

ORIGINAL**The combination of CD8 and TIM3 expression predicts survival outcomes in hepatocellular carcinoma**

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Abstract : **BACKGROUND :** Tumor-infiltrating CD8+ T cells express many immune checkpoint proteins and are currently the focus of therapeutic targeting, although dramatic results have not yet been achieved in hepatocellular carcinoma (HCC). In this study, we performed a statistical analysis of immune checkpoint proteins and prognosis in HCC to evaluate its utility as a biomarker for predicting prognosis. **METHODS :** We analyzed 474 HCC patients with two comprehensive miRNA profiling datasets (GSE76427 : n=115 and TCGA : n=359) with clinical and transcriptomic data to examine associations between mRNA levels of CD8, TIM3, and PD-1/PD-L1. Each gene was divided into low- and high-expression groups using the tertiles of RNA expression. **RESULTS :** CD8 and TIM3 expression were associated with overall survival (OS) ($P=0.021$ and $P=0.025$, respectively), while PD-1/PD-L1 levels were not associated with OS ($P=0.306$ and $P=0.318$, respectively) in TCGA dataset. Among patients with high CD8 expression, low TIM3 expression was associated with better OS (TCGA : $P=0.009$, Whole patient cohort (TCGA+GSE dataset) : $P=0.008$). **CONCLUSIONS :** The combination of CD8 and TIM3 expression which indicates T cell exhaustion was useful biomarker for predicting HCC prognosis than PD-1/PD-L1, suggesting that TIM3 may be a future target for immune checkpoint inhibition. *J. Med. Invest.* 72: 354-360, August, 2025

Keywords : Hepatocellular carcinoma, CD8+ T cells, TIM3, Immune checkpoint inhibitor

INTRODUCTION

Among all cancers, hepatocellular carcinoma (HCC) ranks sixth in terms of incidence worldwide (1), making it the most common primary liver malignancy. The prognosis for hepatocellular carcinoma remains poor, treatments for early-stage HCC include radiofrequency ablation and surgical resection, while for advanced stages, treatments are limited to drug therapies and other options (2, 3).

Immune checkpoint inhibitors (ICIs) are effective against a number of cancers, including malignant melanoma and lung cancer. The discovery of new cancer drugs is also now focused on ICIs, and clinical trials of many ICIs are underway in HCC (4-6). Clinical trials with single-agent nivolumab and pembrolizumab, anti-programmed cell death-1 (PD-1) antibodies, have failed to demonstrate statistically significant efficacy (4, 5), but combination therapy with atezolizumab, an anti-programmed cell death ligand 1 (PD-L1) antibody, and bevacizumab, a vascular endothelial growth factor antibody, has shown efficacy (6) and been approved for the treatment of unresectable advanced HCC. However, ICIs are only effective in a subset of HCC patients, with the response rate to checkpoint inhibitors being approximately 15% to 20%; additionally, patients who initially respond to ICIs may develop resistance (7). Furthermore, it has been reported that approximately 33% of patients with melanoma who had an initial response to PD-1 inhibitors experienced tumor recurrence (8), suggesting that the improvement of T cell function with ICI may only be transient. One reason for this is that inhibition of a single immune checkpoint does not inhibit other types of immune

checkpoints, and therefore may be insufficiently effective (9). The most widely studied biomarkers for anti-PD-1/PD-L1 ICI therapy include PD-L1 expression, tumor mutation burden (TMB), and expression patterns of immune-related genes (10, 11). However, due to the low TMB and PD-L1 expression rates in HCC, there are no biomarkers to accurately predict ICI response or prognosis in HCC (12, 13). Therefore, further studies using ICIs other than anti-PD-1/PD-L1 antibodies are needed.

CD8+ T cells have recently been reported to play an important role in anti-tumor immunity, and high CD8 expression in tumor tissues has been associated with better overall survival (OS) in HCC patients (14). However, when antigen exposure is prolonged in the tumor environment, CD8+ T cells are more likely to differentiate into a stage known as "T cell exhaustion". These exhausted CD8+ T cells lose their ability to produce cytokines (IL2, TNF- α , IFN- γ), and their ability to suppress tumor cells is also reduced (15). It has also been reported that multiple immune checkpoint proteins, including T cell immunoglobulin and mucin domain-containing-3 (TIM3), are expressed on exhausted CD8+ T cells to suppress anti-tumor immunity (16-18) and that their expression levels are positively correlated with the severity of exhaustion (19, 20). As the exhausted T cells showed a decrease of effector cytokine production and cytolytic activity, leading to the tumor progression, TIM3 is one of the most important exhaustion markers. Therefore, in this study, we focused on exhausted T cells and investigated the role of TIM3 and CD8 expression as a prognostic marker using statistical analysis of a comprehensive dataset.

MATERIALS AND METHODS*Patient cohort and comprehensive analyses*

Both clinical and genomic data were obtained from two public datasets (GSE76427 and the Cancer Genome Atlas [TCGA]), and we analyzed gene expression profiles (RNA sequences). Samples

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without gene expression data and survival longer than 3000 days were excluded. In this study, 474 HCC patients with two comprehensive miRNA profiling datasets (GSE76427 : n=115 and TCGA : n=359) were analyzed. CD8, TIM3, PD-1, and PD-L1 expression were assessed for associations with survival. The patient data used in this study are all de-identified.

Statistical analyses

All statistical analyses were performed using JMP Pro 17 (SAS Institute, Cary, NC, USA). Patients were divided into low- and high-expression groups using the tertiles for RNA expression levels of CD8, TIM3, PD-1, and PD-L1 (CD8 and TIM3 : high = upper 1/3 tertile, PD-1 and PD-L1 : high = upper 2/3 tertile). Survival outcomes were analyzed using Kaplan-Meier plots and the log-rank test. A p-value <0.05 was considered statistically significant.

RESULTS

Relationship between the expression of CD8 and immune checkpoint proteins with survival outcomes

To examine the clinical relevance of CD8+ T cell infiltration in HCC, we first examined the association between CD8 expression and survival outcomes. Patients were divided into low and high groups using the tertiles for CD8 expression in the public datasets. There was a significant difference between high CD8 expression and better survival outcomes in TCGA cohort (P=0.021, Fig. 1A). Also, another dataset shows that there was a significant difference between high CD8 expression and better survival outcomes in GSE76427 cohort (P=0.014, Fig. 1B). As the exhausted T cells showed a decrease of effector cytokine production and cytolytic activity, TIM3 is one of the most important exhaustion markers. Then, to investigate the exhaustion of T cells, we focused on the immune checkpoint proteins (TIM3 and PD-1/PD-L1 expression). There was a significant difference between low TIM3 expression and better survival outcomes (TCGA : P=0.025, Fig. 1C ; GSE76427 : P=0.009, Fig. 1D). Interestingly, there were no differences between PD-1 or PD-L1 expressions with survival outcomes (PD-1 TCGA : P=0.306, Fig. 1E ; GSE76427 : P=0.565, Fig. 1F ; PD-L1 : TCGA : P=0.318, Fig. 1G ; GSE76427 : P=0.612, Fig. 1H). In summary, infiltration of CD8+ T cells into the tumor tissue and low TIM3 expression were significantly associated with better survival outcomes in patients with HCC using the comprehensive dates.

Role of the immune checkpoint proteins with survival outcomes among the patients with high CD8 expression

Next, to investigate the role of the relationship between CD8 expression and immune checkpoints, we examined the relationship between TIM3 and PD-1/PD-L1 levels and OS in patients with high CD8 expression (n=239). For TIM3, patients with low TIM3 expression and high CD8 expression had better OS than those with high TIM3 expression (P=0.009, Fig. 2A). Regarding PD-1/PD-L1, no association was found between PD-1 or PD-L1 expression and OS under CD8 high expression (P=0.491 and P=0.772, Fig. 2B, 2C). Univariate and multivariate Cox proportional hazard regression analysis for overall survival revealed that TIM3 levels are most relevant to prognosis (Supplementary Figure). Because CD8 and TIM3 levels are expected to be highly associated with OS, we analyzed the association of the combination of CD8 and TIM3 expression with OS among all HCC patients. High CD8 and low TIM3 expression were associated with improved OS compared with the overall HCC patient population (P=0.008, Fig. 3A). Conversely, low CD8 and high TIM3 expression were associated with significantly worse OS compared with

the overall HCC patient population (P<0.001, Fig. 3B).

Additionally, the group with high CD8 and high TIM3 expression did not have significantly different survival outcomes compared with the group with low CD8 expression (P=0.132, Fig. 4). In other words, regardless of CD8-positive cell infiltration in the tumor, if TIM3 was highly expressed and CD8+ T cells were fatigued, there were no differences in OS from patients with low levels of tumor-infiltrating CD8+ T cells. In summary, regardless of PD-1/PD-L1, tumor-infiltrating CD8+ T cells with low TIM3 expression were associated with a good prognosis, and low TIM3 expression was essential for a good prognosis in HCC.

DISCUSSION

CD8+ T cells are the most important component of anti-tumor immunity. CD8+ T cells directly attack tumor cells *via* perforin and granzymes and induce apoptosis in tumor cells *via* cytokines such as IFN- γ , IL-2, and TNF- α . However, CD8+ T cells usually enter a state of "exhaustion" or "dysfunction" in the tumor microenvironment (TME) (21). CD8+ T cells in the TME express high levels of immune checkpoint proteins such as PD-1, TIM3, and CTLA4 and have reduced release of cytokines and cytotoxic potential and increased expression of IL-10, a cytokine that suppresses immunity (22). Additionally, PD-L1 and galectin9, ligands for the immune checkpoints PD-1 and TIM3, respectively, are primarily expressed on HCC tumor cells and CD68-positive tumor-associated macrophages (TAMs) to promote immune escape (19, 23). Clarifying such mechanisms of the TME, in which tumor-associated macrophages and regulatory T cells are heavily involved, may be important for improving the outcome of ICI therapy (24).

Binding of TIM3 to its ligand galectin-9 inhibits the production of IL-2, TNF α , and IFN- γ , inhibiting the immunity of Th17, Th1, and Tc2 cells (25). In HCC tumor immunity, there is a marked increase in TIM3 expression on CD8+ T cells (19, 20, 26), which suppresses anti-tumor immunity by promoting TME formation and inhibiting T cell function (27, 28). In HCC patients, TIM3 expressions on CD8+ T cells and TAMs and serum TIM3 levels are significantly correlated with shorter OS and more advanced tumor stage (17, 26-28). Moreover, T cells with high TIM3 expression are increased among peripheral blood mononuclear cells (PBMCs) and are also increased in patients with recurrent HCC after hepatectomy as well as in those with treatment-resistant HCC (29). These findings suggest that TIM3 may be an important marker to predict OS and treatment resistance or relapse in HCC. Although there is currently no standard therapy targeting TIM3, anti-TIM3 antibodies have shown anti-tumor effects in preclinical studies (19, 20, 29), and early-stage clinical trials with various anti-TIM3 antibodies are currently underway worldwide (30, 31).

Our statistical analysis showed that regardless of PD-1/PD-L1 expression, high CD8 and low TIM3 expression were associated with better OS in HCC (Figs. 1, 2, 3). We found that both low TIM3 and high CD8 expression were essential for a good prognosis in HCC (Fig. 4). This suggests that TIM3 may be a more useful biomarker for predicting HCC prognosis than PD-1/PD-L1, which is currently a therapeutic target.

In this study, we analyzed mRNA expression levels of CD8 and several immune checkpoint molecules using publicly available datasets. While our findings highlight gene expression patterns that may reflect underlying immune activity and potential therapeutic targets, an important limitation is the lack of direct clinical validation. Future studies integrating protein-level analysis, immunohistochemistry, and patient outcome data are needed to confirm the prognostic and predictive value of these

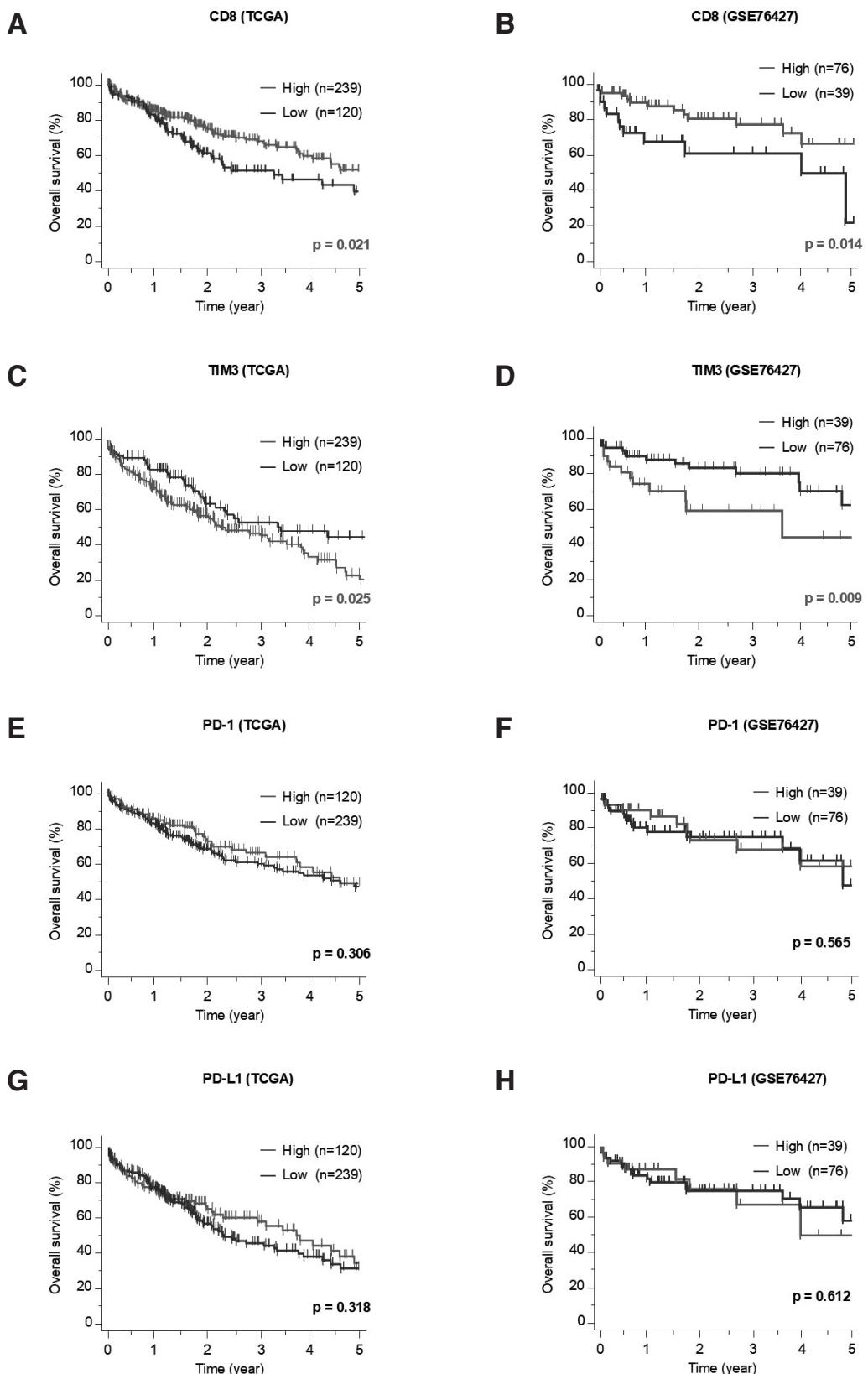


Figure 1. Relationship between the expression of CD8 and other immune checkpoint proteins with OS in patients with hepatocellular carcinoma.

A-B) A comparison of OS between high and low-risk group estimated by the CD8 expression in (A) TCGA : n=359, and (B) GSE76427 : n=115 dataset. C-D) A comparison of OS between high and low-risk group estimated by the TIM3 expression in (C) TCGA : n=359, and (D) GSE76427 : n=115 dataset. E-F) A comparison of OS between high and low-risk group estimated by the PD-1 expression in (E) TCGA : n=359, and (F) GSE76427 : n=115 dataset. G-H) A comparison of OS between high and low-risk group estimated by the PD-L1 expression in (G) TCGA : n=359, and (H) GSE76427 : n=115 dataset. Survival outcomes were analyzed using Kaplan-Meier plots and the log-rank test. A p-value <0.05 was considered statistically significant.

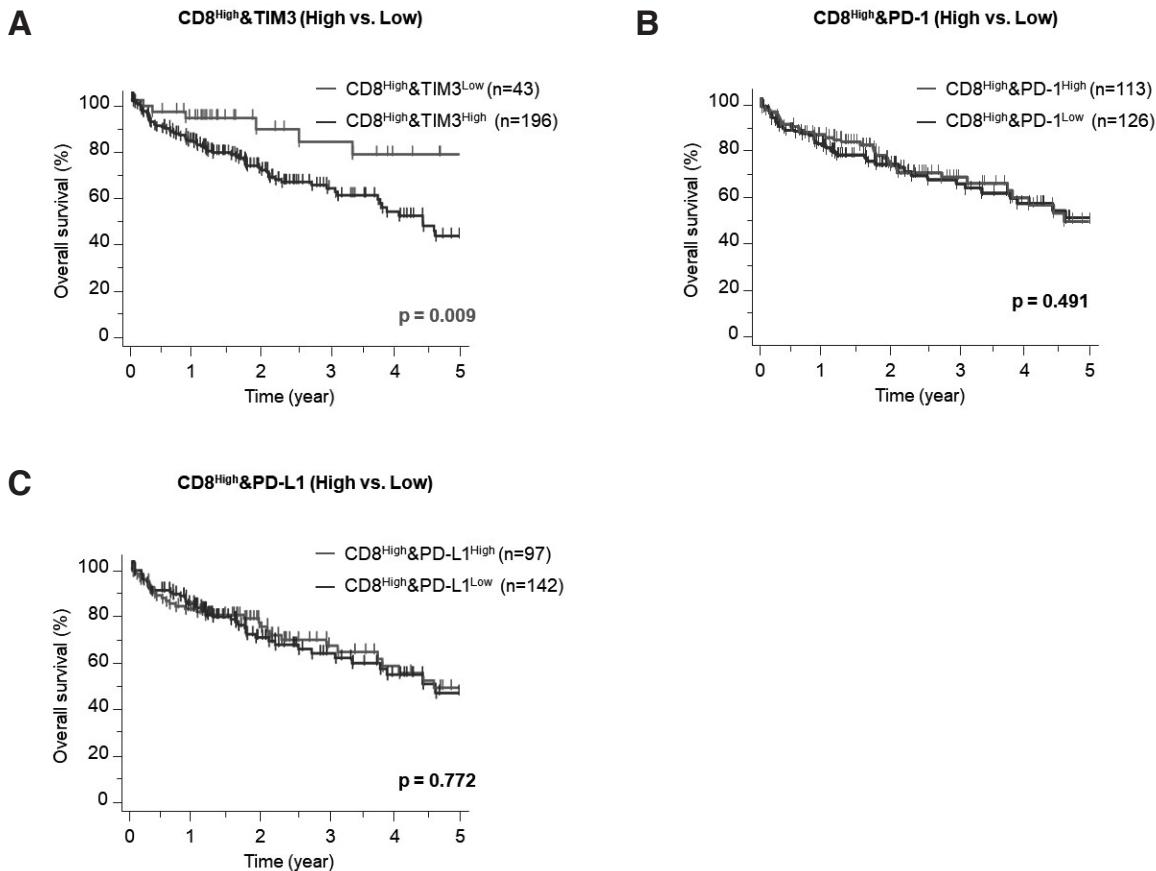


Figure 2. Relationship between the expression of immune checkpoint proteins and OS among the patients with high CD8 expression.

A-C) A comparison of OS between high and low-risk group estimated by the (A) TIM3, (B) PD-1, and (C) PD-L1 expression among the patients with high CD8 expression in TCGA dataset. Survival outcomes were analyzed using Kaplan-Meier plots and the log-rank test. A p-value <0.05 was considered statistically significant.

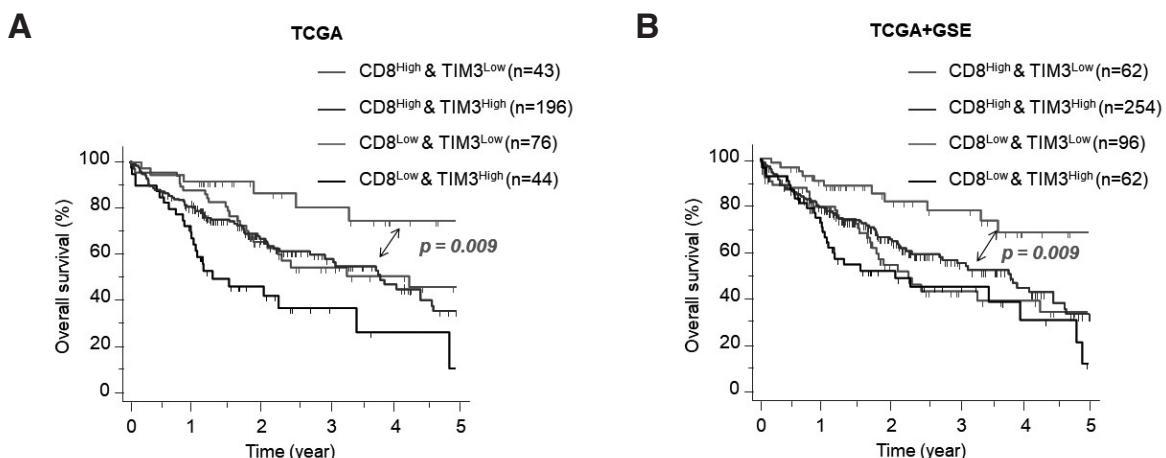


Figure 3. The combination between the CD8 and TIM3 expression with OS in patients with hepatocellular carcinoma.

A-B) A comparison of OS between high and low-risk group combined with CD8 and TIM3 expression in (A) TCGA : n=359, and (B) whole dataset (TCGA+GSE76427 : n=474). Survival outcomes were analyzed using Kaplan-Meier plots and the log-rank test. A p-value <0.05 was considered statistically significant.

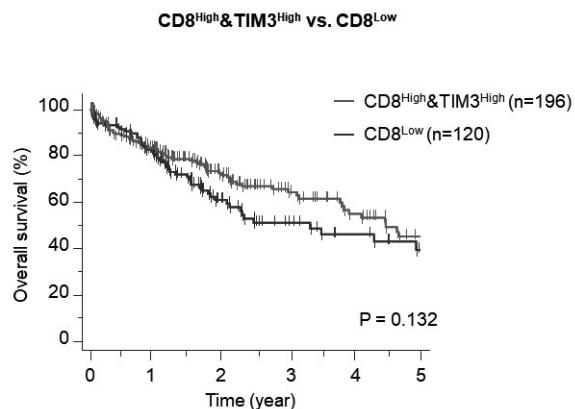


Figure 4. Relationship between the high CD8 and high TIM3 expression compared with low CD8 expression cohort in OS. A comparison of OS between the high CD8 and high TIM3 expression (exhausted CD8+ T cells) compared with low CD8 expression cohort in TCGA dataset. Survival outcomes were analyzed using Kaplan-Meier plots and the log-rank test. A p-value <0.05 was considered statistically significant.

immune markers. In this study, we analyzed mRNA levels of CD8 and immune checkpoint proteins from TCGA dataset and not from single cell RNA sequencing of CD8+ T cells. In the future, we would like to examine the RNA expression levels of CD8+ T cells, where TIM3 data would reflect the actual expression levels of cells and receptors *in vivo*, to examine the relationship between the real expression levels of immune checkpoint proteins in CD8+ T cells with prognosis in patient samples to confirm the results of this study. It has also been reported that blocking PD-1 increases the expression of other immune checkpoint proteins such as TIM3, CTLA-4, and LAG-3 in tumor-infiltrating immune cells (32). Co-expression of PD-1 and TIM3 on CD8+ T cells in the TME has been observed in several tumor types (33, 34), and combined inhibition of both pathways may have synergistic anti-tumor effects (32, 35). We believe that further research into the clinical application of TIM3, which was found to be statistically useful as a prognostic predictor for HCC in this study, and a combined strategy to inhibit the PD-1 and TIM3 pathways in HCC should be explored in the future.

CONCLUSION

Our results show that the patients with high CD8 expression was significantly better OS, reflecting active anti-tumor immunity, while the patients with high TIM3 expression was significantly worse OS, indicating the fatigue of anti-tumor immunity in CD8+ T cells in the patients with HCC. High CD8 and low TIM3 expression (non-exhausted CD8+ T cells) was associated with better OS in HCC than PD-1/PD-L1, suggesting that CD8 and TIM3 are useful biomarkers for predicting HCC prognosis. Restoring exhausted T cells via the suppression of TIM3 is a promising strategy for cancer treatment, which has become a breakthrough in cancer immunotherapy.

DISCLOSURES

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CONFLICT OF INTEREST

The authors declare that they have no conflict of interest.

HUMAN/ANIMAL RIGHTS

All procedures followed were in accordance with the ethical standards of the responsible committee for human experimentation (institutional and national) and with the Helsinki Declaration of 1975, as revised in 2008(5).

INFORMED CONSENT

Informed consent was obtained from all patients to be included in the study.

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No funding was received.

AVAILABILITY OF DATA AND MATERIALS

The data generated in the present study are included in the materials and methods section of this article.

AUTHORS' CONTRIBUTIONS

TN, YW, MN and MS conceived the study idea and designed the study. YM, TI, and YM provided administrative support and study materials or patients. TN, YW, MN, HT, SY, YS, TI, YM and MS collected and assembled the data. TN, YW and MS analyzed and interpreted the data. All authors contributed to the writing of the manuscript. MN, HT and SY confirm the authenticity of all the raw data. All authors have read and approved the final version of the manuscript.

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Factors	Univariate		Multivariate		
	3-year OS (%)	P value	HR	CI	P value
PD-L1 (- / +)	68.6 / 68.2	0.99	0.83	0.49-1.40	0.47
PD-1 (- / +)	66.4 / 67.5	0.80	0.75	0.37-1.51	0.42
TIM-3 (- / +)	83.5 / 63.9	0.02	0.36	0.16-0.79	0.01

Supplementary Figure. Univariate and multivariate Cox proportional hazard regression analysis for overall survival