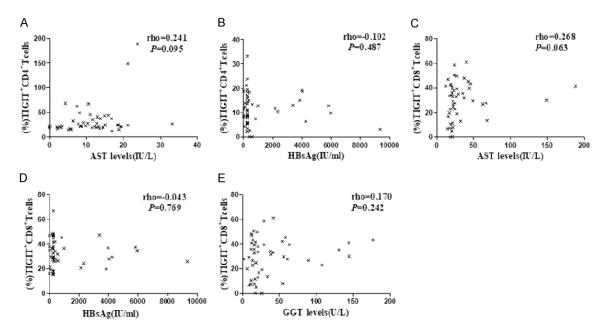
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Supplementary Fiugre 1. The relationship between the expression of elevated TIGIT on T cells and clinical parameters in patients with chronic HBV infection. Graphs show correlation between the percentage of TIGIT⁺CD4⁺ and (A) AST (IU/L) levels, and (B) HBsAg (IU/mL) levels in chronic HBV infection group (n=50). Graphs show correlation between the percentage of TIGIT⁺CD8⁺ and (C) ALT (IU/L) levels, (D) HBsAg (IU/mL) levels, and (E)GGT (U/L) levels in chronic HBV infection group (n=50). Spearman's rho tests were performed for correlations.