

Association of family history with long-term prognosis in patients undergoing liver resection of HBV-related hepatocellular carcinoma

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Background: Family history is a risk factor for the development of hepatocellular carcinoma (HCC). The aim of the current study was to investigate the association between family history of HCC and long-term oncologic prognosis among patients undergoing curative liver resection for hepatitis B virus (HBV)-related HCC.

Methods: Patients who underwent curative liver resection of HBV-related HCC between 2003 and 2013 were consecutively enrolled. Family history was defined as a self-reported history of HCC in a first-degree relative. Propensity score matching (PSM) and multivariable Cox-regression analyses were performed to compare overall survival (OS) and recurrence-free survival (RFS) among patients with and without a family history.

Results: Among 1,112 patients, 183 (16.5%) patients had a family history of HCC. Using PSM, 179 pairs of patients with and without a family history were created that had no differences in the baseline characteristics and operative variables. On matched analysis, family history was associated with decreased OS and RFS after curative-intent resection of HBV-related HCC in the propensity matching cohort (P=0.042 and 0.006, respectively). On multivariable Cox-regression analyses, a family history of HCC was associated with decreased OS (HR: 1.574; 95% CI: 1.171–2.116; P=0.003) and RFS (HR: 1.534; 95% CI: 1.176–2.002; P=0.002) after adjusting for other prognostic risk factors.

Conclusions: Family history was associated with decreased OS and RFS rates among patients undergoing curative liver resection of HBV-related HCC.

Keywords: Hepatocellular carcinoma (HCC); hepatitis B; hepatectomy; survival; recurrence

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Introduction

Hepatocellular carcinoma (HCC) is the most frequent histologic type of primary liver cancer, ranking 6th in incidence and 3rd in mortality worldwide (1). HCC is particularly prevalent in Africa and Southeast Asia, especially in China (2). More than 75% of cases worldwide and 85% of cases in developing countries have been attributed to hepatitis B virus (HBV) and hepatitis C virus (HCV), both of which increase the risk of HCC by approximately 20-fold (3). A familial aggregation of HCC has frequently been reported in Asians, particularly in China (4-9). A meta-analysis, based on 9 case-control and 4 cohort studies, demonstrated that the pooled relative risk for a family history of HCC was 2.50 [95% confidence interval (CI): 2.06-3.03] after adjusting for other confounding factors (10). This meta-analysis also reported that the combination of a family history of HCC and chronic HBV infection was associated with an over 70fold elevated risk of HCC (10).

Despite convincing evidence that a family history of HCC is associated with HCC development, data on the association of family history with the long-term prognosis after HCC diagnosis and treatment are conflicting (11,12). One study reported that patients with HCC and a family history of HCC had better survival after multi-modalities treatment than patients without such a history (11). The authors postulated that the difference was due, in part, to earlier diagnosis, however certain genetic factors may also impact prognosis (11). Another study reported no association of family HCC history with long-term recurrence and survival after resection of HCC (12). These conflicting results may be due to different strategies in patient selection, as well as differences in the definitions of family history and treatment HCC modalities. In particular, unbalanced baseline characteristics (including demographic and clinicopathologic) among patients with and without a family history of HCC may have confounded comparisons.

The current study sought to examine the impact of a family HCC history relative to patient clinicopathologic characteristics, long-term recurrence and survival among patients undergoing curative-intent liver resection of HBVrelated HCC. In particular, we sought to define the impact of family HCC history on long-term oncologic outcomes using propensity matched analysis of a large cohort of HCC patients.

Methods

Study population

Patients presenting to two departments of Hepatic Surgery, Eastern Hepatobiliary Surgery Hospital of Shanghai, the largest tertiary hepatobiliary center in China, between August 2003 and December 2013 were included. Inclusion criteria consisted of (I) 18 years of age or older, (II) medical history of chronic HBV infection, and a positive serology of Hepatitis B surface antigen (HBsAg), (III) newly diagnosed HCC without any previous treatment, and HCC was also confirmed by postoperative histopathological examination, (IV) curative-intent liver resection for HCC, which was defined as R0 resection, and (V) complete medical record on family history and other important prognostic variables. The study was approved by the Institutional Review Board of the Eastern Hepatobiliary Surgery Hospital of Shanghai, China.

Baseline characteristics and operative variables

Baseline patient characteristics and operative variables obtained from review of the medical records were included. Cirrhosis was confirmed by histopathological examination, and portal hypertension was defined as the presence of either esophageal varices, or splenomegaly with a decrease in platelet count ($\leq 100 \times 10^9$ /L). Tumor stage at diagnosis was determined following the Barcelona Clinic Liver Cancer (BCLC) staging system. Operative variables included intraoperative blood loss, requirement of blood transfusion, extent of hepatectomy, and type of liver resection. Major hepatectomy was defined as resection of three or more Couinaud liver segments; minor hepatectomy as resection of fewer than three segments. Anatomical resections were defined by the Brisbane 2000 Nomenclature of Liver Anatomy, while non-anatomical resections included wedge resection or limited resection.

Definition of family bistory

A family history of HCC was defined as a self-reported history of HCC in a first-degree relative. First-degree relatives included parents, siblings, or children, while nieces, nephews, aunts, uncles, or grandparents were excluded.

Use of antiviral therapy and follow-up

Among patients who had a preoperative HBV-DNA

 \geq 1,000 copies/mL, adjuvant antiviral therapy with lamivudine 100 mg, adefovir dipivoxil 10 mg, or entecavir 0.5 mg orally daily was commenced immediately after surgery or after discharge. For patients with renal insufficiency, the daily lamivudine or adefovir dipivoxil dose was adjusted according to creatinine clearance.

The detailed follow-up schedule had previously been reported (13). In general, postoperative surveillance strategy for recurrence consisted of serum alpha-fetoprotein level, ultrasonography or contrast-enhanced computed tomography scan of the chest and abdomen at 2-monthly intervals for the first 6 months and at 3-monthly intervals thereafter. Computed tomography, magnetic resonance imaging, or positron emission tomography was performed when recurrence or distant metastasis was suspected. Tumor recurrence was defined as new appearance of intra- or extrahepatic tumor nodules. Tumor recurrence was divided into early and late recurrences using a cut-off value of 2 years. The management of recurrence was based on the pattern of recurrent tumor, residual hepatic functional reserve, and general condition of the patient. The treatment included re-resection, local ablation therapy, liver transplantation, transcatheter arterial chemoembolization, oral sorafenib or supportive therapy. The dates of recurrence, last follow-up, and death were recorded.

Propensity score matching (PSM)

Patients with and without a family history of HCC were matched using PSM as described by Rubin and Rosenbaum (14,15). This was carried out using the R software version 3.1.0. The propensity score for an individual was calculated using a logistic regression model given the covariates included in Table 1. This method included ordering the case and control subjects, then selecting the first case subject and finding the control subject with the closest propensity score. Afterwards, both subjects were removed from consideration for matching and the next case subject was selected (16). The forward procedure was used, which started out with just the intercept and sequentially added the effect that most improved the fit. Variables were included up to a limit of a monotonized P-to-enter value of <0.2. Thereafter, a 1:1 nearest neighbor matching without replacement was performed so as to ensure any conditional bias was minimized.

Statistical analysis

Baseline patient characteristics and operative variables

among patients with and without a family history of HCC were summarized using frequency and percentage for categorical covariates, and mean ± standard deviation (SD) or median (range) for continuous covariates. Categorical and continuous covariates were compared using the Fisher's exact test and the Wilcoxon rank-sum test, respectively. The primary outcome of the study was overall survival (OS), which was defined as the time from surgery to death resulting from any cause. The secondary outcome of this study was recurrence-free survival (RFS), which was defined as the time from surgery to tumor recurrence or occurrence of a new HCC, or death with evidence of recurrence. OS and RFS were compared among patients with and without a family history before and after PSM using the Kaplan-Meier curves and the log rank test. In order to adjust for other prognostic factors and enhance the accuracy of the model, a robust sandwich variance estimator in the multivariable Cox regression hazard regression analyses in the PSM cohort was used to estimate the hazard ratios and its 95% confidence interval. P<0.05 were considered statistically significant. Statistical analyses were carried out using the IBM SPSS Statistics version 25.0 and R software version 3.1.0.

Results

Among 1,541 patients who were screened, 429 patients did not fit the inclusion criteria and were excluded. The remaining 1,112 patients with chronic HBV infection who underwent curative liver resection for HCC were included in the final analytic cohort (*Figure 1*). There were 996 (89.6%) men and 116 (10.4%) women. The median age at operation was 50 years (range, 19–80 years). There were 183 (16.5%) patients who had a first-degree family history of HCC; 73.2% of patients had cirrhosis and 32.2% had portal hypertension.

Comparison of baseline characteristics and operative variables among patients with and without a family history of HCC are illustrated in *Table 1*. Several clinicopathological features were significantly different among patients with a family history versus patients without a history of HCC such as performance status, largest tumor diameter, tumor number, and presence of satellites (all P<0.05). Early HCC (BCLC A stage) among patients with a family history of HCC was also more common versus patients with no family history of HCC (43.2% *vs.* 26.9%, P<0.001).

PSM was used to create 179 pairs of patients. Patient characteristics and operative variables among patients with

Table 1 Comparisons of patients'	baseline abarratoristics and	oporativo variables before (nd after properties of	oro motohing
Table 1 Comparisons of patients	Dasenne characteristics and	operative variables before a	ind after propensity so	core matching

	The entire cohort		The PSM cohort			
Variables	Without a FH (N=929)	With a FH (N=183)	Р	Without a FH (N=179)	With a FH (N=179)	Р
Age, years	49.9±10.3	49.2±10.0	0.679	49.8±10.9	49.1±10.1	0.278
≤60	779 (83.9)	158 (86.3)	0.438	145 (81.0)	154 (86.0)	0.254
>60	150 (16.1)	25 (13.7)		34 (19.0)	25 (14.0)	
Sex						
Male	827 (89.0)	169 (92.3)	0.233	171 (95.5)	165 (92.2)	0.271
Female	102 (11.0)	14 (7.7)		8 (4.5)	14 (7.8)	
Diabetes mellitus	57 (6.1)	10 (5.5)	0.865	10 (5.6)	10 (5.6)	1.000
Cigarette smoking	290 (31.2)	53 (29.0)	0.600	53 (29.6)	53 (29.6)	1.000
Alcohol drinking	167 (18.0)	35 (19.1)	0.753	33 (18.4)	33 (18.4)	1.000
BMI, kg/m ²	23.9±3.4	24.1±3.4	0.485	23.8±3.3	24.0±3.4	0.256
≤24.0	517 (55.7)	97 (53.0)	0.516	109 (60.9)	97 (54.2)	0.239
>24.0	412 (44.3)	86 (47.0)		70 (39.1)	82 (45.8)	
ASA score						
≤2	833 (89.7)	161 (88.0)	0.512	157 (87.7)	157 (87.7)	1.000
>2	96 (10.3)	22 (12.0)		22 (12.3)	22 (12.3)	
Cirrhosis	674 (72.6)	140 (76.5)	0.315	127 (70.9)	136 (76.0)	0.338
Portal hypertension	292 (31.4)	66 (36.1)	0.226	70 (39.1)	63 (35.2)	0.512
Child-Pugh grade						
A	826 (88.9)	162 (88.5)	0.898	161 (89.9)	158 (88.3)	0.735
В	103 (11.1)	21 (11.5)		18 (10.1)	21 (11.7)	
Preoperative HBV viral load						
≤10,000 copies/mL	459 (49.7)	76 (41.5)	0.043	79 (44.1)	75 (41.9)	0.749
>10,000 copies/mL	470 (50.3)	107 (58.5)		100 (55.9)	104 (58.1)	
Anti-HBV therapy	408 (43.9)	99 (54.1)	0.011	91 (50.8)	93 (52.0)	0.833
HBeAg (+)	238 (25.6)	55 (30.1)	0.233	50 (27.9)	52 (29.1)	0.907
Preoperative AST level, U/L	56.2±47.2	53.3±34.4	0.427	55.8±49.0	52.3±33.9	0.247
≤80	785 (84.5)	152 (83.1)	0.657	142 (79.3)	151 (84.4)	0.273
>80	144 (15.5)	31 (16.9)		37 (20.7)	28 (15.6)	
ECOG performance status						
0	473 (50.9)	112 (61.2)	0.012	119 (66.5)	108 (60.3)	0.273
1–2	456 (49.1)	71 (38.8)		60 (33.5)	71 (39.7)	

Table 1 (continued)

Table 1	(continued)
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	The	The entire cohort			The PSM cohort		
Variables	Without a FH (N=929)	With a FH (N=183)	Р	Without a FH (N=179)	With a FH (N=179)	Р	
Preoperative AFP level							
≤400 μg/L	547 (58.9)	105 (57.4)	0.743	107 (59.8)	105 (58.7)	0.914	
>400 µg/L	382 (41.1)	78 (42.6)		72 (40.2)	74 (41.3)		
Largest tumor diameter							
≤5 cm	413 (44.5)	104 (56.8)	0.003	104 (58.1)	101 (56.4)	0.831	
>5 cm	516 (55.6)	79 (43.2)		75 (41.9)	78 (43.6)		
Tumor number							
Solitary	680 (73.2)	148 (80.9)	0.033	137 (76.5)	144 (80.4)	0.440	
Multiple	249 (26.8)	35 (19.1)		42 (23.5)	35 (19.6)		
Tumor rupture	49 (5.3)	6 (3.3)	0.350	5 (2.8)	6 (3.4)	1.000	
Macroscopic vascular invasion	127 (13.7)	22 (12.0)	0.635	25 (14.0)	21 (11.7)	0.636	
Microscopic vascular invasion	542 (58.3)	104 (56.8)	0.743	97 (54.2)	102 (57.0)	0.671	
Satellites	270 (29.1)	37 (20.2)	0.015	43 (24.0)	37 (20.7)	0.526	
Tumor differentiation							
Well or moderately	152 (16.4)	36 (19.7)	0.281	37 (20.7)	34 (19.0)	0.791	
Poorly	777 (83.6)	147 (80.3)		142 (79.3)	145 (81.0)		
BCLC staging							
A (early)	250 (26.9)	79 (43.2)	<0.001	73 (40.8)	76 (42.5)	0.523	
B (intermediate)	231 (24.9)	26 (14.2)		34 (19.0)	26 (14.5)		
C (advanced)	448 (48.2)	78 (42.6)		72 (40.2)	77 (43.0)		
Intraoperative blood loss							
≤400 mL	549 (59.1)	109 (59.6)	0.935	114 (63.7)	108 (60.3)	0.586	
>400 mL	380 (40.9)	74 (40.4)		65 (36.3)	71 (39.7)		
Blood transfusion	211 (22.7)	45 (24.6)	0.566	45 (25.1)	45 (25.1)	1.000	
Extent of hepatectomy							
Major hepatectomy	267 (28.7)	53 (29.0)	0.952	45 (25.1)	52 (29.1)	0.476	
Minor hepatectomy	662 (71.3)	130 (71.0)		134 (74.9)	127 (70.9)		
Type of resection							
Anatomical	295 (31.8)	56 (30.6)	0.795	63 (35.2)	56 (31.3)	0.501	
Non-anatomical	634 (68.2)	127 (69.4)		116 (64.8)	123 (68.7)		

Values are mean ± standard deviation or median (percentage) unless otherwise indicated. AFP, alpha-fetoprotein; ASA, American Society of Anesthesiologists; AST, aspartate aminotransferase; BCLC, Barcelona Clinic Liver Cancer; BMI, body mass index; ECOG, Eastern Cooperative Oncology Group; FH, family history; HBeAg, hepatitis B e antigen; HBV, hepatitis B virus; PSM, propensity score matching.

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Figure 1 CONSORT diagram of study population. HBV, hepatitis B virus; HCC, hepatocellular carcinoma; FH, family history; PSM, propensity score matching.

and without a family history after PSM are illustrated in *Table 1*. Of note, there were no differences in any of the baseline characteristics among patients with and without a family history of HCC (all P>0.2) after PSM.

Comparisons of long-term outcomes among patients with and without a family history of HCC are illustrated in *Table 2*. After PSM, there were no differences in early recurrence among patients with and without a family history (40.2% *vs.* 34.6%, P=0.326). However, the total recurrence, late recurrence, and mortality among patients with a family history were higher versus patients without such a history (75.4% *vs.* 53.6%, P<0.001, 35.2% *vs.* 19.0%, P=0.001, and 60.9% *vs.* 47.5%, P=0.015, respectively).

Before PSM, the 3-, 5-, and 10-year OS among patients with and without a family history of HCC were 70.5%, 57.1%, and 29.9%, and 67.2%, 56.5%, and 33.0%, respectively (*Figure 2A*). The 3-, 5-, and 10-year RFS among patients with and without a family history of HCC were 47.5%, 35.4%, and 13.1%, and 49.6%, 38.7%, and 19.1%, respectively (*Figure 2B*). Family history was not associated with increased risk of OS [hazard ratio (HR): 0.999; 95% CI: 0.814–1.226; P=0.994] and RFS (HR: 1.076; 95% CI:

0.897–1.292; P=0.428). After PSM, the 3-, 5-, and 10-year OS among patients with and without a family history of HCC were 69.8%, 56.1%, and 30.4%, and 72.5%, 64.5%, and 43.7%, respectively (*Figure 3A*). The 3-, 5-, and 10-year RFS among patients with and without a family history of HCC were 46.9%, 34.5%, and 12.7%, and 58.0%, 49.6%, and 26.0%, respectively (*Figure 3B*). Of note, after PSM, family history was associated with decreased risks of OS (HR: 1.342; 95% CI: 1.010–1.784; P=0.042) and RFS (HR: 1.420; 95% CI: 1.420–1.826; P=0.006). *Figure 4* showed the comparisons of late recurrence (>2 years after surgery) rate between patients with and without a family history in the PSM cohort (P<0.001).

Univariable and multivariable Cox-regression analyses of OS and RFS after curative-intent liver resection of HBV-related HCC in the PSM cohort are shown in *Tables 3,4*. After univariable analysis, variables with P<0.1 were entered in the multivariable analysis. On multivariable Coxregression analyses with robust estimator, after adjustment for other prognostic factors, a family history was independently associated with decreased OS and RFS after curative liver resection of HBV-related HCC. The adjusted HRs for OS

Table 2 Comparisons of long-term outcomes before and after propensity score matching

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The entire cohort			The PSM cohort		
Without a FH (N=929)	With a FH (N=183)	Ρ	Without a FH (N=179)	With a FH (N=179)	Р
58.8±40.7	62.0±41.5	0.250	64.5±43.4	62.0±41.6	0.566
586 (63.1)	136 (74.3)	0.004	96 (53.6)	135 (75.4)	<0.001
381 (41.0)	72 (39.3)	0.742	62 (34.6)	72 (40.2)	0.326
205 (22.1)	64 (35.0)	<0.001	34 (19.0)	63 (35.2)	0.001
531 (57.2)	111 (60.7)	0.413	85 (47.5)	109 (60.9)	0.015
77.0±4.4	72.8±8.5	0.994	109.0±12.0	68.8±9.3	0.042
88.4	88.5		88.3	88.3	
67.2	70.5		72.5	69.8	
56.5	57.1		64.5	56.1	
33.0	29.9		43.7	30.4	
35.5±3.1	33.1±2.6	0.428	57.7±12.0	33.1±2.7	0.006
70.1	74.9		71.5	74.3	
49.6	47.5		58.0	46.9	
38.7	35.4		49.6	34.5	
19.1	13.1		26.0	12.7	
	The Without a FH (N=929) 58.8±40.7 586 (63.1) 381 (41.0) 205 (22.1) 531 (57.2) 77.0±4.4 88.4 67.2 56.5 33.0 35.5±3.1 70.1 49.6 38.7 19.1	The entire cohortWithout a FH (N=929)With a FH (N=183) 58.8 ± 40.7 62.0 ± 41.5 586 (63.1) 136 (74.3) 381 (41.0) 72 (39.3) 205 (22.1) 64 (35.0) 531 (57.2) 111 (60.7) 77.0 ± 4.4 72.8 ± 8.5 88.4 88.5 67.2 70.5 56.5 57.1 33.0 29.9 35.5 ± 3.1 33.1 ± 2.6 70.1 74.9 49.6 47.5 38.7 35.4 19.1 13.1	The entire cohortWithout a FH (N=929)With a FH (N=183)P 58.8 ± 40.7 62.0 ± 41.5 0.250 586 (63.1) 136 (74.3) 0.004 381 (41.0) 72 (39.3) 0.742 205 (22.1) 64 (35.0) <0.001 531 (57.2) 111 (60.7) 0.413 77.0 ± 4.4 72.8 ± 8.5 0.994 88.4 88.5 67.2 70.5 56.5 57.1 33.0 29.9 35.5 ± 3.1 33.1 ± 2.6 0.428 70.1 74.9 49.6 47.5 38.7 35.4 19.1 13.1	$\begin{tabular}{ c c c c c } \hline The entire cohort & The entire cohort & The Without a FH (N=1929) & With a FH (N=183) & P & Without a FH (N=179) & 58.8 \pm 40.7 & 62.0 \pm 41.5 & 0.250 & 64.5 \pm 43.4 & 586 & (63.1) & 136 & (74.3) & 0.004 & 96 & (53.6) & 381 & (41.0) & 72 & (39.3) & 0.742 & 62 & (34.6) & 205 & (22.1) & 64 & (35.0) & <0.001 & 34 & (19.0) & 531 & (57.2) & 111 & (60.7) & 0.413 & 85 & (47.5) & 77.0 \pm 4.4 & 72.8 \pm 8.5 & 0.994 & 109.0 \pm 12.0 & 88.4 & 88.5 & 88.3 & 67.2 & 70.5 & 72.5 & 56.5 & 57.1 & 64.5 & 33.0 & 29.9 & 43.7 & 35.5 \pm 3.1 & 33.1 \pm 2.6 & 0.428 & 57.7 \pm 12.0 & 70.1 & 74.9 & 71.5 & 49.6 & 47.5 & 58.0 & 38.7 & 35.4 & 49.6 & 19.1 & 13.1 & 26.0 & \end{tabular}$	$\begin{tabular}{ c c c c c } \hline The entire cohort & The PSM cohort \\ \hline Without a FH (N=929) & With a FH (N=183) & P & Without a FH (N=179) & With a FH (N=179) \\ \hline 8.8 ± 40.7 & 62.0 ± 41.5 & 0.250 & 64.5 ± 43.4 & 62.0 ± 41.6 \\ \hline $586 (63.1) & 136 (74.3) & 0.004 & $96 (53.6)$ & $135 (75.4)$ \\ \hline $381 (41.0) & 72 (39.3) & 0.742 & $62 (34.6)$ & $72 (40.2)$ \\ \hline $205 (22.1) & $64 (35.0)$ & <0.001 & $34 (19.0)$ & $63 (35.2)$ \\ \hline $531 (57.2) $ & $111 (60.7)$ & 0.413 & $85 (47.5)$ & $109 (60.9)$ \\ \hline 77.0 ± 4.4 & 72.8 ± 8.5 & 0.994 & 109.0 ± 12.0 & 68.8 ± 9.3 \\ \hline 88.4 & 88.5 & 88.3 & 88.3 \\ \hline 67.2 & 70.5 & 72.5 & 69.8 \\ \hline 56.5 & 57.1 & 64.5 & 56.1 \\ \hline 33.0 & 29.9 & 43.7 & 30.4 \\ \hline 35.5 ± 3.1 & 33.1 ± 2.6 & 0.428 & 57.7 ± 12.0 & 33.1 ± 2.7 \\ \hline 70.1 & 74.9 & 71.5 & 74.3 \\ \hline 49.6 & 47.5 & 58.0 & 46.9 \\ \hline 38.7 & 35.4 & 49.6 & 34.5 \\ \hline 19.1 & 13.1 & 26.0 & 12.7 \\ \hline \end{tabular}$

*, values are mean ± standard deviation; **, values are median ± standard error. FH, family history; OS, overall survival; PSM, propensity score matching; RFS, recurrence-free survival.



Figure 2 Overall survival (A) and recurrence-free survival (B) curves comparisons between patients with and without a family history (FH) in the entire cohort.



Figure 3 Overall survival (A) and recurrence-free survival (B) curves comparisons between patients with and without a family history (FH) in the propensity score matching (PSM) cohort.



Figure 4 Comparisons of late recurrence (>2 years after surgery) rate between patients with and without a family history (FH) in the propensity score matching (PSM) cohort.

and RFS were 1.574 (95% CI: 1.171–2.116; P=0.003) and 1.534 (95% CI: 1.176–2.002; P=0.002), respectively.

Discussion

In recent years, the association between family history and long-term oncologic prognosis has been studied in many malignant tumors, including colon cancer, stage III or IV gastric cancer, breast cancer, prostate cancer, and lung cancer (17-21). A large number of studies have identified that a family history of HCC increases the risk of HCC development and such a history increases the risk of developing HCC by 4- to 32-folds in patients with chronic HBV infection (4,7,10,22-27). However, very few studies have evaluated the impact of family history on longterm prognosis after liver resection of HCC. This large cohort study demonstrated that a family history of HCC was indeed associated with decreased OS and RFS after liver resection of HBV-related HCC, even after adjusting for the potential risk factors of patient's demographic, environmental and clinicopathological characteristics.

Family history was based on self-reported information from the patients and only those patients with a family history of HCC in their first-degree relatives were enrolled and analyzed. The 2001 population-based Connecticut family health study noted that reports from first-degree relatives were more accurate than information from the seconddegree relatives, with positive predictive values varying between 78% and 80% for lung and breast cancers (28). Numerous studies have also reported that only a family history of HCC in the first-degree relative increased the risk of developing HCC, but not the second-degree relatives (4,10,22). As a consequence, only patients with a family history of HCC in their first-degree relatives were enrolled in this study and second-degree relatives were excluded. In fact, 183 of 1,112 patients (16.4%) had a family history of HCC among their first-degree relatives. This proportion of patients with a family HCC history was similar to other reports. For example, a case-control study, Turati et al. (10) observed that 25 of 204 HCC patients (12.3%) had a first-degree relative with HCC in Western population of patients. In another study, Yu et al. (4) from Taiwan, observed that 17.5% of 553 male patients who had chronic

Table 3 Univariable and multivariable Cox regression analyses of overall survival in the propensity score matching cohort

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Variables		Univariable analy	sis	Multivariable analysis		
	HR companson	HR (95% CI)	Р	HR (95% CI)	P*	
Family history	Yes vs. no	1.342 (1.010–1.784)	0.042	1.574 (1.171–2.116)	0.003	
Age	≤60 <i>vs</i> . >60 years	1.256 (0.874–1.805)	0.216	_	-	
Sex	Male vs. female	1.118 (0.591–2.114)	0.732	-	-	
Diabetes mellitus	Yes vs. no	1.307 (0.711–2.404)	0.387	-	-	
Cigarette smoking	Ever vs. never	1.125 (0.827–1.531)	0.453	-	-	
Alcohol drinking	Ever vs. never	1.197 (0.863–1.660)	0.280	-	-	
BMI	≤24.0 <i>vs</i> . >24.0 kg/m ²	1.201 (0.901–1.462)	0.105	_	-	
ASA score	≤2 <i>v</i> s. >2	0.973 (0.634–1.492)	0.900	_	-	
Cirrhosis	Yes vs. no	2.191 (1.503–3.192)	<0.001	NS	0.178	
Portal hypertension	Yes vs. no	1.433 (1.078–1.904)	0.013	1.661 (1.224–2.254)	0.001	
Child-Pugh grade	A vs. B	1.501 (0.992–2.272)	0.053	NS	0.440	
Preoperative HBV viral load	$\leq 10^4$ vs. >10 ⁴ copies/mL	1.667 (1.220–2.277)	0.001	1.660 (1.185–2.326)	0.003	
Anti-HBV therapy	Yes vs. no	1.573 (1.052–2.033)	0.037	NS	0.103	
HBeAg (+)	Yes vs. No	1.281 (0.944–1.738)	0.112	_	-	
Preoperative AST level	≤80 <i>vs</i> . >80 U/L	1.760 (1.254–2.472)	0.001	NS	0.892	
ECOG performance status	0 vs. 1–2	2.035 (1.530–2.707)	<0.001	NS	0.588	
Preoperative AFP level	≤400 <i>v</i> s. >400 μg/L	1.643 (1.239–2.180)	0.001	1.396 (1.014–1.924)	0.041	
Largest tumor diameter	≤5 <i>v</i> s. >5 cm	2.011 (1.515–2.669)	<0.001	1.864 (1.088–3.192)	0.023	
Tumor number	Solitary vs. multiple	3.221 (2.354–4.406)	<0.001	1.994 (1.178–3.375)	0.010	
Tumor rupture	Yes vs. no	2.894 (1.527–5.485)	0.001	NS	0.111	
Macroscopic vascular invasion	Yes vs. no	7.868 (5.365–11.537)	<0.001	4.670 (3.004–7.258)	<0.001	
Microscopic vascular invasion	Yes vs. no	1.833 (1.370–2.452)	<0.001	1.355 (1.036–1.772)	0.027	
Satellites	Yes vs. no	3.398 (2.489–4.638)	<0.001	2.262 (1.235–4.141)	0.008	
Tumor differentiation	Well or moderately vs. poorly	2.174 (1.437–3.289)	<0.001	1.894 (1.214–2.955)	0.005	
Intraoperative blood loss	≤400 <i>v</i> s. >400 mL	2.027 (1.524–2.696)	<0.001	NS	0.701	
Blood transfusion	Yes vs. no	2.887 (2.118–3.936)	<0.001	NS	0.134	
Extent of hepatectomy	Major vs. minor	2.583 (1.919–3.476)	<0.001	NS	0.129	
Type of resection	Anatomical vs. non-anatomical	1.073 (0.797–1.446)	0.642	-	-	

*, Those variables found significant at P<0.1 in univariable analyses were entered into multivariable Cox regression models. AFP, alphafetoprotein; ASA, American Society of Anesthesiologists; AST, aspartate aminotransferase; BMI, body mass index; CI, confidence interval; ECOG, Eastern Cooperative Oncology Group; HBeAg, hepatitis B e antigen; HBV, hepatitis B virus; HR, hazard ratio; NS, not significant.

Table 4 Univariable and multivariable Cox regression analyses of recurrence-free survival in the propensity score matching cohort

Veriables		Univariable analys	sis	Multivariable analysis	
variables	nn companson	HR (95% CI)	Р	HR (95% CI)	P*
Family history	Yes vs. no	1.420 (1.104–1.826)	0.006	1.997 (1.499–2.659)	<0.001
Age	≤60 <i>vs</i> . >60 years	1.021 (0.725–1.438)	0.905	-	-
Sex	Male vs. female	0.918 (0.544–1.548)	0.748	-	-
Diabetes mellitus	Yes vs. no	1.540 (0.927–2.558)	0.093	NS	0.118
Cigarette smoking	Ever vs. never	1.098 (0.836–1.442)	0.503	-	-
Alcohol drinking	Ever vs. never	1.300 (0.980–1.725)	0.068	NS	0.182
BMI	≤24.0 vs. >24.0 kg/m ²	1.190 (0.926–1.529)	0.174	-	-
ASA score	≤2 <i>vs.</i> >2	1.077 (0.739–1.570)	0.700	-	-
Cirrhosis	Yes vs. no	1.811 (1.326–2.473)	<0.001	NS	0.412
Portal hypertension	Yes vs. no	1.334 (1.035–1.720)	0.025	1.394 (1.063–1.827)	0.016
Child-Pugh grade	A vs. B	1.989 (1.390–2.848)	<0.001	NS	0.162
Preoperative HBV viral load	$\leq 10^4$ vs. $> 10^4$ copies/mL	1.821 (1.381–2.402)	<0.001	1.534 (1.176–2.002)	0.002
Anti-HBV therapy	Yes vs. no	1.437 (1.077–2.253)	0.019	NS	0.175
HBeAg (+)	Yes vs. no	1.355 (1.036–1.771)	0.026	NS	0.543
Preoperative AST level	≤80 <i>vs</i> . >80 U/L	1.811 (1.335–2.456)	<0.001	NS	0.798
ECOG performance status	0 vs. 1–2	1.606 (1.244–2.072)	<0.001	NS	0.455
Preoperative AFP level	≤400 <i>v</i> s. >400 µg/L	1.241 (0.964–1.599)	0.093	1.783 (1.065–2.985)	0.028
Largest tumor diameter	≤5 <i>vs</i> . >5 cm	1.835 (1.430–2.357)	<0.001	1.464 (1.095–1.958)	0.010
Tumor number	Solitary vs. multiple	2.643 (1.981–3.526)	<0.001	1.678 (1.251–2.250)	0.001
Tumor rupture	Yes vs. no	2.458 (1.300–4.646)	0.004	1.801 (1.076–3.014)	0.025
Macroscopic vascular invasion	Yes vs. no	8.055 (5.643–11.496)	<0.001	6.140 (4.177–9.026)	<0.001
Microscopic vascular invasion	Yes vs. no	1.493 (1.160–1.922)	0.002	1.463 (1.091–1.962)	0.011
Satellites	Yes vs. no	2.660 (1.996–3.544)	<0.001	1.621 (1.181–2.225)	0.003
Tumor differentiation	Well or moderately vs. poorly	1.497 (1.080–2.074)	0.015	NS	0.213
Intraoperative blood loss	≤400 <i>vs</i> . >400 mL	1.760 (1.366–2.267)	<0.001	NS	0.340
Blood transfusion	Yes vs. no	2.644 (1.988–3.516)	<0.001	NS	0.185
Extent of hepatectomy	Major vs. minor	2.284 (1.744–2.990)	<0.001	NS	0.558
Type of resection	Anatomical vs. non-anatomical	1.121 (0.862–1.459)	0.393	-	-

*, Those variables found significant at P<0.1 in univariable analyses were entered into multivariable Cox regression models. AFP, alphafetoprotein; ASA, American Society of Anesthesiologists; AST, aspartate aminotransferase; BMI, body mass index; CI, confidence interval; ECOG, Eastern Cooperative Oncology Group; HBeAg, hepatitis B e antigen; HBV, hepatitis B virus; HR, hazard ratio; NS, not significant. HBV infection and HCC reported to a family history of HCC in a first-degree relative. Similarly, in another study from China, 169 of 1,313 HCC patients (12.9%) gave a history of HCC in a first-degree relative (12).

The strengths of the present study included the large sample size, the long-term follow-up, the standard definition used in the family history of HCC (restricted to the firstdegree relatives), and the attempt to control for potential confounders by using the PSM method and multivariable Cox-regression analysis. PSM analysis was carried out to balance the differences in baseline variables among patients with and without a family history of HCC (16). After PSM, the real impact of family history on the prognosis of HCC after resection was more able to be determined. In addition, to further adjust for other confounding prognostic factors, a multivariable Cox regression analysis was applied to the PSM cohort. A family history was independently associated with decreased OS or RFS after resection for patients with HBV-related HCC. In the current study, rates of early recurrence (≤ 2 years after HCC resection) among patients with and without a family history appeared to be similar. Many previous studies have reported consistently that early recurrence within 2 years after resection is most likely the consequence of occult metastasis from the initial tumor, which is associated with aggressive tumor pathologic factors, such as large tumor size, multiple tumors, poor differentiation, macro- and microvascular invasion, and satellite lesions (29-31). In this way, it is understandable that family history was not associated with early recurrence after curative liver resection of HCC, as revealed by the result of this present study. However, the rate of late recurrence (>2 years after HCC resection) among patients with a family history was much higher (35.2% vs. 19.0%, P=0.001). Interestingly, the RFS curves among patients with or without a family history gradually separated after 2 years from the date of surgery (Figure 3B). These data suggest that late recurrence in patients with a family history may be higher than in patients without such a history. Apart from a small portion of late recurrence due to occult metastasis from the initial tumor, most late recurrence after 2 years of resection are commonly considered to develop from new malignant clones of HCC or de novo carcinogenesis (29). Thus, it is reasonable to assume that the genetic make-up of an individual contributes not only to the development of HCC, but also to the long-term prognosis after liver resection (28,32). Therefore, this significant difference in RFS, OS, and late recurrence between patients with and without a family history probably calls for closer and

more stringent recurrence surveillance for patients with a family history in the late period of postresection followup, which may be helpful to early find and early treat those recurrent HCCs and improve the long-term prognosis after recurrence in our clinical practice.

A familial clustering of HCC in HBV carriers can be explained by shared HBV contagious infection among the first-degree relatives, inherent genetic contributions, and environmental or health-behavioral risk factors (4-9,25). In the present study, data on the environmental variables were prospectively collected such that their potential impact on tumor recurrence and survival after HCC resection could be investigated. Of note, there were no differences in the proportions of diabetes mellitus, cigarette smoking, alcohol drinking, and obesity among patients with and without a family history. Moreover, none of these factors were associated with RFS and OS after HCC resection, although some of these factors had been reported to be closely related to the development of HCC in previous epidemiological studies (7,32).

The analytic cohort in the current study consists of patients with HBV-related HCC. Previous studies by our group and others have demonstrated that preoperative HBV-DNA level over 10,000 copies/mL was an independent risk factor of OS and RFS after liver resection and anti-HBV therapy decreased HCC recurrence and prolonged survival for these patients (33,34). In the present study, a preoperative HBV-DNA level of over 10,000 copies/mL was indeed demonstrated to be an independent risk factor of OS and RFS. However, we did not find a beneficial impact of anti-HBV therapy on OS and RFS after PSM. The reasons for this difference are likely multifactorial and may be related to specific treatment regimens, treatment duration, discontinuation or adjustment after viral reactivation (35). It's well-recognized that the eradication of HBV plays an important role in decreasing the development of HCC, as well as reducing HCC recurrence after curative resection. A recent randomized controlled trial study even showed that anti-HBV therapy can be effective for anti-recurrence in patients with low HBV-DNA load (<10,000 copies/mL) (35). However, in the study period [2003–2013], there were no guidelines in China even in the world which recommended antiviral therapy for those patients with low HBV load. The cost of antiviral drugs every day is not a small economic burden for most patients with chronic HBV infection in China as a developing country, especially for those who don't have full health insurance. Although we recommend antiviral therapy for all patients with HBV-related HCC,

regardless of preoperative HBV load, this issue is actually very complicated in the real clinical practice.

The current study had several limitations. Self-reporting of the family history was used and HCC family history was not confirmed with any relatives. This method may have resulted in under-reporting of the family history of HCC. As reporting of a family history is always more accurate for the first-degree relatives, we purposely did not extend to the second- and third-degree relatives. Although selfreporting the family history was limited to the first-degree relatives, this allowed greater accuracy (36,37). There was also a lack of information in the family history of chronic HBV infection in the first-degree relatives with HCC. HBV transmission among family members, together with other shared environmental risk factors, may be responsible for part of the observed familial aggregation of HCC. In addition, although the study indicated that the poor prognosis associated with a family history of HCC might be attributed to genetic contributions, defining the underlying mechanism was beyond the scope of the present study. Further studies involving tumor biology and genetic contributions in this group of patients with HCC and family history are required, especially for HBV-related HCC.

In conclusion, family history of HCC in a first-degree relative was associated with a worse OS and RFS after curative-intent resection among patients with HBV-related HCC. The genetic contributions of a family history might increase the risk of HCC recurrence after resection. To better assess HCC susceptibility in a population where HBV is endemic, future studies exploring the underlying mechanisms on the impact of genetic contributions on the development and recurrence of HCC in patients with chronic HBV infection are warranted.

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Footnote

Conflicts of Interest: The abstract of this study has been presented as oral presentation at the 13th World Congress of the International Hepato-Pancreato-Biliary Association (4-7 Sep 2018, Gevena, Switzerland).

Ethical Statement: The study was approved by the

Institutional Review Board of the Eastern Hepatobiliary Surgery Hospital of Shanghai, China.

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