# Does microvascular invasion in Barcelona Clinic Liver Cancer stage A multinodular hepatocellular carcinoma indicate earlystage behavior?

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**Background:** To identify the impact of tumor number on Barcelona Clinic Liver Cancer (BCLC) earlystage hepatocellular carcinoma (HCC) and the impact of microvascular invasion (MVI) on multinodular HCC (MHCC).

**Methods:** We retrospectively analyzed 1,548 patients who had early-stage HCC [solitary HCC (SHCC, n=1,481) and MHCC (n=67)], according to the BCLC classification, after curative resection. Recurrence-free survival (RFS) and overall survival (OS) were compared. Propensity score matching (PSM) was used to balance potential confounding factors.

**Results:** Both before and after PSM, significant differences were noted between the MHCC group and the SHCC group in RFS but not in OS. For the PSM cohort, the 5-year RFS rates were 7.5% and 41.2% for the MVI-positive MHCC group and the SHCC group, respectively (P<0.001). The 5-year OS rates were 48.9% and 75.2% for the MVI-positive MHCC group and the SHCC group, respectively (P=0.017). The RFS and OS were not significantly different between the MVI-negative MHCC group and the SHCC group. MVI (P=0.029) and multiple nodules (P=0.029) were associated with early recurrence.

**Conclusions:** The presence of MVI in BCLC early-stage MHCC was highly suggestive of a poor prognosis and should not be classified as early-stage biological behavior.

**Keywords:** Hepatocellular carcinoma (HCC); Barcelona Clinic Liver Cancer (BCLC); early stage; microvascular invasion (MVI); multiple nodules

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## Page 2 of 12

## Introduction

Hepatocellular carcinoma (HCC) is one of the most common causes of cancer-related deaths worldwide (1,2). To date, numerous HCC staging systems have been proposed to classify patients for better prognostic prediction and treatment decisions (3,4). Among these classification systems, the Barcelona Clinic Liver Cancer (BCLC) classification is the most widely used system and is recommended by the European Association for the Study of the Liver (EASL) and the American Association for the Study of Liver Diseases (AASLD) (5-7). Radiofrequency ablation, surgical resection and liver transplantation are regarded as curative treatment methods and are recommended for patients with BCLC early-stage (Stage 0 and Stage A) disease (8). Multinodular HCC (MHCC) with up to 3 nodules, each less than 3 cm in size, but without major vascular invasion and extrahepatic metastasis was also divided into early stages, and patients with MHCC are suggested for liver transplantation (9). However, since transplantation is limited by donor organ availability, surgical resection remains an appropriate choice for these patients (10,11).

Recent progress has led to the recognition that both intrahepatic metastasis (IM) and multicentric occurrence (MO) contribute to multiple tumor nodules in the liver (12). IM-type HCC is accompanied by a worse biological behavior than MO-type HCC and does not benefit from curative therapy (13,14). In addition, a previous study demonstrated that tumor biology and the condition of the underlying liver were better prognostic factors than tumor size and should be given closer attention (15). As such, we hypothesized that the IM-type original pattern also exists in BCLC early-stage MHCC and represents a poor prognosis. Interestingly, microvascular invasion (MVI), which is identified as the presence of tumor emboli in a vascular space on microscopy, highly indicates the possibility of early recurrence and metastasis (16,17). Numerous studies have shown meaningful associations between the presence of MVI and IM-type HCC (13,18). Therefore, in this study, we aimed to analyze the clinicopathological data of BCLC early-stage HCC patients to explore the significance of tumor number and MVI on prognostic and biological behavior and to illustrate whether IM-type MHCC exists in these patients, which might enable clinicians to better assess whether patients are suitable for surgery and to implement reasonable pre- and post-operative management.

## **Methods**

# Patients

The consecutive patients who underwent hepatic resection for BCLC early-stage HCC from December 2009 to December 2010 at the Eastern Hepatobiliary Surgery Hospital were identified. The inclusion criteria were as follows: (I) HCC within solitary nodules or up to 3 multiple nodules, with no nodules >3 cm, (II) Child-Pugh A-B, (III) Eastern Cooperative Oncology Group (ECOG) score =0, and (IV) underwent curative resection. The exclusion criteria were as follows: (I) major vascular invasion, (II) extrahepatic metastasis, (III) preoperative anticancer treatments, and (IV) a previous history of other malignancies.

A 7-point baseline sampling protocol was performed to evaluate the pathological parameters (19). HCC was diagnosed by two pathologists in all cases. MVI was defined as the presence of tumor cells in a portal vein, hepatic vein, or large capsular vessel of the surrounding hepatic tissue lined by endothelium that was visible only on microscopy (19). The clonal origin of MHCC was evaluated by the comprehensive criteria pertaining to tumor grade and histological type (20). The reported diameter and capsule are based on the tumor with the largest diameter. The reported Edmondson-Steiner grade (III-IV or I-II) is the highest grade found in the specimen.

# Follow-up

The patients were followed up once every 2 months in the first year and once every 3 months thereafter. The follow-up investigations consisted of ultrasonographic scans, computed tomography scans, or magnetic resonance imaging with serum alpha-fetoprotein (AFP). The study endpoints were recurrence-free survival (RFS) and overall survival (OS). RFS was calculated from the date of surgery to the date when recurrence was diagnosed. OS was defined as the duration between surgery and the last follow-up visit or HCC-related death. The patients were followed up for 84 months.

## Statistical analysis

Continuous variables are expressed as the mean  $\pm$  standard deviation. Chi-squared tests or Fisher's exact tests were used to compare categorical data. Continuous data were analyzed with Student's *t*-tests or Mann-Whitney U tests,

when appropriate. Survival analyses of RFS and OS were performed by the Kaplan-Meier method with the log-rank test. A Cox proportional hazards regression model was used to develop a multivariable model. Variables with a P value less than 0.1 in the univariable analysis were entered into the multivariable model. Logistic regression models were used to determine the predictors associated with the presence of MVI. Variables with a P value less than 0.2 in the univariable analysis were entered into the multivariable model.

The effect of selection bias and confounding factors was reduced by using propensity score matching (PSM) (21). All variables with potential differences (P<0.2) were entered into the PSM model, including sex, aspartate aminotransferase (AST), alkaline phosphatase (ALP), white blood cells (WBCs), platelets (PLTs), international normalized ratio (INR), hepatitis B e antibody (HBeAb), use of Pringle maneuver, use of transfusion, capsule, and presence of cirrhosis. Considering the strong correlation between prothrombin time (PT) and INR, we selected INR for propensity matching. Furthermore, we did not include tumor diameter for propensity matching as BCLC earlystage solitary HCC (SHCC) was not bound by tumor size. A logistic regression analysis was performed using nearest neighbor matching to estimate the propensity score. The ratio for matching was established at 1:2 using a caliper width of 0.1 of the standard deviation of the logit of the propensity score. The absolute standardized difference was computed to assess the balance of matched variables and to confirm whether the values were lower than 0.1 (22). In addition, the discrimination of the propensity score model was assessed using the area under the receiver operating characteristic (ROC) curve. Calibration was assessed using the Hosmer-Lemeshow goodness-of-fit test. The Hosmer-Lemeshow test compared model performance (observed vs. expected) across deciles of risk to test whether the model was biased (i.e., performed differently at the extremes of risk). A nonsignificant value for the Hosmer-Lemeshow test suggested an absence of such bias (23).

All P values were 2 tailed, and P<0.05 was considered statistically significant. All statistical analyses were conducted with SPSS 24.0 (IBM, New York, USA) and R software (version 3.4.2, http://www.r-project.org/).

## Results

# **Baseline characteristics**

Our selection criteria identified 1,548 patients with BCLC

early-stage HCC. Of these patients, 1,481 and 67 patients had SHCC and MHCC, respectively. PSM was performed to overcome the imbalances between these two groups and resulted in 126 patients with SHCC and 64 patients with MHCC. No significant differences existed between the two groups (all P>0.05). The baseline characteristics of the patients are listed in *Table 1*. The effectiveness of PSM is shown in *Table 1* and *Figure S1*. *Figure S2* shows the correlation between MVI and the clonal origin pattern of MHCC.

#### Risk factors for RFS and OS

The survival analysis of the crude cohort is shown in *Table 2*. The multivariate Cox proportional hazards model identified a high total bilirubin (TBIL), high  $\gamma$ -glutamyl transpeptidase (GGT), high INR, HBV DNA load >10<sup>3</sup> IU/mL, positive hepatitis B e antigen (HBeAg), large tumor size, multiple tumor nodules, and MVI as risk factors for RFS. A high TBIL, low albumin (ALB), high GGT, high AFP, positive HBeAg, large tumor size, presence of cirrhosis, poor tumor differentiation and MVI remained independent risk factors for poor OS. *Table S1* shows the Cox regression analysis of the PSM cohort.

## Impact of tumor number before and after PSM

Before PSM, the 1-, 3-, 5-, and 7-year RFS rates in the SHCC group were 71.3%, 52.9%, 44.3%, and 38.0% and those in the MHCC group were 65.9%, 41.2%, 29.6%, and 24.7%, respectively (Figure 1A, P=0.012). The mean RFS of the SHCC and MHCC groups were 45.71 and 35.57 months, respectively. The 1-, 3-, 5-, and 7-year OS rates of the SHCC group were 91.7%, 79.2%, 70.0%, and 63.8% and those of the MHCC group were 86.3%, 75.4%, 65.9%, and 61.1%, respectively (Figure 1B, P=0.573). The mean OS of the SHCC and MHCC groups were 65.76 and 63.06 months, respectively. After PSM, the 1-, 3-, 5-, and 7-year RFS rates of the SHCC group were 74.2%, 54.6%, 41.2% and 36.1% and those of the MHCC group were 64.2%, 39.9%, 27.8%, and 22.6%, respectively (Figure 1C, P=0.029). The mean RFS of the SHCC group was 45.00 months compared with 34.29 months of the MHCC group. The 1-, 3-, 5-, and 7-year OS rates of the SHCC group were 93.6%, 85.5%, 75.2%, and 63.2% and those of the MHCC group were 85.7%, 74.2%, 64.2%, and 60.9%, respectively (Figure 1D, P=0.493). The mean OS of the SHCC and MHCC groups were 68.74 and 62.30 months,

# Page 4 of 12

Table 1 Ba	aseline character	ristics of BCL	C early stag	e HCC	patients be	fore and after	propensit	v score matching
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		Crude cohort				PSM cohort		
Characteristics	SHCC (n=1,481)	MHCC (n=67)	P value	ASD	SHCC (n=126)	MHCC (n=64)	P value	ASD
Sex			0.054	0.275			1.000	0.030
Male	1,240 (83.7%)	62 (92.5%)			117 (92.9%)	59 (92.2%)		
Female	241 (16.3%)	5 (5.7%)			9 (7.1%)	5 (7.8%)		
Age, year	51.99±10.62	52.99±9.74	0.371	0.098	53.17±8.27	53.16±9.92	0.994	0.024
TBIL, µmol/L	16.25±23.86	15.60±7.96	0.507	0.036	15.67±8.47	15.62±8.12	0.601	0.002
TP, g/L	73.89±5.71	73.72±5.79	0.817	0.029	72.94±5.72	73.62±5.76	0.306	0.113
ALB, g/L	42.32±3.93	42.57±4.32	0.549	0.059	42.38±3.72	42.52±4.37	0.818	0.036
ALT, U/L	42.79±31.70	44.86±30.12	0.240	0.067	42.98±32.49	40.03±18.33	0.507	0.105
AST, U/L	40.83±28.14	34.06±15.15	0.185	0.300	35.49±21.19	33.40±15.12	0.790	0.109
GGT, U/L	92.41±121.41	75.01±88.77	0.410	0.164	67.76±57.17	75.75±90.77	0.899	0.103
ALP, U/L	93.21±56.84	81.48±31.36	0.039	0.256	80.13±32.08	82.11±31.90	0.747	0.060
AFP, ng/mL	371.75±499.68	351.68±458.91	0.479	0.042	305.90±482.08	362.90±465.60	0.437	0.127
CA199, ng/mL	28.72±56.97	24.55±24.68	0.977	0.095	24.28±22.13	24.93±24.98	0.750	0.039
WBC, 10 <sup>9</sup> /L	5.30±1.77	4.94±1.37	0.108	0.229	5.08±1.69	4.99±1.37	0.910	0.063
RBC, 10 <sup>9</sup> /L	4.66±0.52	4.66±0.49	0.652	0.001	4.71±0.46	4.67±0.50	0.527	0.097
PLT, 10 <sup>9</sup> /L	162.20±70.06	135.25±51.56	0.004	0.437	139.30±68.83	136.72±51.29	0.744	0.062
INR	1.00±0.08	1.02±0.10	0.058	0.269	1.02±0.09	1.02±0.10	0.987	0.050
PT, s	12.01±0.97	12.29±1.16	0.057	0.267	12.25±1.03	12.29±1.18	0.987	0.043
HBV DNA load			0.561	0.072			0.475	0.095
>10 <sup>3</sup> IU/mL	739 (49.9%)	31 (46.3%)			64 (50.8%)	29 (45.3%)		
≤10 <sup>3</sup> IU/mL	742 (50.1%)	36 (53.7%)			62 (49.2%)	35 (54.7%)		
HBsAg			0.290	0.143			0.974	0.000
Positive	1,282 (86.6%)	61 (91.0%)			114 (90.5%)	58 (90.6%)		
Negative	199 (13.4%)	6 (9.0%)			12 (9.5%)	6 (9.4%)		
HBsAb			0.761	0.037			0.921	0.022
Positive	223 (15.1%)	11 (16.4%)			19 (15.1%)	10 (15.6%)		
Negative	1,258 (84.9%)	56 (83.6%)			107 (84.9%)	54 (84.4%)		
HBeAg			0.211	0.153			0.715	0.034
Positive	405 (27.3%)	23 (34.3%)			40 (31.7%)	22 (34.4%)		
Negative	1,076 (72.7%)	44 (65.7%)			86 (68.3%)	42 (65.6%)		
HBeAb			0.037	0.247			0.555	0.068
Positive	1,137 (76.8%)	44 (65.7%)			88 (69.8%)	42 (65.6%)		
Negative	344 (23.2%)	23 (34.3%)			38 (30.2%)	22 (34.4%)		
HBcAb			0.622	0.182			1.000	0.126
Positive	1,457 (98.4%)	67 (100.0%)			125 (99.2%)	64 (100.0%)		
Negative	24 (1.6%)	0 (0.0%)			1 (0.8%)	0 (0.0%)		

Table 1 (continued)

Table 1 (continued)

Characteristics		Crude cohort				PSM cohort		
Characteristics	SHCC (n=1,481)	MHCC (n=67)	P value	ASD	SHCC (n=126)	MHCC (n=64)	P value	ASD
Child-Pugh			1.000	0.089			1.000	0.057
А	1,440 (97.2%)	66 (98.5%)			123 (97.6%)	63 (98.4%)		
В	41 (2.8%)	1 (1.5%)			3 (2.4%)	1 (1.6%)		
Pringle maneuver			0.077	0.255			0.735	0.108
Yes	1,282 (86.6%)	63 (94.0%)			121 (96.0%)	60 (93.8%)		
No	199 (13.4%)	4 (6.0%)			5 (4.0%)	4 (6.3%)		
Transfusion			0.155	0.191			0.777	0.049
Yes	279 (18.8%)	8 (11.9%)			14 (11.1%)	8 (12.5%)		
No	1,202 (81.2%)	59 (88.1%)			112 (88.9%)	56 (87.5%)		
Capsule			0.011	0.351			0.651	0.063
Complete	434 (29.3%)	10 (14.9%)			23 (18.3%)	10 (15.6%)		
Incomplete	1,047 (70.7%)	57 (85.1%)			103 (81.7%)	54 (84.4%)		
Cirrhosis			0.101	0.209			0.692	0.082
Yes	777 (52.5%)	42 (62.7%)			75 (59.5%)	40 (62.5%)		
No	704 (47.5%)	25 (37.3%)			51 (40.5%)	24 (37.5%)		
ES grade			0.723	0.045			0.769	0.054
1-11	391 (26.4%)	19 (28.4%)			31 (24.6%)	17 (26.6%)		
III-IV	1,090 (73.6%)	48 (71.6%)			95 (75.4%)	47 (73.4%)		
MVI			0.938	0.009			0.831	0.016
Yes	656 (44.3%)	30 (44.8%)			57 (45.2%)	30 (46.9%)		
No	825 (55.7%)	37 (55.2%)			69 (54.8%)	34 (53.1%)		
Diameter, cm	5.37±3.68	2.40±0.46	<0.001	1.131	4.33±2.88	2.41±0.47	<0.001	0.932
Tumor number			NA				NA	
1	1,481 (100.0%)	0 (0.0%)			126 (100.0%)	0 (0.0%)		
2	0 (0.0%)	62 (92.5%)			0 (0.0%)	60 (93.8%)		
3	0 (0.0%)	5 (7.5%)			0 (0.0%)	4 (6.3%)		
Location			NA				NA	
Same lobe	0 (0.0%)	50 (74.6%)			0 (0.0%)	47 (73.4%)		
Different lobe	0 (0.0%)	17 (25.4%)			0 (0.0%)	17 (26.6%)		
Very early stage			NA				NA	
Yes	170 (11.5%)	0 (0.0%)			17 (13.5%)	0 (0.0%)		
No	1,311 (88.5%)	0 (0.0%)			109 (86.5%)	0 (0.0%)		

TBIL, total bilirubin; TP, total protein; ALB, albumin; ALT, alanine transaminase; AST, aspartate aminotransferase; GGT, γ-glutamyl transpeptidase; ALP, alkaline phosphatase; AFP, alpha fetal protein; CA199, carbohydrate antigen 19-9; WBC, white blood cells; RBC, red blood cells; PLT, platelets; INR, international normalized ratio; PT, prothrombin time; HBsAg, hepatitis B surface antigen; HBsAb, hepatitis B surface antibody; HBeAg, hepatitis B e antigen; HBeAb, hepatitis B e antibody; HBcAb, hepatitis B core antibody; ES, Edmondson-Steiner; MVI, microvascular invasion; NA, no answer; ASD, absolute standard difference.

		Recurrence-t	free survival			Overall	lsurvival	
Characteristics		Univariate		Multivariate		Univariate		Multivariate
	P value	HR (95% CI)	P value	HR (95% CI)	P value	HR (95% CI)	P value	HR (95% CI)
Sex, male	0.005	1.306 (1.086, 1.570)						
TBIL, µmol/L	<0.001	1.005 (1.003, 1.007)	0.001	1.004 (1.002, 1.006)	<0.001	1.005 (1.004, 1.007)	<0.001	1.004 (1.002, 1.006)
TP, g/L	0.007	0.984 (0.973, 0.996)			0.043	0.984 (0.970, 1.000)		
ALB, g/L	<0.001	0.945 (0.929, 0.961)			<0.001	0.914 (0.894, 0.935)	<0.001	0.946 (0.925, 0.968)
ALT, U/L	<0.001	1.004 (1.002, 1.006)			<0.001	1.004 (1.002, 1.006)		
AST, U/L	<0.001	1.007 (1.005, 1.009)			<0.001	1.009 (1.007, 1.011)		
GGT, U/L	<0.001	1.001 (1.001, 1.002)	0.003	1.001 (1.000, 1.001)	<0.001	1.002 (1.001, 1.002)	<0.001	1.001 (1.000, 1.001)
ALP, U/L	<0.001	1.003 (1.002, 1.003)			<0.001	1.004 (1.003, 1.005)		
AFP, ng/mL	<0.001	1.000 (1.000, 1.000)			<0.001	1.000 (1.000, 1.001)	0.002	1.000 (1.000, 1.000)
CA199, ng/mL	<0.001	1.002 (1.001, 1.003)			<0.001	1.003 (1.002, 1.004)		
RBC, 10 <sup>9</sup> /L					<0.001	0.713 (0.603, 0.841)		
INR	<0.001	10.816 (4.954, 23.617)	<0.001	8.375 (3.718, 18.864)	<0.001	9.145 (3.317, 25.216)		
PT, s	<0.001	1.217 (1.141, 1.299)			<0.001	1.206 (1.108, 1.312)		
HBV DNA load, >10 <sup>3</sup> IU/mL	<0.001	1.489 (1.308, 1.695)	<0.001	1.291 (1.120, 1.487)	<0.001	1.396 (1.176, 1.656)		
HBsAg, positive	0.037	1.236 (1.013, 1.508)						
HBeAg, positive	<0.001	1.380 (1.203, 1.583)	0.001	1.280 (1.100, 1.488)	0.016	1.251 (1.042, 1.501)	0.004	1.317 (1.090, 1.591)
Child-Pugh, B	<0.001	1.967 (1.382, 2.800)			<0.001	2.775 (1.884, 4.089)		
Transfusion, yes	<0.001	1.619 (1.385, 1.894)			<0.001	2.038 (1.684, 2.465)		
Diameter, cm	<0.001	1.074 (1.057, 1.092)	<0.001	1.078 (1.060, 1.097)	<0.001	1.121 (1.101, 1.141)	<0.001	1.102 (1.079, 1.125)
Number, multiple	0.012	1.454 (1.084, 1.950)	<0.001	1.754 (1.300, 2.365)				
Capsule, incomplete					<0.001	1.446 (1.184, 1.767)		
Cirrhosis, yes	0.001	1.248 (1.097, 1.421)			0.016	1.234 (1.040, 1.465)	<0.001	1.401 (1.165, 1.685)
ES grade, III-IV	0.001	1.297 (1.118, 1.505)			<0.001	1.932 (1.547, 2.411)	0.046	1.273 (1.004, 1.614)
MVI, yes	<0.001	1.478 (1.300, 1.681)	<0.001	1.396 (1.226, 1.591)	<0.001	1.871 (1.577, 2.220)	<0.001	1.577 (1.317, 1.888)
TBIL, total bilirubin; TP, tota AFP, alpha fetal protein; C <sup>A</sup>	l protein, AL \199, carbo	LB albumin; ALT, alanine tr hydrate antigen 19-9; RB	ransaminase 3C. red bloo	e; AST, aspartate aminoti	ransferase; al normaliz	GGT, γ-glutamyl transpe	ptidase; ALF	) alkaline phosphatase; An henatitis R surface

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Figure 1 Cumulative incidence of recurrence-free survival and overall survival curves showing a comparison between SHCC patients and MHCC patients before (A,B) and after (C,D) propensity score matching. SHCC, solitary hepatocellular carcinoma; MHCC, multinodular hepatocellular carcinoma.

respectively.

## Subgroup analyses based on tumor number and MVI

For all patients selected for the PSM cohort, the mean RFS and OS of the SHCC group, the MVI-negative MHCC group, and the MVI-positive MHCC group were 45.00, 45.46, and 21.70 months as well as 68.74, 71.77, and 52.05 months, respectively. The 1-, 3-, 5-, and 7-year RFS rates of the SHCC group were 74.2%, 54.6%, 41.2%, and 36.1%, those of the MVI-negative MHCC group were 68.5%, 52.2%, 45.7%, and 39.1%, and those of the MVI-positive MHCC group were 59.6%, 26.1%, 7.5%, and 3.7%, respectively. The 1-, 3-, 5-, and 7-year OS rates of the SHCC group were 94.4%, 85.5%, 75.2%, and 63.2%, those of the MVI-negative MHCC group were 94.1%, 87.8%, 78.3%, and 75.0%, and those of the MVI-positive MHCC group were 76.7%, 59.4%, 48.9%, and 45.4%, respectively. The MVI-positive MHCC group had a worse RFS and OS than

the SHCC group (RFS P<0.001, *Figure 2A*; OS P=0.017, *Figure 2B*). Nevertheless, the RFS and OS were not significantly different between the MVI-negative MHCC group and the SHCC group (RFS P=0.917, *Figure 2A*; OS P=0.272, *Figure 2B*). The predictors of MVI in the MHCC group are shown in *Table S2*.

# Risk factors for early and late recurrence

The independent risk factors for early tumor recurrence (<2 years) were analyzed among all 190 patients in the PSM cohort, while the factors associated with late recurrence were assessed among the 104 patients who had a postoperative period of at least 2 years and did not have early recurrence (*Table 3*). The multivariate analysis demonstrated that multinodular tumors [P=0.029, hazard ratio (HR) 1.655; 95% confidence interval (CI), 1.053–2.602] and MVI (P=0.029, HR 1.646; 95% CI, 1.053–2.572) were associated with early recurrence. A high ALT (P=0.045,

#### Page 8 of 12

## Wang et al. MVI for BCLC stage A MHCC



Figure 2 Cumulative incidences of recurrence-free survival (A) and overall survival (B) curves showing comparisons between SHCC patients and MHCC patients with/without MVI after propensity score matching. SHCC, solitary hepatocellular carcinoma; MHCC, multinodular hepatocellular carcinoma; MVI, microvascular invasion.

Table 3 Risk facto	ors of earl	y and late tumor recurr	ence						
		Early tumor recu	urrence (	n=190)	Late tumor recurrence (n=104)				
Characteristics		Univariate		Multivariate		Univariate		Multivariate	
	P value	HR (95% CI)	P value	HR (95% CI)	P value	HR (95% CI)	P Value	HR (95% CI)	
ALT, U/L					0.043	1.012 (1.000, 1.024)	0.045	1.012 (1.000, 1.024)	
AST, U/L					0.001	1.028 (1.012, 1.045)			
GGT, U/L	0.007	1.003 (1.001, 1.006)			0.022	1.007 (1.001, 1.012)			
ALP, U/L	0.001	1.008 (1.003, 1.013)							
AFP, ng/mL	<0.001	1.001 (1.000, 1.001)							
PLT, 10 <sup>9</sup> /L	0.094	1.003 (1.000, 1.006)							
Capsule, incomplete	0.015	2.613 (1.201, 5.684)							
Diameter, cm					0.027	1.130 (1.014, 1.258)	0.027	1.132 (1.014, 1.264)	
Tumor number, multiple	0.030	1.651 (1.051, 2.595)	0.029	1.655 (1.053, 2.602)					
ES grade, III-IV	0.021	2.016 (1.111, 3.659)							
MVI, yes	0.030	1.642 (1.050, 2.566)	0.029	1.646 (1.053, 2.572)	0.028	1.914 (1.073, 3.414)			

ALT, alanine transaminase; AST, aspartate aminotransferase; GGT, γ-glutamyl transpeptidase; ALP, alkaline phosphatase; AFP, alpha fetal protein; PLT, platelets; ES, Edmondson-Steiner; MVI, microvascular invasion.

HR 1.012; 95% CI, 1.000–1.024) and large tumor size (P=0.027, HR 1.132; 95% CI, 1.014–1.264) were significant risk factors for late recurrence. For SHCC patients, the 6-month, 1-, and 2-year RFS rates were 84.7%, 74.2%, and 62.8%, respectively; for MHCC patients, the 6-month, 1-, and 2-year RFS rates were 71.0%, 64.2%, and 46.8%, respectively (P=0.028) (*Figure 3*). The recurrence pattern of MHCC is shown in *Table S3*.

#### **Discussion**

Initially, when Llovet *et al.* proposed the BCLC staging classification, stage A4 (early stage) disease was defined as MHCC with up to 3 nodules smaller than 3 cm, and other MHCCs were classified as stage B (intermediate stage) (5).



Figure 3 Cumulative incidence of early recurrence-free survival curve showing a comparison between SHCC patients and MHCC patients after propensity score matching. SHCC, solitary hepatocellular carcinoma; MHCC, multinodular hepatocellular carcinoma.

The authors believed that stage A4 HCC patients could achieve a beneficial prognosis through curative therapy, which was mainly referred to as the Milan criteria (9). However, the study that presented the Milan criteria included only 23 patients with MHCC, and all of the patients had unresectable MHCC; hence, the conclusion of the study might be biased. Furthermore, substantial studies have detected that multiple tumor nodules are important risk factors for early recurrence. For instance, Li et al. demonstrated that HCC patients classified with the Milan criteria achieved a poorer disease-free survival with an increased tumor number (24). Li et al. illustrated that multiple tumors were associated with early recurrence for patients who underwent R0 resection (25). These results compelled us to reappraise whether all BCLC early-stage MHCC patients were suitable for curative therapy. Namely, we considered that defining early-stage MHCC by only size and number of tumors was not rigorous. Combined with individualized and intensification treatments, the parameters that reflect the biological behaviors of tumor clearly offer the strongest evidence. Therefore, histopathological features have been the most valuable basis for retrospective explorations of the misjudgments of the BCLC staging classification.

One of the significant discoveries in the molecular pathology of HCC is the clonal origin pattern of MHCC (26). Two major clonal origin patterns of MHCC have been suggested; one model is the monoclonal origin of IM-type MHCC, and the other is the polyclonal origin of MO-type MHCC (27,28). Among cases of MHCC, in cases where the tumor factors indicated high malignancy, the original tumor was hypothesized to lead to IM-type HCC, whereas MO-type HCC would be produced in other portions of the liver, such



Figure 4 Schematic diagram of reappraisal for the BCLC early-stage patients. BCLC, Barcelona Clinic Liver Cancer.

as in environments with poor background liver factors (29). Remarkably, MVI is a pathological phenomenon highly suggestive of early recurrence and unfavorable prognoses of HCC, and MVI is also a pathological factor highly correlated with IM of HCC (30). In our previous study of the clonal origin analysis of 40 recurrent HCCs, IMtype HCC had a higher frequency of vascular invasion than MO-type HCC (13). Kim et al. analyzed 198 MHCC patients and reported that MVI was the foremost factor for discriminating between the IM-type group and the MOtype group (18). Interestingly, our study also suggested a close correlation between MVI and IM-type MHCC. Since no consensus exists on the technology and criteria for determining the clonal origin of HCC, we believe that MVI represents malignant biological behavior and is a practical indicator for identifying IM-type MHCC.

The first observation of this study with a large cohort of patients with BCLC early-stage HCC was that the MHCC group had a higher recurrence rate than the SHCC group both before and after PSM. Furthermore, the presence of multiple tumor nodules was an independent risk factor for early recurrence but not late recurrence. These results prompted us to determine the possibility of earlystage MHCC forming IM and having MVI. Therefore, a subgroup analysis was conducted to explore the impact of MVI on the prognosis of MHCC. The Kaplan-Meier analysis implied that only MHCC patients without MVI had similar clinical outcomes to SHCC patients, while MHCC patients with MVI were associated with a decreased clinical outcome compared with SHCC patients. Considering the 44% detection rate of MVI, we believe that patients with BCLC early-stage HCC and MVI should not be classified as the same stage as those with SHCC and MHCC without MVI. The underlying cause is that MHCC with MVI is probably formed by IM, so the biological behavior of these tumors is highly malignant and therefore, these tumors are unable to benefit from curative treatment (Figure 4).

Current technology still cannot accurately predict MVI preoperatively, which is an inevitable limitation of our study. Notably, some research has proposed some efficient prediction models for MVI (31-33). Thus, we also created a prediction model for MVI based on our data to provide a reference for treatment selection for BCLC early-stage MHCC patients. More accurate prediction models for MVI for these patients are anticipated in the future.

To the best of our knowledge, our study was the first to analyze the impact of tumor number on BCLC earlystage patients and to suggest the role of MVI in MHCC. Moreover, we used PSM to balance the baseline data of the included patients, thus avoiding interference bias from other factors on the prognosis (21). Furthermore, the detection rate of MVI in our study was more authentic than that in other studies because a 7-point baseline sampling protocol was performed for all specimens. Tissue specimens sampled at the junction of the tumor and the adjacent liver tissues can typically reflect the parameters that indicate biological tumor behaviors such as MVI, capsule, and grade.

Some limitations exist in our study; first, our study was a single-center study. Considering the different treatment experiences of different hospitals, multicenter research is still essential. Second, the number of BCLC early-stage MHCC patients in our study was relatively small, and although PSM could overcome bias to a certain extent, our conclusions still need to be verified in a larger cohort. In addition, liver transplantation is also a selective treatment option for BCLC early-stage MHCC, but these patients were valuable to the analyses of our study.

## Conclusions

The presence of MVI in BCLC early-stage MHCC was highly suggestive of poor postoperative outcomes, which is possibly due to the malignant biological behavior of tumors caused by IM. The BCLC staging system should reappraise MHCC based on parameters that reflect the presence of MVI and distinguished the biological behavior level of early-stage MHCC.

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# Footnote

*Conflicts of Interest:* The authors have no conflicts of interest to declare.

*Ethical Statement:* The authors are accountable for all aspects of the work in ensuring that questions related

to the accuracy or integrity of any part of the work are appropriately investigated and resolved. This study was conducted in accordance with the Declaration of Helsinki and approved by the Clinical Research Ethics Committee of the Eastern Hepatobiliary Surgery Hospital of Shanghai, China (No. EHBHKY2015-02-001). Informed consent was obtained from all patients for their dates to be used for research.

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## Wang et al. MVI for BCLC stage A MHCC

## Page 12 of 12

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# Supplementary





Figure S2 Correlation between microvascular invasion and clonal origin in early-stage MHCC patients. MHCC, multinodular hepatocellular carcinoma.

Figure S1	Evaluation	of	effectiveness	about	propensity	score
matching.						

## Table S1 Cox regression analysis of the propensity score matching cohort

		Recurrence-	free surviva	al	Overall survival			
Characteristics		Univariate		Multivariate		Univariate		Multivariate
	P value	HR (95% CI)	P value	HR (95% CI)	P value	HR (95% CI)	P value	HR (95% CI)
Sex, male	0.037	2.593 (1.059,6.350)			0.087	3.417 (0.837, 13.956)		
ALB, g/L					0.031	0.934 (0.878, 0.994)	0.009	0.916 (0.858, 0.978)
ALT, U/L	0.019	1.007 (1.001,1.013)						
AST, U/L	0.025	1.008 (1.001,1.016)			0.001	1.012 (1.005, 1.020)	0.004	1.012 (1.004, 1.020)
GGT, U/L	0.001	1.004 (1.002,1.006)			0.045	1.003 (1.000, 1.005)		
ALP, U/L	0.005	1.007 (1.002,1.012)			0.019	1.006 (1.001, 1.011)		
AFP, ng/mL	0.012	1.000 (1.000,1.001)			<0.001	1.001 (1.000, 1.001)	0.029	1.001 (1.000, 1.001)
INR					0.033	18.805 (1.258, 281.059)		
PT, s					0.03	1.283 (1.024, 1.608)		
HBV DNA load, >10 <sup>3</sup> IU/mL	0.080	1.371 (0.963,1.952)	0.024	1.520 (1.057, 2.187)				
Transfusion, yes					0.015	2.117 (1.156, 3.875)		
Diameter, cm	0.018	1.077 (1.013, 1.145)	<0.001	1.135 (1.065, 1.209)	0.003	1.115 (1.039, 1.198)		
Tumor number, multiple	0.030	1.498 (1.040, 2.158)	0.001	2.003 (1.345, 2.984)				
ES grade, III-IV	0.038	1.572 (1.026, 2.409)						
MVI, yes	0.002	1.739 (1.220, 2.479)	0.001	1.787 (1.250, 2.553)	<0.001	2.560 (1.556, 4.212)	<0.001	2.612 (1.531, 4.457)

ALB, albumin; ALT, alanine transaminase; AST, aspartate aminotransferase; GGT, γ-glutamyl transpeptidase; ALP, alkaline phosphatase; AFP, alpha fetal protein; INR, international normalized ratio; PT, prothrombin time; ES, Edmondson-Steiner; MVI, microvascular invasion.

# Table S2 Predictors of MVI for the early-stage MHCC

Characteristics		Univariate		Multivariate
Characteristics	P value	OR (95% CI)	P value	OR (95% CI)
TBIL, μmol/L	0.085	1.074 (0.990, 1.164)		
HBsAb, negative	0.028	10.741 (1.287, 89.608)	0.010	21.506 (2.073, 223.126)
Capsule, no	0.105	3.862 (0.754, 19.794)	0.030	7.182 (1.206, 42.756)
AFP, ng/mL	0.044	1.001 (1.000, 1.002)	0.024	1.002 (1.000, 1.003)

MHCC, multinodular hepatocellular carcinoma; MVI, microvascular invasion; MHCC, multinodular hepatocellular carcinoma; TBIL, total bilirubin; HBsAb, hepatitis B surface antibody; AFP, alpha fetal protein.

Table S3 Recurrence pattern of the early-stage MHCC

Characteristics	MVI negative	MVI positive	P value
Recurrence rate	20 (54.1%)	27 (90.0%)	0.001
Recurrence <2 year	14 (70.0%)	19 (70.4%)	0.978
Intrahepatic tumor number			1.000
No	1 (5.0%)	1 (3.7%)	
Single	13 (65.0%)	18 (66.7%)	
Multiple	6 (30.0%)	8 (29.6%)	
Extrahepatic recurrence			1.000
Yes	1 (5.0%)	2 (7.4%)	
No	19 (95.0%)	25 (92.6%)	
Recurrent AFP, ng/mL	82.01±203.14	182.14±314.62	0.554
Treatment			0.847
Re-resection	2 (10.0%)	3 (11.1%)	
Ablation	8 (40.0%)	8 (29.6%)	
TACE	8 (40.0%)	14 (51.9%)	
Others	2 (10.0%)	2 (7.4%)	

MHCC, multinodular hepatocellular carcinoma; AFP, alpha fetal protein; TACE, transcatheter arterial chemoembolization.