ORIGINAL ARTICLE

B7-H3 is expressed in human hepatocellular carcinoma and is associated with tumor aggressiveness and postoperative recurrence

Tai-Wei Sun · Qiang Gao · Shuang-Jian Qiu · Jian Zhou · Xiao-Ying Wang · Yong Yi · Jie-Yi Shi · Yong-Feng Xu · Ying-Hong Shi · Kang Song · Yong-Sheng Xiao · Jia Fan

Received: 17 September 2011/Accepted: 27 April 2012/Published online: 22 May 2012 © Springer-Verlag 2012

Abstract B7-H3, a novel B7 family member, positively or negatively regulates T-cell responses. We investigated the clinical relevance and prognostic significance of B7-H3 in hepatocellular carcinoma (HCC). Western blotting showed B7-H3 upregulation in 17 of 24 (70.8 %) HCC tissues compared with nontumor liver tissues (p = 0.028). B7-H3 immunostaining on tissue microarrays containing 240 HCC patient samples indicated that 225 (93.8 %) tumors had aberrant B7-H3 expression, with strong intensity in 79 (32.9 %) cases, whereas B7-H3 expression in

Tai-Wei Sun, Qiang Gao, and Shuang-Jian Qiu contributed equally to this work.

Electronic supplementary material The online version of this article (doi:10.1007/s00262-012-1278-5) contains supplementary material, which is available to authorized users.

T.-W. Sun · Q. Gao · S.-J. Qiu · J. Zhou · X.-Y. Wang · Y. Yi · J.-Y. Shi · Y.-H. Shi · K. Song · Y.-S. Xiao · J. Fan (\boxtimes)

Liver Cancer Institute, Zhongshan Hospital and Shanghai Medical School, Fudan University, 180 Feng Lin Road, Shanghai 200032, People's Republic of China e-mail: jiafan99@yahoo.com

 $\begin{array}{l} T.-W. \; Sun \cdot Q. \; Gao \cdot S.-J. \; Qiu \cdot J. \; Zhou \cdot X.-Y. \; Wang \cdot \\ Y. \; Yi \cdot J.-Y. \; Shi \cdot Y.-H. \; Shi \cdot K. \; Song \cdot Y.-S. \; Xiao \cdot J. \; Fan \\ Key \; Laboratory \; of \; Carcinogenesis \; and \; Cancer \; Invasion, \\ Ministry \; of \; Education, \; Shanghai, \; People's \; Republic \; of \; China \\ \end{array}$

J. Zhou · J. Fan Institute of Biomedical Sciences, Fudan University, Shanghai, People's Republic of China

Y.-F. Xu

Department of Pancreas and Hepatobiliary Surgery, Cancer Hospital and Shanghai Medical School, Fudan University, Shanghai, People's Republic of China peritumor liver cells was weak in most cases (226; 94.2 %). Notably, patients with high/moderate tumor cell B7-H3 expression showed significantly poorer survival (p = 0.009) and increased recurrence (p = 0.002). After multivariable adjustment, high/moderate B7-H3 expression remained significant for an increased risk of recurrence (hazard ratio = 1.79; 95 % confidence interval = 1.19-2.70; p = 0.005). B7-H3 expression correlated with invasive phenotypes like vascular invasion and advanced tumor stage, and the metastatic potential of HCC cell lines. Flow cytometry showed that B7-H3 expression is inversely correlated with proliferation and interferon- γ production by infiltrating T cells. Interferon- γ stimulation significantly upregulated B7-H3 expression in HCC cells in vitro, implicating B7-H3 expression as a feedback mechanism to evade anti-tumor immunity. Importantly, the prognostic value of B7-H3 expression was validated in an independent cohort of 206 HCC patients. Collectively, our data suggest that B7-H3 was abundantly expressed in HCC and was associated with adverse clinicopathologic features and poor outcome. Thus, B7-H3 represents an attractive target for diagnostic and therapeutic manipulation in human HCC.

Keywords Hepatocellular carcinoma · B7-H3 · Prognosis · Immunohistochemistry · Tumor immunity

Abbreviations

- HCC Hepatocellular carcinoma
- TNM Tumor-node-metastasis
- SD Standard deviation
- OS Overall survival
- TIL Tumor-infiltrating lymphocytes
- HR Hazard ratio
- CI Confidence interval

Introduction

Hepatocellular carcinoma (HCC) is the fifth most common cancer and the third most common cause of cancer death worldwide [1]. Most patients present with disease too advanced for curative treatment by resection, transplantation, or local ablation. Although a survival advantage for the multitargeted kinase inhibitor sorafenib in patients with advanced HCC has been suggested, this effect is modest [2]. No other systemic therapy has shown a survival benefit, and recurrence is common, even for curative treatment, highlighting the need for identification of prognostic biomarkers and new therapeutic strategies.

Accumulating evidence suggests that immunotherapy may become a potent therapeutic option for patients with HCC [3]. Stimulating T-cell activation or eliminating inhibitory T-cell signaling to enforce antitumor responses is a prerequisite for immune-based therapies to eradicate human cancers. It is evident that T-cell priming, activation, and tolerance are ultimately governed by costimulatory and coinhibitory signals, in which the B7/CD28 family of coregulatory molecules plays an indispensable role [4]. The prototype ligands of this family, B7-1 and B7-2, have welldefined roles as essential costimulatory molecules involved in T-cell activation. Within the past decade, new members of the B7 family (B7-H1, PD-L2, B7-H3, and B7-H4) have been identified; however, their exact roles in tumor immunity remain unclear. We and others have reported that several of these B7-H molecules (e.g., B7-H1 and B7-H4) are expressed in human cancers and are implicated as coregulatory inhibitors that may induce T-cell anergy or apoptosis upon antigen recognition [5-7].

B7-H3 was discovered in 2001 and was initially identified as a costimulatory molecule on T cells [8]. However, the precise function of B7-H3 remains debatable [9]. Both stimulatory properties, such as promotion of T-cell proliferation and interferon (IFN)- γ production [8, 10], along with inhibitory properties, including impairment of Th1 responses and protection from natural killer cell-mediated lysis [11], have been described. Studies regarding the clinical significance of B7-H3 in human cancers have suggested that higher B7-H3 expression indicates increased risk of recurrence in some cancers [12-14], or paradoxically, correlates with better postoperative survival in others [15, 16], or has no clinical significance [17]. Although expression of B7-H3 in human cancers appears to be a fairly general phenomenon, the literature is currently devoid of clinical observations of B7-H3 in human HCC.

Thus, we investigated the prognostic significance and clinical relevance of B7-H3 expression in a cohort of 240 HCC patients treated with curative resection. We found that tumor cell instead of tumor vasculature expression of B7-H3 was significantly associated with an increased risk of recurrence and aggressive clinicopathologic features. The results were further validated in an independent cohort of 206 HCC patients.

Materials and methods

Cell lines

Eight human HCC cell lines were used, including HepG2, Hep3B, PLC/PRE/5, SMCC-7721, MHCC97L, MHCC97H, HCCLM3, and HCCLM6. The latter four HCC cell lines were established at our institute, all from the same parental cell line MHCC97, with stepwise pulmonary metastatic potential (MHCC97L < MHCC97H < HCCLM3 < HCCLM6) as reported elsewhere [18, 19]. Cell lines were all maintained in Dulbecco's modified Eagle's Medium supplemented with 10 % fetal bovine serum and 1 % penicillin-streptomycin at 37 °C with 5 % CO₂.

Patient cohort and clinical follow-up

Archived tissues were obtained for a cohort of 240 patients who received curative resection of HCC at Liver Cancer Institute, Zhongshan Hospital of Fudan University between 2002 and 2006 [7, 20]. Patients neither showed signs of distant metastasis nor received anticancer therapy before surgery. Follow-up procedures and postoperative treatments according to uniform guidelines were described previously [21]. Conventional clinicopathologic variables are detailed in Supplementary Table S1. The tumor stage was determined according to the 2002 AJCC/UICC tumor-node-metastasis (TNM) classification system [22]. Tumor differentiation was graded using the Edmondson grading system. Liver function was assigned by the Child-Pugh scoring system. Data were censored at the last follow-up for patients without recurrence or death. Time to recurrence or overall survival (OS) was defined as the interval between the time of surgery to that of recurrence or death, respectively.

The median follow-up period was 39.0 months (range, 1.5–95.0 months; standard deviation [SD], 22.7 months). At the last follow-up (March 31st, 2010), 127 (52.9 %) patients were confirmed to have relapsed, including 89 with intrahepatic recurrence, 17 with extrahepatic metastasis, and 21 with both events. A total of 122 (50.8 %) patients had died, including 52 due to liver failure or bleeding from the gastrointestinal tract and the remaining 70 cases due to tumor recurrence.

Fresh tissues from an additional series of 24 HCC patients who underwent resection in 2010 were used for isolation of tumor-infiltrating lymphocytes (TILs) and for flow cytometry and Western blot analyses. Among these 24 patients, 12, 7, and 5 were classified as TNM stage I, II, and IIIA, respectively. Informed consent was obtained, and the study was approved by the Zhongshan Hospital Research Ethics Committee.

Tissue microarray and immunohistochemistry

Tissue microarrays were performed as described previously [7, 20]. Briefly, representative areas, away from necrotic and hemorrhagic materials, were premarked in the paraffinembedded blocks by hematoxylin and eosin staining. Triplicates of 1-mm-diameter cylinders from the center of the tumor and noncancerous liver tissue, along with samples from different controls (spleen, lymph node, artery, and glioma), were included in each case to ensure reproducibility and homogeneity. Serial sections (4-µm thick) were placed on slides coated with 3-aminopropyltriethoxysilane.

Immunohistochemistry was performed using a two-step protocol (Novocastra) as previously described [7, 20]. Briefly, sections were dewaxed, hydrated, and washed. After neutralization of endogenous peroxidase (0.3 % H₂O₂ for 20 min) and antigen retrieval (121 °C for 4 min), slides were preincubated with blocking serum and then incubated at 4 °C overnight with each primary antibody. The primary antibodies were as follows: rabbit polyclonal anti-human 4Ig-B7-H3 (Sigma), mouse monoclonal antihuman CD4 (Novocastra), CD8 (Novocastra), and FOXP3 (Abcam). Subsequently, the sections were serially rinsed, incubated with secondary antibodies, and treated with horseradish peroxidase-conjugated streptavidin. Reaction products were visualized using 3,3'-diaminobenzidine tetrahydrochloride and counterstained with hematoxylin. All tissue microarray stainings, including negative-staining controls using phosphate-buffered saline in place of the primary antibodies, were performed in a single experiment.

Evaluation of immunohistochemical staining

Microarrays were evaluated using light microscopy by two investigators blinded to the clinicopathologic data of the patients. Sections were considered to be positive when tumor cells showed cytoplasmic or membranous B7-H3 immunostaining. The B7-H3 immunostaining intensities were scored as follows: 0, negative; 1, weak; 2, moderate; 3, strong [14]. A negative grade represented no tumor cells showing positive immunostaining. For analysis, the B7-H3 immunostaining intensities were classified as follows: the sections scored as 0 and 1 were defined as the low expression group, and sections scored as 2 and 3 were defined as the high expression group. In addition, B7-H3 expression in the tumor vasculature and the vasculature of adjacent nontumor liver tissue was evaluated and recorded as being absent, focal, moderate, or diffuse in nature [12].

Regarding the TIL densities, the entire 1-mm-diameter core was counted manually under $\times 200$ magnification. The results were expressed as the mean \pm standard error (SE) of the number of cells/1-mm core for each triplicate.

Western blot analysis

Twenty-four cases with paired tumor and peritumoral liver tissues (five cases with venous tumor thrombi), as well as eight human HCC cell lines were analyzed, including six normal liver tissues from patients with benign hepatic tumors as controls. Western blot analysis was performed as described previously [7, 20] with rabbit polyclonal antihuman 4Ig-B7-H3 (Sigma) and β -actin (Cell Signaling Technology) as an internal loading control.

TIL isolation and flow cytometry

TILs were isolated as described previously [23]. The following antibodies were used: fluorescein isothiocyanateconjugated anti-CD4, allophycocyanin-conjugated anti-CD8, and R-phycoerythrin-conjugated anti-Ki-67, anti-IFN- γ , and anti-interleukin (IL)-4 (BD Pharmingen). Cells were stimulated for 5 h with Leukocyte Activation Cocktail (BD Pharmingen). Cells were stained with surface markers, fixed and permeabilized with IntraPre Reagent (Beckman Coulter), and finally stained. A portion of each specimen was stained with isotype-matched fluorescent control monoclonal antibodies. Data were acquired on a FACS-Vantage SE instrument and analyzed with CellQuest software (BD Biosciences).

Induction of B7-H3 in cultured HCC cells and flow cytometry analysis

To determine the effects of IFN- γ , IL-4, and transforming growth factor (TGF)- β_1 on expression of B7-H3 in cultured HCC cells, HCCLM3 and HepG2 cells were seeded in 6-well plates (100,000 cells per well) and incubated with 40 ng/ml of recombinant IFN- γ , IL-4, and TGF- β_1 (all from PeproTech). After 24 and 48 h of stimulation, cells were harvested and stained with mouse monoclonal antihuman B7-H3-PE (Biolegend) or control mouse monoclonal anti-human IgG-PE (Biolegend). Single-color flow cytometry was analyzed with FlowJo Software (TreeStar).

Independent validation

For further validation, we evaluated B7-H3 expression in another set of tissue microarrays containing a consecutive cohort of 206 HCC patients who underwent liver transplantation between 2001 and 2006 at Liver Cancer Institute, Zhongshan Hospital of Fudan University. Immunohistochemistry, quantification of markers, and statistics were conducted in the same manner. Clinicopathologic characteristics are described in Supplementary Table S1. The median duration of follow-up for this cohort was 48.0 months (range, 3.0–113.5 months; SD, 27.8 months). Of the 206 patients examined, 96 (46.6 %) died before the end of the observation period, and 93 (45.1 %) were diagnosed with tumor recurrence.

Statistics

All statistical analyses were performed using SPSS 15.0 for Windows (SPSS, Inc.). Pearson's χ^2 test and Fisher's exact test were applied to compare qualitative variables, and quantitative variables were analyzed using Student's *t* test or Spearman's ρ rank correlation coefficient determination. Univariate analysis was calculated by the Kaplan–Meier method (the log-rank test). Multivariate analysis was done using the Cox multivariate proportional hazards regression model in a stepwise manner (backward, conditional). The model included all clinicopathologic variables found to have significant prognostic value in univariate analysis. A two-tailed p < 0.05 was deemed to indicate statistical significance.

Results

B7-H3 is elevated in HCC tissue

Representative immunostaining images of B7-H3 are presented in Fig. 1. We found that 225 (93.8 %) HCC cases had positive tumor cell B7-H3 expression, with various intensities. A total of 57 (23.8 %) cases showed weak tumor cell B7-H3 intensity, 89 (37.1 %) showed moderate intensity, and 79 (32.9 %) showed marked intensity. In addition, tumor vasculature expression of B7-H3 was apparent in 86.2 % of the tumors. There were 43 (17.9 %), 80 (33.3 %), and 84 (35.0 %) specimens with focal, moderate, and diffuse tumor vasculature B7-H3 expression, respectively. By contrast, peritumor liver cell B7-H3 expression was weak in most cases (226; 94.2 %), with 14 (5.8 %) cases showing moderate intensity and no cases showing marked B7-H3 intensity. However, peritumor liver sinusoid endothelial cells and resident Kupffer cells showed constitutively strong B7-H3 expression. These results were different from the findings in renal cell carcinoma where 360 out of 378 nontumor renal tissues (95.2 %) showed no vasculature B7-H3 expression [12].

To extend our observations, we tested 24 fresh HCC tissues and 8 human HCC cell lines for B7-H3 expression

using Western blot analysis (Fig. 2a). Consistent with our observations in paraffin-embedded HCC tissues, all these HCC tissues and cell lines exhibited expression of B7-H3. In cell lines with a similar genetic background (MHCC97L, MHCC97H, HCCLM3, and HCCLM6), the least aggressive, MHCC97L, displayed the lowest level of B7-H3, whereas the most aggressive, HCCLM3 and HCCLM6, showed the highest levels of B7-H3. In 24 paired samples, B7-H3 was obviously upregulated in 17 of 24 (70.8 %) HCC tissues (Fig. 2b) compared with peritumoral tissues (p = 0.028; Fig. 2c), although peritumor liver sinusoid endothelial cells and Kupffer cells constitutively expressed B7-H3. Specifically, in five cases with macroscopic tumor thrombi, the levels of B7-H3 showed a successive increase from peritumor to tumor tissue, and to the corresponding tumor thrombi (Fig. 2d).

B7-H3 expression inversely correlated with T-cell proliferation, but showed no association with infiltrating T-cell number

We examined the relationship between tumor B7-H3 expression levels and the functional status of tumor-infiltrating T cells. According to Western blot results, the 24 HCC cases were divided into two groups: B7-H3 high (high B7-H3 expression) and B7-H3 low (low B7-H3 expression) groups. Flow cytometry indicated that the B7-H3 high group was associated with a significantly low proliferation rate and IFN- γ production by CD4+ and CD8+ TILs, whereas IL-4 production of CD4+ TILs did not differ significantly between the two groups (Fig. 3).

We further investigated whether tumor B7-H3 expression correlated with the density of tumor-infiltrating T cells. Tissue microarrays containing 240 HCC cases were immunostained for CD4+, CD8+, and FOXP3+ T cells (Supplementary Fig. S1A). No significant difference was found in the density of CD8+ T cells (p = 0.95) between B7-H3 high and B7-H3 low tumors (Supplementary Fig. S1B). Although B7-H3 high HCC contained relatively high numbers of CD4+ T cells (p = 0.065) and FOXP3+ regulatory T cells (p = 0.082) compared with B7-H3 low HCC, the differences did not reach statistical significance (Supplementary Fig. S1B).

IFN- γ markedly increased B7-H3 expression in HCC cells

Treatment with 40 ng/ml recombinant IFN- γ (Supplementary Fig. S2A) markedly increased the expression of B7-H3 in HCCLM3 (1.97-fold) and HepG2 cells (2.17-fold) after 24 h. Additionally, in HCCLM3 cells, the effect induced by IFN- γ treatment almost disappeared after 48 h (1.12-



Fig. 1 Representative immunostaining for B7-H3 expression in HCC and peritumor liver tissue. Peritumor liver cells were unstained (a_1) or weakly stained (a_2) with anti-4Ig-B7-H3 Ab, with limited cases of moderate (b_2) staining, whereas liver sinusoid endothelial cells and Kupffer cells (*red arrow*) show constitutive positive staining. HCC

cells show weak (**b**₁), moderate (**b**₂), or strong (**b**₃) staining. Focal (**c**₁), moderate (**c**₂), or diffuse (**c**₃) tumor vasculature staining is observed. Specific cases show strong staining in both tumor cells and tumor vessels (**d**₁), in venous tumor thrombi (**d**₂; *green arrow*), or in intrahepatic spreading nodules (**d**₂). Magnification, $400 \times$

fold), whereas in HepG2 cells, IFN- γ treatment resulted in a further increase in B7-H3 expression at 48 h (2.84-fold) compared with that at 24 h. By contrast, IL-4 or TGF- β_1 treatment induced no significant change in B7-H3 expression in HCCLM3 and HepG2 cells after 24 or 48 h (Supplementary Fig. S2B and C).

Tumor B7-H3 expression and prognosis

Comparisons of clinicopathologic features by B7-H3 expression in tumor cells are summarized in Table 1. B7-H3 expression in tumor cells was significantly associated with the presence of aggressive features, including a higher Fig. 2 Western blot analysis of B7-H3 expression in HCC tissues and cell lines. a Western blot detection of 4Ig-B7-H3 protein in 24 paired tumors (T) and peritumoral tissues (pT) (including five cases with tumor thrombi [TT]), eight HCC cells lines, and six normal liver tissues (N). β -Actin served as a loading control. b The bar graph illustrates the ratio of the B7-H3 protein level (T/pT). Seventeen of 24 tumors (70.8 %) display upregulated B7-H3 protein levels, compared with the corresponding peritumoral liver tissues. c The B7-H3 protein level obtained by densitometric scanning is significantly upregulated in tumors compared with peritumoral tissues. d B7-H3 expression shows a sequential increase from peritumor tissues to tumor tissues, and to the corresponding tumor thrombi



 α -fetoprotein level, advanced TNM stage, poor differentiation, the presence of liver cirrhosis, and vascular invasion. Univariate analysis revealed that patients with tumors with high B7-H3 expression were more likely to experience cancer recurrence (p = 0.002) and death (p = 0.009) compared with patients with tumors with low B7-H3 expression (Table 2). The rates of recurrence-free survival and OS (in parentheses) at 1, 3, and 5 years following resection were 79 % (85 %), 61 % (68 %), and 53 % (58 %) for patients with tumors with low B7-H3 expression compared with 60 % (79 %), 40 % (49 %), and 35 % (41 %) for those in the B7-H3 high group, respectively (Fig. 4a, b).

The associations of tumor B7-H3 intensity with patient outcome after adjusting for clinicopathologic features are shown in Table 2. High tumor B7-H3 intensity was significantly associated with an adverse outcome even after multivariate adjustment: patients in the B7-H3 high group were about twice as likely to suffer from tumor recurrence (hazard ratio [HR], 1.79; 95 % confidence interval [CI], 1.19–2.70; p = 0.005) compared with those in the B7-H3 low group.

B7-H3 expression in the tumor vasculature has been shown to be associated with adverse clinical outcome in patients with renal cancer [12] and ovarian carcinoma [24]. However, in the present study, moderate or diffuse B7-H3 expression in the tumor vasculature was not significantly associated with an increased risk of recurrence or death from HCC compared with patients with no B7-H3



Fig. 3 Flow cytometry analysis of proliferation and cytokine production by tumor-infiltrating lymphocytes (TILs). TILs were isolated from 24 fresh HCC tissues. The 24 HCC cases were dichotomized into B7-H3 high and B7-H3 low groups, according to the median level of B7-H3 expression as determined by western blot assays. The percentage of Ki-67+ (**a**) and IFN- γ + (**b**) cells in CD4+

expression or focal B7-H3 expression in the tumor vasculature (Table 2; Fig. 4c, d).

Independent validation

The prognostic ability of the B7-H3 expression profile in tumor cells was validated in an independent dataset with multivariate analysis (Supplementary Table S2). The 1-, 3-, and 5-year recurrence-free survival and OS (in parentheses) rates were 79 % (87 %), 73 % (71 %), and 67 % (66 %) for patients with lower B7-H3 expression, compared with 68 % (75 %), 50 % (51 %), and 47 % (47 %) for patients with higher B7-H3 expression, respectively (Fig. 4e, f).



and CD8+ T cells, as well as the frequency of IL-4+CD4+ T cells (c) are shown. The gates were based on isotype-matched negative controls. The data shown are representative *dot* plots from more than five independent experiments. **d** shows the statistical analysis of these samples

Notably, tumor B7-H3 expression allowed us to subclassify TNM stage II patients, with the median recurrence-free survival and OS (in parentheses) times of 96.0 (84.0) versus 84.0 (54.0) months for patients with low versus high expression levels, respectively (p = 0.018, p = 0.026, respectively; log-rank test).

Discussion

We showed that expression of the T-cell coregulatory ligand B7-H3 was low in normal liver parenchyma cells and increased significantly in HCC cells. High B7-H3

Table 1 Correlation of clinicopathologic characterist with B7-H3 expression

Table 1Correlation ofclinicopathologic characteristicswith B7-H3 expression	Characteristics	Tumor c	cell B7-H3		Tumor vessel B7-H3		
		Low	High	р	Low	High	р
	Age, years						
	<52	32	92	0.143	41	83	0.63
	- >52	40	76		35	81	
	Gender						
	Male	66	138	0.058	57	147	0.003
	Female	6	30		19	17	
	Hepatitis history						
	No	7	9	0.214	6	10	0.604
	Yes	65	159		70	154	
	α-fetoprotein (ng/ml)						
	≤20	44	38	< 0.0001	24	58	0.565
	>20	28	130		52	106	
	γ-GT (U/l)						
	≤54	31	73	0.955	35	69	0.563
	>54	41	95		41	95	
	Liver cirrhosis						
	No	14	14	0.014	10	18	0.624
	Yes	58	154		66	146	
	Tumor size (cm)						
	≤5	38	79	0.414	39	78	0.588
	>5	34	89		37	86	
	Tumor encapsulation						
	Complete	41	78	0.135	32	87	0.115
	None	31	90		44	77	
	Tumor number						
	Single	58	126	0.351	55	129	0.284
	Multiple	14	42		21	35	
	Vascular invasion						
	No	49	82	0.006	46	85	0.208
	Yes	33	86		30	79	
	Tumor differentiation						
	I–II	50	85	0.007	38	97	0.184
	III–IV	22	83		38	67	
Statistical analyses were	TNM stage						
performed using the chi-square	Ι	41	65	0.014	34	72	0.786
test	II	21	55		22	54	
γ -GT γ -glutamyl transferase, TNM tumor-node-metastasis	IIIA	10	48		20	38	

staining was observed in a subset of 70 % of tumor specimens and was significantly associated with multiple adverse prognostic features of HCC, including a higher α -fetoprotein level, advanced TNM stage, poor differentiation, liver cirrhosis, and vascular invasion. More importantly, after multivariate adjustment for conventional prognostic factors, high B7-H3 expression persisted as a significant predictor of tumor recurrence following curative resection. The results were further validated in an independent cohort of HCC patients who underwent liver transplantation. Thus, B7-H3 may prove useful for clinical evaluation of HCC patients who undergo curative surgical treatment for their cancer, particularly to identify high-risk patients who are at increased risk of cancer progression.

Substantive research has been performed regarding the role of B7-H3 in tumor immunity; however, the pathophysiologic function of B7-H3 has yet to be fully elucidated. Both negative and positive immunologic functions for B7-H3 have been reported. Although, in the present study, we have suggested a detrimental role for B7-H3 in

Table 2 Univariate and multivariate analyses of prognostic factors associated with recurrence and survival among 240 HCC patients

Variables	Time to recurrence				Overall survival			
	Univariate p	Multivariate			Univariate	Multivariate		
		HR	95 % CI	р	р	HR	95 % CI	р
Age, years (>52 vs. ≤52)	0.217			NA	0.304			NA
Gender (male vs. female)	0.078			NA	0.254			NA
Hepatitis history (yes vs. no)	0.313			NA	0.627			NA
α-fetoprotein (ng/ml) (>20 vs. ≤20)	0.001			NS	0.0002			NS
γ-GT (U/l) (>54 vs. ≤54)	0.066			NA	0.0004	1.72	1.18-2.49	0.004
Liver cirrhosis (yes vs. no)	0.143			NA	0.015			NS
Tumor differentiation (poor vs. well)	0.073			NA	0.001	1.65	1.16-2.34	0.005
Tumor size (cm) (>5 vs. \leq 5)	< 0.0001			NS	< 0.0001			NS
Tumor multiplicity (multiple vs. single)	< 0.0001	2.10	1.45-3.06	0.0001	0.001	1.80	1.23-2.64	0.003
Tumor encapsulation (none vs. complete)	< 0.0001	1.56	1.09-2.23	0.015	0.009			NS
Vascular invasion (yes vs. no)	< 0.0001	3.43	2.37-4.95	< 0.0001	< 0.0001	3.93	2.70-5.73	< 0.0001
TNM stage (IIIA vs. II vs. I)	< 0.0001			NA	< 0.0001			NA
Tumor cell B7-H3 (high vs. low)	0.002	1.793	1.19-2.70	0.005	0.009			NS
Tumor vessel B7-H3 (high vs. low)	0.376			NA	0.627			NA

Univariate analysis was performed using the Kaplan-Meier method (log-rank test). Multivariate analysis was performed using the Cox multivariate proportional hazards regression model in a stepwise manner (backward, conditional)

y-GT y-glutamyl transferase, TNM tumor-node-metastasis, NS not significant, NA not adopted, HR hazard ratio, CI confidence interval

HCC patients, in murine synergistic HCC models, intratumoral administration of a B7-H3-expressing plasmid and arsenic trioxide or a vasostatin-expressing plasmid synergized to completely eradicate established tumors [25, 26]. In the mouse HCC models, B7-H3 therapy activated CD8+ and natural killer cells and increased their infiltration into tumors [25, 26]. Likewise, opposing roles regarding B7-H3 in mice experiments versus human clinical settings have also been recorded in colon cancer [27, 28]. The contrasting roles of B7-H3 can likely be attributed to multiple receptors on different cells. Indeed, a costimulatory receptor for mouse B7-H3 that preferentially enhances mouse CD8+ T-cell activation (TREM-like transcript 2) has been identified recently [10]. However, Leitner and colleagues found no evidence for such an interaction between TREM-like transcript 2 and B7-H3 in both mice and humans [29], and to date, cognate coinhibiting receptors for B7-H3 have not been elucidated.

Additionally, earlier observations of human gastric [16] and pancreatic carcinoma [15] have implicated that B7-H3positive tumor cells are associated with improved patient survival, whereas subsequent studies in the same cancers yielded opposite results [30, 31]. More recently, B7-H3 has been overwhelmingly implicated as a potent inhibitor of T-cell activity. For example, in neuroblastoma [32], breast [14], lung [33], and prostate [6, 34] cancers, patients with strong B7-H3 expression are more likely to show disease spread. Additionally, the B7-H3-positive vasculature of ovarian carcinoma implies increased recurrence and reduced survival [24]. Furthermore, either B7-H3 expression in tumor cells or diffuse B7-H3 expression in the tumor vasculature significantly predicts an increased risk of death from renal cell carcinoma [12]. Evidence from mechanistic studies has revealed that both mouse and human B7-H3 inhibit T-cell activation and production of effector cytokines such as IFN- γ through regulation of NFTA, NF- κ B, and AP-1, three major signaling pathways through which the T-cell receptor regulates gene transcription [8, 10]. Based on the strong correlation between B7-H3 intensity and adverse clinicopathologic features of human HCC, we surmise that B7-H3 functions as an inhibitor of T cell-mediated immunity that promotes aggressive clinical behavior of HCC. Supporting our hypothesis, microRNA miR-29, a newly identified regulator of B7-H3 protein expression that decreases B7-H3 translation [35], is markedly downregulated in HCC and plays an important role in apoptosis and molecular etiology of HCC [36]. Besides HCC, downregulation of miR-29 was reported in neuroblastoma [35], ovarian carcinoma [37], and other malignancies [38, 39], which may lead to feedback upregulation of B7-H3 and consequent immune evasion in these tumors.

To further provide insight into the clinical significance of B7-H3 in HCC, we performed Western blotting on 24 pairs of tumor and peritumor liver tissues, including five cases with tumor thrombi. Protein expression of B7-H3 was significantly higher in tumors than in peritumor liver



Fig. 4 Kaplan–Meier curves of survival differences among HCC patients. Recurrence-free and overall survival for B7-H3 expression in tumor cells (**a**, **b**), instead of B7-H3 expression in tumor vessels (**c**, **d**), are statistically significant (log-rank test). Significant differences

in recurrence-free and overall survival are validated in an independent cohort according to tumor cell B7-H3 expression (\mathbf{e} , \mathbf{f}). Cores of one case were detached from the tissue microarray section during immunostaining in the validation cohort

tissues, with the most prominent expression detected in the corresponding tumor thrombi. This result was consistent with the strong staining of B7-H3 in intrahepatic metastases and venous tumor thrombosis of HCC by immunhistochemistry (Fig. 1d). Furthermore, although the numbers of CD4+, CD8+, and FOXP3+ TILs showed no significant difference between B7-H3 high and B7-H3 low tumors, flow cytometry results indicated that B7-H3 high tumors were associated with reduced proliferation and IFN- γ production by TILs. Therefore, tumor cells spreading regionally in the peritumor liver and systemically in the peripheral circulation that express B7-H3 may have a significant survival advantage. Conversely, the presence of IFN-y could significantly upregulate B7-H3 expression in HCC cells, suggesting that B7-H3 may act as a feedback mechanism to evade anti-tumor immunity. We envision a mechanistic role for B7-H3 in promoting HCC progression, based on its potential to impair T cell-mediated immunity.

In addition to cancer cells, B7-H3 is highly expressed in endothelial cells of the HCC tumor vasculature and is constitutively expressed in sinusoid endothelial cells of normal liver tissue. However, our present study failed to reveal any significant prognostic value of B7-H3 expression in the tumor vasculature of HCC. This does not exclude the biological significance of B7-H3 expression in tumor vasculature of HCC. Conversely, the question of preempting toxicity to the normal liver sinusoid endothelial cells has to be addressed in future B7-H3-targeted therapeutic strategies.

The main limitation of the present study was its retrospective nature. To compensate for this limitation, we validated the prognostic value of B7-H3 in an independent cohort of 206 HCC patients who underwent liver transplantation. Similar to hepatectomy, liver transplantation is a curative option that simultaneously removes both HCC and the diseased liver. However, as with all initial series that propose the prognostic significance of molecular biomarkers, our current findings still require independent external validation.

In conclusion, the current study indicates that B7-H3 is highly expressed in human HCC and is associated with multiple aggressive tumor features. Particularly, B7-H3 serves as an independent prognostic factor for recurrence in HCC patients who receive curative treatments like tumor resection or liver transplantation. Given the proposed immune inhibitory mechanisms of B7-H3, B7-H3 may not only help to predict which patients are at a higher risk of recurrence, thus facilitating patient selection for more aggressive treatment, but also may represent an attractive target for therapeutic manipulation in the future for multimodal management of HCC. Acknowledgments This study was supported by the Major Program of NSFC (No.81030038), National Key Sci-Tech Project (2008ZX 10002-019), Shanghai Rising-Star Program (No. 10QA1401300), National Natural Science Foundation of China (No. 30872379 & No. 30901432), and "Chen Guang" project supported by Shanghai Municipal Education Commission and Shanghai Education Development Foundation (No. 11CG02).

Conflict of interest The authors declare no competing interests to this paper.

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