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Reduced Siglec-7 expression on NK cells predicts NK cell dysfunction in primary hepatocellular carcinoma

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Summary

Major histocompatibility complex class I (MHC-I)-dependent inhibitory receptors on natural killer (NK) cells have been found to contribute to NK cell dysfunction in hepatocellular carcinoma (HCC). However, the roles of MHC-I-independent inhibitory receptors on NK cells in HCC remain poorly defined. In this study, we analyzed the expression of the MHC-Iindependent inhibitory receptors sialic acid-binding immunoglobulin-like lectin (Siglec)-7 and Siglec-9 on NK cells by analyzing the peripheral blood of 35 HCC patients and 63 healthy donors. We observed that HCC patients had lower frequencies and total numbers of NK cells in the peripheral blood. Importantly, both the expression levels of Siglec-7 on NK cells and the frequencies of Siglec-7⁺ NK cells were significantly reduced in HCC patients, which was accompanied by a decrease in activating receptor and an increase in inhibitory receptor expression on NK cells. Moreover, Siglec-7+ NK cells expressed higher levels of activating receptors and displayed stronger effector functions, compared with Siglec-7⁻ NK cells. Our findings demonstrate for the first time that reduced Siglec-7 expression predicts NK cell dysfunction in HCC patients, suggesting that Siglec-7 may be a potential marker of functional NK cell subset in HCC patients.

Keywords: hepatocellular carcinoma, natural killer cells, Siglec-7

Introduction

Hepatocellular carcinoma (HCC) is currently the fifth most common cancer but the second leading cause of cancer-related deaths worldwide [1]. Unfortunately, only a limited choice of effective therapeutic regimens for HCC exist to date. Increasing evidence indicates that immunotherapy may represent a promising strategy for HCC treatment [2]. The human liver is an organ enriched in diverse lymphocyte populations, of which natural killer (NK) cells account for 25-50% [3]. NK cells have potent anti-tumor activity through direct cell lysis and/ or cytokine production. Accumulating studies have shown that the decrease in NK cell numbers and their dysfunction (also called NK cell exhaustion) is associated with the progression and poor clinical outcome of HCC [4]. This suggests that NK cell-based immunotherapies to improve NK cell effector function including, but not limited to, checkpoint blockade, cytokine therapy or NK cell adaptive transfer, have the potential to be applied

to HCC treatment. However, the lack of detailed knowledge relating to the exact mechanisms underlying NK cell exhaustion has deterred the use of NK cell immunotherapy in HCC.

NK cells comprise ~15% of total lymphocytes in the human peripheral blood mononuclear cell (PBMC) population, and are divided into two subsets (CD56^{bright}CD16⁻ and CD56^{dim}CD16⁺), according to the expression of CD56 (dim or bright) and CD16 (presence or absence). CD56^{dim}CD16⁺ NK cells account for ~90-95% of the total NK cell populaexhibit higher cytotoxic activity than tion and CD56^{bright}CD16⁻ NK cells [5]. NK cell function is regulated by (i) activating receptors, such as natural killer group (NKG)2D, NKp46, NKp44 and NKp30; and (ii) inhibitory receptors such as NKG2A, lymphocyte-activation gene (LAG-3), T cell immunoglobulin mucin-3 (TIM-3), T cell immunoglobulin and immunoreceptor tyrosine-based inhibition motif (ITIM) domain (TIGIT) and CD96 [6]. Healthy cells protect themselves against NK cell-mediated killing via the interaction between their self-major histocompatibility

complex class I (MHC-I) molecules and the inhibitory receptors on NK cells [7]. This strategy is also utilized by cancer cells to escape NK cell-mediated anti-tumor immunity [8]. Recently, accumulating studies have demonstrated that MHC-I-independent inhibitory receptors also contribute to NK cell dysfunction during cancer progression [9,10]. Deciphering the roles of these inhibitory receptors in cancer pathogenesis will therefore enhance our understanding of the mechanisms implicated in NK cell exhaustion and benefit the development of effective HCC therapies.

Sialic-acid-binding immunoglobulin-like lectin-7 (Siglec-7) belongs to the CD33-related Siglec family, predominantly expressed on human NK cells [11]. The cytoplasmic domain of Siglec-7 has an ITIM, which defines its action as an inhibitory receptor on NK cells [12]. Although the ligation of Siglec-7 with specific antibodies or ligands inhibits NK cell function [13], Siglec-7 displays a similar expression pattern to that of NK cell-activating receptors. The down-regulation of Siglec-7 on NK cells, which represents an early marker of NK cell dysfunction, has been observed during chronic viral infection [14,15]. The expression of Siglec-7 is also reduced on NK cells in obese humans, whose NK cell anti-tumor activity is impaired [16,17]. Thus, cell surface Siglec-7 expression defines a highly functional NK cell phenotype, despite its role in suppressing NK cell activity. However, the changes in Siglec-7 expression experienced by NK cells and their association with the clinical outcomes of HCC remain largely unknown.

In this study, we observed that both the level of Siglec-7 expression on NK cells and the frequency of Siglec-7⁺ NK cells were significantly reduced in HCC patients, which was accompanied by a decrease in activating receptor and an increase of inhibitory receptor expression on NK cells, as well as overall NK cell functional impairment. Therefore, reduced Siglec-7 expression on NK cells predicts circulating NK cell dysfunction in HCC patients, suggesting that Siglec-7 may be a potential marker of functional NK cell subset in HCC patients.

Materials and methods

Human samples

None of the HCC patients selected for this study were receiving chemotherapy or radiotherapy. Venous blood was collected from 35 untreated patients with primary HCC and 63 healthy donors (HDs) between 2017 and 2019. Patients were recruited from the First Affiliated Hospital of Anhui Medical University. This study was approved by the Medical Ethics Committee of the First Hospital of Anhui Medical University, and the protocols were carried out in accordance with the approved guidelines. All participants gave written informed consent to participate in the study.

Isolation and activation of PBMCs

PBMCs were isolated from the peripheral blood of HCC patients or HDs by gradient centrifugation with Ficoll-Paque Plus (Solarbio, Beijing, China), according to the manufacturer's instructions. To assess intracellular cytokine production, PBMCs were isolated and stimulated with phorbol 12-myristate 13-acetate (PMA), ionomycin and monensin for 4 h.

Flow cytometry

Single-cell suspensions were stained with the fluorescently-labeled antibodies, including fluorescein isothiocyanate (FITC)-anti-human CD3 (clone OKT3), Brilliant Violet 421 anti-human CD56 (clone 5.1H11), phycoerythrin (PE) anti-human CD366 (TIM-3; clone F38-2E2), peridinin chlorophyll/cyanine 5.5 (PerCP/Cy5.5) antihuman CD337 (NKp30; clone P30-15), PE/Cy7-CD335 (NKp46; clone 9E2), PE anti-human CD328 (Siglec-9; clone K8), APC-anti-human CD328 (Siglec-7; clone 6-434), APC/Fire[™] 750 anti-human CD16 (clone 3G8), PE-anti-human tumor necrosis factor (TNF)- α (clone MAb11), PerCP/Cy5.5 anti-human CD107a [lysosomalassociated membrane protein 1 (LAMP-1); clone H4A3] and PE/Cy7 anti-human interferon (IFN)- γ (clone B27). Isotype controls were used to determine cut-off levels for positive staining. All antibodies and isotypes were purchased from Biolegend (San Diego, CA, USA). Data were acquired using a flow cytometer (Beckman Coulter; Brea, CA, USA) and analyzed using FlowJo version 10 software (Tree Star; Ashland, OR, USA).

Statistical analysis

Data are shown as means \pm standard deviation (s.d.). The difference between groups was analyzed using a paired- or unpaired-samples *t*-test. Any potential correlations between variables were analyzed using Pearson's correlation test. Analyses yielding *P*-values of less than 0.05 were considered statistically significant; **P* < 0.05; ***P* < 0.01; ****P* < 0.001; *****P* < 0.0001. Statistical analyses were performed using PRISM software, version 6 (GraphPad).

Results

Abnormal frequencies and functions of NK cells observed in the peripheral blood of HCC patients

To analyze NK cells in the peripheral blood, $CD3^-CD56^+$ NK cells were first identified within the lymphocyte gate and then subdivided into $CD56^{dim}CD16^+$ and $CD56^{bright}CD16^-$ NK subsets (Fig. 1a). Our results show that the frequencies of total NK cells in the peripheral

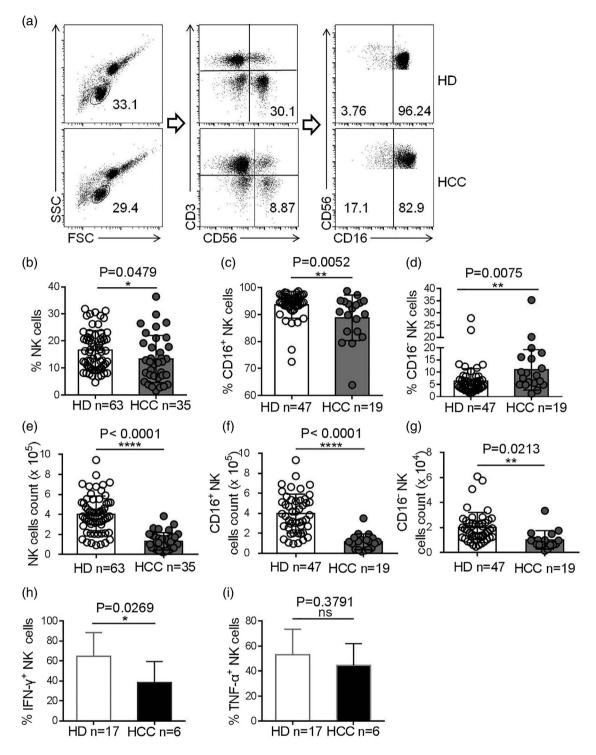


Fig. 1. Lower natural killer (NK) cell numbers detected in the peripheral blood of hepatocellular carcinoma (HCC) patients. (a) The gating strategy used for the cytometric analysis of NK cell subsets in peripheral blood mononuclear cell (PBMC) samples. (b–d) Frequencies of NK cells within the lymphocyte population (b), CD16⁺ NK cell subset (c) and CD16⁻ NK cells subset (d) in total NK cells of peripheral blood from patients with HCC patients and healthy donors (HDs). (e–g) The numbers of NK cells (e), CD16⁺ NK cells (f) or CD16⁻ NK cells (g) in the peripheral blood of HCC patients and HDs. (h,i) The proportion of interferon (IFN)- γ^+ NK cells (h) or tumor necrosis factor (TNF)- α^+ NK cells (i) in HCC patients and HDs. Data are shown as means ± standard deviation (s.d.). Analyses yielding *P*-values of less than 0.05 were considered statistically significant. **P* < 0.05; ***P* < 0.001; ****P* < 0.0001.

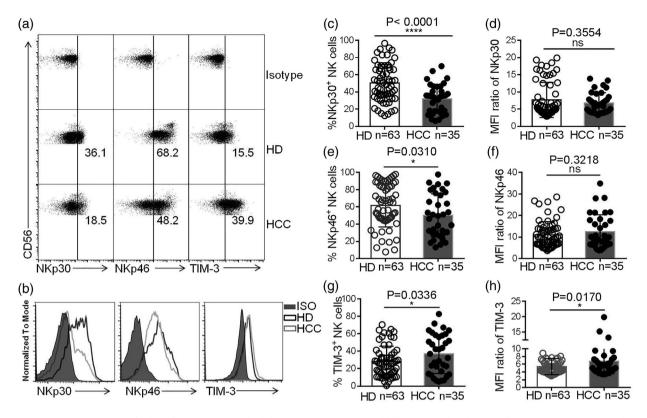


Fig. 2. Abnormal expression of cell-surface activating and inhibitory receptors on natural killer (NK) cells in hepatocellular carcinoma (HCC). (a,b) Dot-plot (a) and histogram (b) representations of T cell immunoglobulin mucin-3 (TIM-3), NKp30 and NKp46 expression on NK cells in HCC patients and healthy donors (HDs). (c,e,g) The proportion of NKp30⁺ NK cells (c), NKp46⁺ NK cells (e) or TIM-3⁺ NK cells (g) in HCC patients and HDs. (d,f,h) The median fluorescence intensity (MFI) ratios of NKp30 (d), NKp46 (f) or TIM-3 (h) on NK cells. MFI ratio of X = X-MFI/isotype-MFI. Data are shown as means \pm standard deviation (s.d.). Analyses yielding *P*-values of less than 0-05 were considered statistically significant. **P* < 0-05; ***P* < 0-001; ****P* < 0-001.

blood of HCC patients were lower compared to HDs (Fig. 1b). Moreover, the proportion of the CD56^{dim}CD16⁺ NK subset within the total NK cell population decreased, while the proportion of the CD56^{bright}CD16⁻ NK subset increased in HCC patients compared to HD controls, implying a shift in the NK cell composition of HCC patients (Fig. 1c,d). Accordingly, the absolute number of NK cells in the peripheral blood of HCC patients was markedly lower than in that of HDs (Fig. 1e). The same was true for the absolute numbers of both CD56^{dim}CD16⁺ and CD56^{dim}CD16⁻ NK cells of HCC patients displayed impaired IFN-γ production (Fig. 1h,i). Taken together, our data reveal a reduction in the frequency, number and function of NK cells in HCC patients.

Abnormal expression of activating and inhibitory receptors on NK cells in HCC patients

NK cell anti-tumor activity is regulated by activating and inhibitory receptors, which led us to investigate the expression characteristics of these receptors on NK cells in HCC patients compared to HDs. We found that the frequencies of NKp30⁺ and NKp46⁺ NK cells were lower in the peripheral blood of HCC patients compared to HDs (Fig. 2a–f). In contrast, the frequency of TIM-3⁺ NK cells was higher in HCC patients (Fig. 2g,h). In addition, the frequency of programmed cell death 1 (PD-1⁺) NK cells was slightly higher in HCC patients, although the frequency was very low in both HCC patients and HDs (Supporting information, Fig. S1a–c). The frequency of TIGIT⁺ NK cells in HCC patients was similar to that in HDs (Supporting information, Fig. S1d–f). No significant difference of frequencies of NKp44⁺ and NKG2D⁺ NK cells was observed in HCC patients and HDs (Supporting information, Fig. S1g–l). Taken together, our data reveal that the expression of activating and inhibitory receptors on NK cells was abnormal in HCC patients, indicating NK cell dysfunction.

Reduced Siglec-7 expression on NK cells in the peripheral blood of HCC patients

Siglec-7 and Siglec-9 are MHC-I-independent NK cell inhibitory receptors. We found that the frequency of Siglec-9⁺ NK cells in HCC patients was similar to that

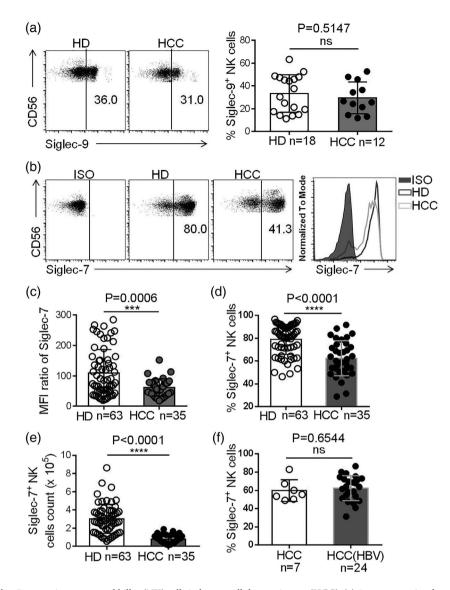


Fig. 3. Decreased Siglec-7 expression on natural killer (NK) cells in hepatocellular carcinoma (HCC). (a) A representative dot-plot showing Siglec-9⁺ expression on the NK cells of HCC patients and healthy donors (HDs), assessed by flow cytometry (left panel) and comparison between Sigleg-9 expression on the NK cells of HCC patients and HDs (right panel). (b) Representative dot-plot (left panel) and histogram (left panel) of surface Siglec-7 expression on NK cells in HCC patients and HDs. (c) The Siglec-7 median fluorescence intensity (MFI) ratio on the NK cells of HCC patients and HDs. (d) The proportion of Siglec-7⁺ NK cells in HCC patients and HDs. (e) Siglec-7⁺ NK cell counts in HCC patients and HDs. (f) The comparison of Siglec-7⁺ NK cells in HCC patients with or without hepatitis B virus (HBV) infection. Data are shown as means ± standard deviation (s.d.). Analyses yielding *P*-values less than 0-05 were considered as statistically significant. **P* < 0-05; ***P* < 0-01; ****P* < 0-0001.

in HDs (Fig. 3a) However, the median fluorescence intensity (MFI) level of Siglec-7 expression on HCC patient NK cells was significantly lower compared with HDs (Fig. 3b,c). The frequency and number of Siglec-7⁺ NK cells were also lower in the peripheral blood of HCC patients (Fig. 3d,e). Moreover, the number of Siglec-7⁺ NK cells in HCC patients with alpha-fetoprotein (AFP) (> 25) was significantly lower than that in HCC patients with AFP (\leq 25) (Table 1). It had been reported that the frequency of Siglec-7⁺ NK cells is reduced in hepatitis B virus (HBV)-infected individuals [15]. However, we found no difference in the frequency of Siglec-7⁺ NK cells between HBV⁻ and HBV⁺ HCC patients (Fig. 3f), suggesting that HBV infection was not the reason for Siglec-7 downregulation in HCC patient NK cells. The expression of activating and inhibitory receptors on NK cells was further investigated on different NK cell subsets. The frequencies of CD56^{dim}CD16⁺NKp30⁺, CD56^{dim}CD16⁺NKp46⁺ and CD56^{dim}CD16⁺Siglec-7⁺ NK cells were all reduced in HCC patients compared to HDs (Fig. 4a–e). Taken together,

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Table 1. Clinical characteristics of HCC patients

	Mean with s.e.m. (Siglec-7 ⁺ NK cell		
Characteristics	Number	count)	P-value (unpaired Student's t-test)
Age (years)			
≤ 56	19	$81\ 864 \pm 11\ 701$	0.5996
> 56	16	$73\ 520\pm 10\ 039$	
Gender			
Male	31	$79\ 218\pm 8505$	0.6816
Female	4	$68\ 998 \pm 1819$	
AFP (ng/ml)			
≤ 25	15	99 116 ± 1241	0.0163
> 25	20	$62\ 249\pm 8526$	
ALT, U/l			
≤ 40	18	93 122 ± 10 065	0.0438
> 40	17	$62\ 091\pm 10\ 880$	
Tumor size, cm			
≤ 5	16	77 624 ± 10 953	0.9608
> 5	19	$78\ 408\pm 11\ 189$	
Tumor multiplicity			
Solitary	22	76 363 ± 9099	0.7820
Multiple	13	$80\ 904 \pm 14\ 587$	
Vascular invasion			
Absent	19	81 197 ± 11 309	0.6651
Present	16	$74\ 312\pm 10\ 697$	
Intrahepatic metastasis			
No	23	77 662 ± 8323	0.9462
Yes	12	78 791 ± 16 641	

*Analyses yielding P-values of less than 0.05 were considered as statistically significant.

AFP = a-fetoprotein; ALT = alanine aminotransferase; HCC = hepatocellular carcinoma; NK = natural killer; s.e.m. = standard error of the mean.

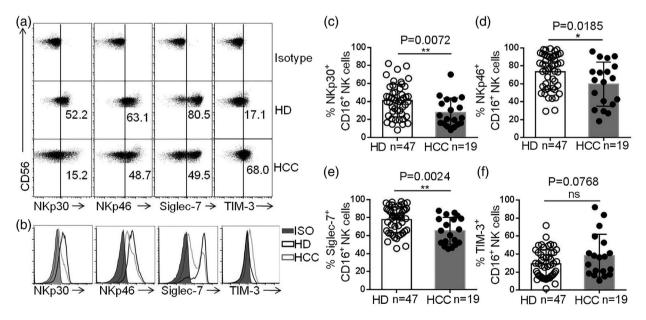


Fig. 4. Decreased Siglec-7 expression on the CD16⁺ natural killer (NK) cells subset in hepatocellular carcinoma (HCC). (a,b) Dot-plot (a) and histogram (b) representation of NKp30, NKp46, Siglec-7 and T cell immunoglobulin mucin-3 (TIM-3) expression on CD16⁺ NK cells. (c-f) The proportions of NKp30⁺CD16⁺ NK cells (c), NKp46⁺CD16⁺ NK cells (d), Siglec-7⁺CD16⁺ NK cells (e) and TIM-3⁺CD16⁺ NK cells (f) within the total NK cell population in the peripheral blood of HCC patients and healthy donors (HDs). Data are shown as means ± standard deviation (s.d.). Analyses yielding *P*-values less than 0.05 were considered statistically significant. **P* < 0.01; ****P* < 0.001; *****P* < 0.0001.

our data demonstrate that the expression of Siglec-7 was lower on NK cells in the peripheral blood of HCC patients.

Siglec-7⁺ NK cells exhibit the hyperresponsiveness in HCC patients

Lower numbers of Siglec-7⁺ NK cells were observed in the peripheral blood of HCC patients compared to HDs. In order to more clearly characterize Siglec-7⁺ NK cells in HCC patients, we evaluated their activating and inhibitory receptor expression profiles and measured their effector functions. Our data suggest that Siglec-7⁺ NK cells express higher levels of the activating receptors NKp30 and NKp46 compared with Siglec-7⁻ NK cells, both in HCC patients and HDs (Fig. 5a-f). Moreover, Siglec-7⁺ NK cells produced more TNF- α compared with Siglec-7⁻ NK cells after stimulation with by PMA and ionomycin (Fig. 6a,b). No significant difference was observed in the production of IFN- γ and the expression of CD107a by the Siglec-7⁺ NK cells and Siglec-7⁻ NK cells (Fig. 6c–f). Thus, our data demonstrate that Siglec-7⁺ NK cells express higher levels of activating receptors and exhibit hyperresponsiveness in HCC patients.

Discussion

Siglec-7 is an MHC-I-independent inhibitory receptor on human NK cells that recognizes sialic acid-containing carbohydrates [18]. It has been reported that the expression of Siglec-7 on NK cells is reduced in HBV- or hepatitis C virus (HCV)-infected patients [15]. In the present study,

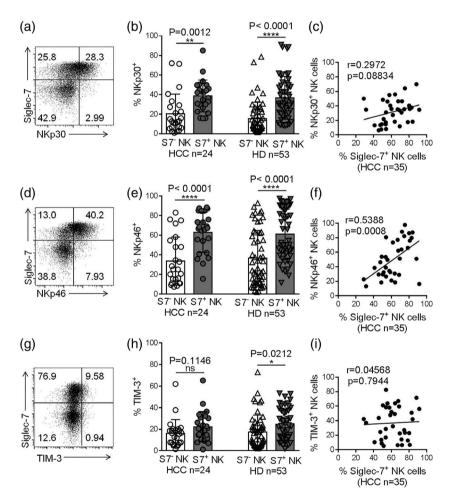


Fig. 5. Siglec-7⁺ natural killer (NK) cells express higher levels of activating receptors. (a) A representative dot-plot showing NKp30 expression on Siglec-7⁺ and Siglec-7⁻ NK cells, assessed by flow cytometry. (b) The proportions of NKp30⁺Siglec-7⁺ NK cells and NKp30⁺Siglec-7⁻ NK cells in hepatocellular carcinoma (HCC) patients and healthy donors (HDs). (c) The correlation between Siglec-7 and NKp30 expression on NK cells. (d) Dot-plot representation of NKp46 expression on Siglec-7⁺ and Siglec-7⁻ NK cells in HCC patients and HDs. (f) The correlation between Siglec-7 and NKp46 expression on NK cells in term siglec-7⁻ NK cells. (g) A representative dot-plot showing T cell immunoglobulin mucin-3 (TIM-3) expression on Siglec-7⁺ and Siglec-7⁻ NK cells. (h) The proportions of TIM-3⁺Siglec-7⁺ and TIM-3⁺Siglec-7⁻ NK cells in HCC patients and HDs. (i) The correlation between Siglec-7 and TIM-3 expression on NK cells. (b) The proportions of NK cells. Data are shown as means ± standard deviation (s.d.). Analyses yielding *P*-values less than 0-05 were considered statistically significant. **P* < 0-05; ***P* < 0-01; ****P* < 0-001;

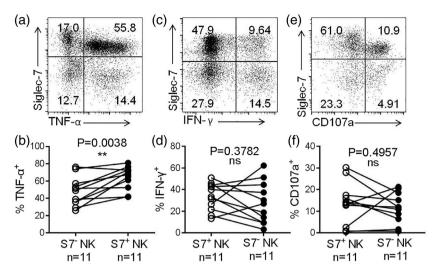


Fig. 6. Siglec-7⁺ natural killer (NK) cells exhibit stronger effector functions than Siglec-7⁻ NK cells. (a) A representative dot-plot showing TNF- α production by Siglec-7⁺ and Siglec-7⁻ NK cells, assessed by flow cytometry. (b) Comparison of TNF- α production by Siglec-7⁺ NK cells and Siglec-7⁻ NK cells. (c) A representative dot-plot showing the production of IFN- γ by Siglec-7⁺ and Siglec-7⁻ NK cells. (d) Comparison of IFN- γ production by Siglec-7⁺ and Siglec-7⁻ NK cells. (e) A representative dot-plot showing CD107a expression by Siglec-7⁺ and Siglec-7⁻ NK cells. (f) Comparison of CD107a expression by Siglec-7⁺ and Siglec-7⁻ NK cells. (d) Analyses yielding *P*-values less than 0.05 were considered statistically significant. **P* < 0.05; ***P* < 0.01; ****P* < 0.001; ****P* < 0.001.

we found that Siglec-7 expression was also reduced on the NK cells of HCC patients. Siglec-7⁺ NK cells displayed higher levels of activating receptor expression compared with Siglec-7⁻ NK cells. Therefore, reduced Siglec-7 expression on NK cells appears to predict NK cell dysfunction in the context of primary HCC.

Like cytotoxic CD8+ T cells, NK cells play important roles in host immunity against cancer. NK immunotherapy may therefore provide a feasible, if not an improved, alternative to the more conventional type of cancer immunotherapy involving T cells [19-21]. A specific NK cell-mediated immune response has been confirmed in diverse cancer types, including hematological malignancies, non-small-cell lung cancer, colorectal cancer and HCC [22-25]. It has been widely reported that the peripheral and tumor-infiltrating NK cells of HCC patients are associated with an abnormal frequency, phenotype and function. Our study has further confirmed these findings. We found that the frequency and absolute number of NK cells in the peripheral blood of HCC patients was lower than in that of HDs. Moreover, the proportion of the CD56^{dim}CD16⁺ NK cell subset was reduced, while the proportion of the CD56^{dim}CD16⁻ NK cell subset was elevated in HCC patients. In addition, the IFN-y production capacity of NK cells of HCC patients was also impaired compared to HD controls. The activity and homeostasis of NK cells is regulated by activating, inhibitory and cytokine receptors, which are also utilized as a means of escape from the NK cell-mediated immune response [26]. In HCC, NK dysfunction can be caused by a number of cell-specific receptors and their molecular interactions, including: (i) myeloid-derived suppressor cells (MDSC), via NKp30 [27]; (ii) monocytes/macrophages, via the CD48–2B4 interaction [28]; (iii) HCC-associated fibroblasts, via prostaglandin E_2 (PGE₂) and indoleamine 2,3 dioxygenase (IDO) [29]; (iv) the soluble form of MHC-I-related chain A (sMICA); and (v) the MHC-I-dependent inhibitory receptor NKG2A [30]. Therefore, one of the central areas of current NK cell-based immunotherapy research is to decipher the potential immune checkpoints modulating NK cell activity in HCC.

Siglecs are a family of sialic acid-binding immunoglobulin-like co-receptors that recognize sialic acid-containing ligands and regulate immune cell function [31-33]. The interaction of Siglecs with ligands helps cancer cells to escape the anti-tumor immune attack [34]. Recently, Siglec-15 became recognized as an immune suppressor and potential target for normalization cancer immunotherapy [35]. Furthermore, tumor-expressed CD24 mediates immune evasion through its interaction with Siglec-10 on macrophages [36]. Siglec-10 is also reported to be associated with lower rates of survival among HCC patients, and Siglec-10⁺ NK cells exhibit impaired effector functions [37]. Siglec-9 has been found to regulate effector memory CD8⁺ T cells in the context of melanoma [38]. Human NK cells ubiquitously express Siglec-7, whereas a smaller subset expresses Siglec-9 [39]. Both Siglec-7 and -9 contain ITIMs to recruit Src homology-2 (SH2) phosphatases and suppress kinase phosphorylation [31]. Siglec-9 defines a subset of mature cytotoxic NK cells with enhanced chemotactic potential, and CD56^{dim}Siglec-9⁺ NK cell numbers have been shown to be reduced in the peripheral blood of patients with colon adenocarcinoma or malignant melanoma [39]. The study authors have reported that Siglec-7 is expressed at similar levels on the peripheral blood NK cells of these patients and HDs [39]. In our study, however, there was no significant difference in Siglec-9 expression on the NK cells of HCC patients and HDs. Instead, both the frequency Siglec-7⁺ peripheral NK cells and the levels of NK cell surface Siglec-7 expression were significantly reduced in HCC patients compared to HDs. We also observed that Siglec-7⁺ NK cells express higher levels of activating receptors and exhibit stronger effector functions. Moreover, the reduction in Siglec-7 expression on peripheral blood NK cells in HCC patients was not associated with HBV infection, despite previous reports of Siglec-7 being significantly reduced on the NK cells of HBV-infected patients [15]. Therefore, reduced Siglec-7 levels on NK cells appear to predict NK cell dysfunction in the context of HCC.

Siglec-7 has a preference for $\alpha 2$, 6-linked sialic acid and $\alpha 2$, 8-linked sialic acid [18]. Cao et al. reported that α2,6-linked sialic acid was positive on HCC tissues, while normal hepatocytes and preneoplastic foci of altered hepatocytes did not express $\alpha 2,6$ -linked sialic acid [40]. Gong *et al.* also reported that α 2,6-sialic acids increased significantly on the hepatocyte membrane after the carcinomatous change [41]. These findings demonstrate that Siglec-7 ligands exist in HCC tissues. Accumulating studies report that Siglec-7 ligands inhibit NK cell activation and cytotoxicity, which have been used by glycocalyx engineering to protect against NK cell killing [13]. Siglec-7 ligands expressed on K562 cells significantly suppressed cytolysis of K562 cells by peripheral NK cells [39]. Disialosyl globopentaosylceramide (DSGb5) expressed on renal cell carcinoma (RCC) cells inhibited NK cell cytotoxicity in a DSGb5-Siglec-7-dependent manner [42]. Therefore, it is suggested that Siglec-7 ligands in HCC tissues might inhibit NK cell function in the hepatic location. However, the precise mechanisms responsible for: (i) the changes in Siglec-7 expression on NK cells in HCC patients; and (ii) the role of Siglec-7 in modulating NK cell-mediated anti-tumor activity and ultimately HCC patient survival, still require further investigation.

In summary, our study has revealed for the first time, to our knowledge, that the extent and expression levels of Siglec-7 on peripheral blood NK cells were significantly reduced in HCC patients, and that in HCC, Siglec-7⁺ NK cells exhibit powerful anti-tumor activity. These findings serve to deepen our understanding of Siglec-7⁺ NK cell properties in the context of HCC and provide new avenues for NK-based cancer immunotherapy.

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Author contributions

J. L. and X. W. conceived the study, designed the experiments and wrote the paper. L. T. and S. W. performed the experiments, analyzed the data and wrote the paper. L. Y. and L. J. provided technical assistance in certain experiments.

Disclosures

The authors declare no competing interests.

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Supporting Information

Additional supporting information may be found in the online version of this article at the publisher's web site:

Fig. S1. The alteration of activating receptors and inhibitory receptors on peripheral NK cells in HCC patients.