

# A Novel Nomogram Model to Predict the Recurrence-Free Survival and Overall Survival of Hepatocellular Carcinoma

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**Background:** Treatments for patients with early-stage hepatocellular carcinoma (HCC) include liver transplantation (LT), liver resection (LR), radiofrequency ablation (RFA), and microwave ablation (MWA), are critical for their long-term survival. However, a computational model predicting treatment-independent prognosis of patients with HCC, such as overall survival (OS) and recurrence-free survival (RFS), is yet to be developed, to our best knowledge. The goal of this study is to identify prognostic factors associated with OS and RFS in patients with HCC and develop nomograms to predict them, respectively.

**Methods:** We retrospectively retrieved 730 patients with HCC from three hospitals in China and followed them up for 3 and 5 years after invasive treatment. All enrolled patients were randomly divided into the training cohort and the validation cohort with a 7:3 ratio, respectively. Independent prognostic factors associated with OS and RFS were determined by the multivariate Cox regression analysis. Two nomogram prognostic models were built and evaluated by concordance index (C-index), calibration curves, area under the receiver operating characteristics (ROC) curve, time-dependent area under the ROC curve (AUC), the Kaplan–Meier survival curve, and decision curve analyses (DCAs), respectively.

**Results:** Prognostic factors for OS and RFS were identified, and nomograms were successfully built. Calibration discrimination was good for both the OS and RFS nomogram prediction models (C-index: 0.750 and 0.746, respectively). For both

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nomograms, the AUC demonstrated outstanding predictive performance; the DCA shows that the model has good decision ability; and the calibration curve demonstrated strong predictive power. The nomograms successfully discriminated high-risk and low-risk patients with HCC associated with OS and RFS.

**Conclusions:** We developed nomogram survival prediction models to predict the prognosis of HCC after invasive treatment with acceptable accuracies in both training and independent testing cohorts. The models may have clinical values in guiding the selection of clinical treatment strategies.

Keywords: hepatocellular carcinoma, Prognosis, nomogram, OS, RFS

# INTRODUCTION

Liver cancer is one of the most prevalent and aggressive tumors, as well as the third leading cause of cancer-related mortality, with roughly 906,000 new cases and 83,0000 deaths reported in 2020 (1). Hepatocellular carcinoma (HCC) is the most common primary liver cancer comprising 75%–85% of all liver cancers. At present, there is no effective method for the treatment of advanced liver cancer, so the early treatment of liver cancer is very important for the prognosis of patients (2).

Early-stage HCC has been defined as Barcelona Clinic Liver Cancer (BCLC) 0 and A stages (BCLC 0/A), with curative therapy being the primary therapeutic option (2, 3). The treatment of early-stage HCC is feasible, but most patients with intermediate or advanced liver cancer have limited treatment opportunities and receive palliative treatment. The main curative treatments for early-stage HCC are liver resection (LR) and liver transplantation (LT). Radiofrequency ablation (RFA) or microwave ablation (MWA) is a viable minimally invasive therapy option for very early and early HCC, with equivalent outcomes to surgical resection (4–7). The high postoperative recurrence rate of liver cancer is the main obstacle affecting the low survival rate of patients with liver cancer (8–10).

There are several staging and grading systems for HCC, most notably the BCLC classification and the AJCC/TNM eight edition (11, 12). Several other factors were reported as potential predictor for the postoperative outcomes of patients with HCC, such as aspartate aminotransferase-to-platelet ratio index, albumin-bilirubin score (ALBI), the Model for Endstage Liver Disease (MELD) score and the Child-Pugh score (11, 13, 14). In addition, there are a few machine learning models for predicting the prognosis of cancer patients for other types of cancers based on histopathological images and multi-omics data (15-18). However, to our best knowledge, an accurate model predicting treatment-independent prognosis of patients with HCC, such as overall survival (OS) and recurrence-free survival (RFS), is yet to be developed. Although the treatment technology of HCC has made progress, the OS and RFS of patients with HCC are still relatively low. At present, there is still an urgent need for accurate models to predict OS and RFS in patients with HCC, guide individualized treatment, and prolong survival.

Nomogram is a user-friendly graphical prediction model tool that can help with therapeutic decision-making by quantifying the impacts of a variety of parameters (19). Therefore, we aimed to identify prognostic factors associated with OS and RFS in patients with HCC with different invasive treatments, including LT, LR, and minimally invasive approach (RFA or MWA) and develop nomograms to estimate 3-year and 5-year OS and RFS, respectively.

# MATERIALS AND METHODS

## **Study Population**

We retrospectively retrieved 730 patients with HCC underwent LT, LR, and RFA or MWA in three Chinese medical centers (Tianjin Medical University Cancer Institute and Hospital, Tianjin, China; The First Central Clinical School, Tianjin Medical University, Tianjin, China; and Clinical School of the Second People's Hospital, Tianjin Medical University, Tianjin, China), and the follow-up deadline was on May 2019.

The inclusion criteria were as follows: (1) patients were validated by pathological diagnosis with primary HCC and assessed at BCLC 0/A; (2) patients underwent LT, LR, and RFA or MWA; (3) patients with complete clinic-pathological follow-up data; and (4) distant metastasis was not found.

## **Data Collection and Follow-Ups**

In our study, we collected the following clinical data from patients with HCC: (1) demographic characters, including age, gender, body mass index (BMI), and cirrhosis; (2) tumor size (largest tumor diameter), number, and location were estimated using magnetic resonance imaging (MRI) and/or computed tomography (CT) before treatment; (3) curative options, including LT, LR, and RFA or MWA; (4) microvascular invasion and differentiation grade were assessed postoperatively by postoperative pathology; and (5) BCLC classification was used to identify tumor stage. BMI was calculated using the following formula: BMI = weight (kg)/ height (m<sup>2</sup>).

OS was defined as the time from the date of surgery to the date of death, and RFS was defined as the time from surgery to the date of first recurrence. All data were obtained from the first

laboratory examination after admission. hepatitis C virus (HCV) and/or hepatitis B virus (HBV) infection as the presence of absence of anti-HCV or HBV surface antigen, respectively. Laboratory tests included routine blood tests, liver function tests, and alpha fetoprotein (AFP). Subgroup analysis will be performed based on Laboratory tests as follows: gammaglutamyl transpeptidase (GGT) (<45 versus  $\geq$ 45, U/L), albumin (ALB) (<35 versus  $\geq$ 35, g/L), prothrombin time (PT)  $(\leq 13 \text{ versus} > 13, \text{ s})$ , aspartate aminotransferase (AST)  $(\leq 40$ versus>40, U/L), total bilirubin (TBIL) (<20 versus ≥20, µmol/ L), and AFP (<400 versus  $\geq$ 400, ng/ml). MELD score grade, Child-Pugh Classification, and ALBI grade were also recorded. MELD score has been proved to be a predictor of survival in different end-stage liver diseases (20). Liver function was evaluated using Child-Pugh classification system. ALBI grade I:  $\leq$ -2.60 score; ALBI grade II: >-2.60 to  $\leq$ -1.39 score; ALBI grade III: >-1.39 score.

## **Statistical Analysis**

HCV

Statistical analysis was performed by R software (R Statistical Software, version 4.1.2). Independent prognostic factors were identified using multivariate Cox regression analyses. The results are presented as hazard ratio (HR) with 95% confidence intervals (CIs). Nomogram and calibration plots were constructed using R software. The C-index, ROC curve, calibration curve, and DCA were used to assess the nomogram. On the basis of the nomogram risk scores, Kaplan-Meier (K-M) survival curves were plotted for patients in the high-risk and low-risk groups.

**TABLE 1** | Demographic and clinical characteristics of patients.

# RESULTS

## Patients' Demographics and Clinical Characteristics

As shown in Table 1, a total of 730 patients diagnosed with primary HCC were included in our research. After using R software, patients were randomly allocated in a 7:3 ratio between the training cohort (512 patients) and the validation cohort (218 patients). In the OS analysis, the median follow-up period of the entire study cohort was 56.9 months (interquartile range, 2.8-116.3 months). In the RFS analysis, the median follow-up period for the overall research population was 41.3 months (interquartile range, 1.7-116.3 months). The training cohort was used to build the nomogram and internally validate the model, whereas the validation cohort was utilized for external verification. Both the training and the validation cohort had no statistical differences in their baseline characteristics in OS or RFS groups (Table 1).

In the entire cohort, 68.8% were aged < 60 years, 18.7% of the population were women, 58.8% were BMI < 25, and 87.8% of patients had cirrhosis. For all evaluated tumors, CT, MRI, or pathological examination results were available. Tumor size was defined as the largest diameter and 69.9% were less than 3 cm. There were 576 cases with a single tumor and 154 cases with multiple tumors. According to tumor location, 127 cases were in the left lobe, 571 cases were in the right lobe, and 32 cases were in both lobes. Pathological examination revealed microvascular invasion in 13.3% of all patients, whereas 55.3% were not. Child-Pugh grade and ALBI grade were used for assessment of

Characters		Total patients (n = 730)	OS			RFS	
			Training Cohort (n = 512)	Validation Cohort (n = 218)	P- value	Training Cohort (n = 512)	Validation Cohort (n = 218)
Age (year)	<60/≥60	502/228	349/163	153/65	0.590	352/160	150/68
Gender	Female/Male	137/593	103/409	34/184	0.153	89/432	48/170
BMI (kg/m <sup>2</sup> )	<25/≥25	429/301	293/219	136/82	0.195	299/213	130/88
Cirrhosis	No/Yes	89/641	66/446	23/195	0.377	67/445	22/196
Tumor location	Left Lobe/Right Lobe/Both Lobe	127/571/32	90/396/26	37/175/6	0.690	84/405/23	43/166/9
Tumor Number	Solitary/Multiple	576/154	399/113	177/41	0.323	405/107	171/47
Tumor Size (cm)	≤3/3 < R ≤ 5	510/220	361/151	149/69	0.561	362/50	148/70
Operation	LT/LR/RFA or MWA	252/249/229	185/170/157	67/79/72	0.233	178/164/170	74/85/59
MELD score grade	<9/9-15/>15	459/218/53	319/154/39	140/64/14	0.577	319/156/37	140/62/16
Child-Pugh	A/B	553/177	387/125	166/52	0.872	382/130	171/47
Classification							
ALBI grade	1/11/111	344/344/42	236/249/27	108/95/15	0.564	240/240/32	104/104/10
Laboratory examina	ation						
GGT (U/L)	<45/≥45	315/415	221/291	94/124	0.991	221/291	94/124
ALB (g/L)	<35/≥35	194/536	146/366	48/170	0.069	136/376	58/160
PT (S)	≤13/>13	347/383	243/269	104/114	0.952	246/266	101/117
AST (U/L)	≤40/>40	444/286	301/211	143/75	0.085	309/203	135/83
TBIL (µmol/L)	<20/≥20	412/318	267/225	125/93	0.749	288/224	124/94
AFP (ng/ml)	<400/≥400	632/98	443/69	189/29	0.950	440/72	192/26
HBV	No/Yes	136/594	100/412	36/182	0.338	95/417	41/177

670/60

BMI, body mass index; MELD, model for end-stage liver disease; ALBI, albumin-bilirubin; GGT, gamma-glutamyl transpeptidase; ALB, albumin; PT, prothrombin time; AST, aspartate aminotransferase: TBIL, total bilirubin: AFP, alpha fetoprotein: HBV, hepatitis B virus; HCV, hepatitis C virus,

204/14

0.249

468/44

466/46

No/Yes

202/16

P-

value

0.988

0.142

0.757

0.258

0.301

0.842

0 4 4 9

0.433

0.663

0.270

0.689

0.991 0.991

0.671 0.690 0.875

0.439

0.936

0.573

hepatic function, and MELD score wasused to assess disease severity. After evaluation, most patients with liver cancer have good liver function.

The other components included GGT < 45 U/L (43.2% versus 56.8%), ALB < 35 g/L (26.6% versus 73.4%), PT  $\leq$  13 s (47.5% versus 52.5%), AST $\leq$  40 U/L (60.8% versus 39.2%), TBIL < 20 µmol/L (56.4% versus 43.6%), AFP < 400 ng/ml (86.6% versus 13.4%), HCV negative (18.6% versus 81.4%), and HBV negative (91.8% versus 8.2%). In the BCLC classification, grades 0/A accounted for 15.6% and 84.4%, respectively.

# Identification Prognostic Risk Factors For OS and RFS in 512 Patients With HCC

Multivariate analysis was conducted to identify the prognostic risk factors for OS and RFS. The results are shown in **Table 2**. In the OS analysis, gender (HR: 1.680; 95% CI: 1.002, 2.817; P = 0.049), BMI (HR: 1.621; 95% CI: 1.131, 2.324; P = 0.009), tumor number (HR: 2.165; 95% CI: 1.323, 3.544; P = 0.002), tumor size (HR: 2.180; 95% CI: 1.412, 3.391; P < 0.001), and operation (LR: HR: 3.905, 95% CI: 2.068, 7.372; P < 0.001), and operation (LR: HR: 3.905, 95% CI: 3.906, 13.033; P < 0.001) were statistically significant differences. In the RFS analysis, tumor number (HR: 1.829; 95% CI: 1.223, 2.736; P = 0.003), operation (LR: HR: 6.019; 95% CI: 3.588, 10.098; P < 0.001; RFA or MWA: HR: 12.089; 95%

CI: 7.417, 19.703; P < 0.001), GGT (HR: 1.650; 95% CI 1.199, 2.269; P = 0.002), and HCV (HR: 0.430; 95% CI: 0.213, 0.870; P = 0.019) were considered statistically different.

# Nomogram for OS and RFS Construction and Performance Evaluation

The identification prognostic risk factors of OS and RFS were included to create prognostic nomograms to assess the 3-year and 5-year OS and RFS of patients with HCC (Figure 1). Nomograms predicted 3-year and 5-year OS and RFS indicated that operation factors had major impacts on patient prognosis. The 3-year and 5-year AUCs for OS were 0.757 and 0.795, respectively, and were 0.788 and 0.801 for RFS, respectively (Figures 2A, B, E, F). In the OS testing cohort, the AUCs of 3-year and 5-year AUCs were 0.686 and 0.774, respectively (Figures 2C, D). In the RFS testing cohort, the AUCs of 3year and 5-year AUCs were 0.768 and 0.786, respectively (Figures 2G, H). On the 3-year and 5-year calibration plots of OS and RFS, calibration curves revealed the consistency of the nomogram between predicted and actual observed and showed that the nomograms were highly consistent in both training and validation cohorts (Figure 3). In addition, the DCA curves revealed that the nomogram had a high prediction efficiency for CSS of patients with HCC in both OS and RFS (Figure 4).

TABLE 2 | Multivariate analyses for OS and RFS in patients with 512 HCC (training cohort).

$\begin{tabular}{ c c c c c c c c c c c c c c c c c c c$
Age (year) <60/260
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$
BMI (kg/m <sup>2</sup> ) <25/≥25 1.621 (1.131–2.324) 0.009* 1.272 (0.959–1.686) 0.06   Cirrhosis No/Yes 0.705 (0.393–1.265) 0.241 0.954 (0.613–1.483) 0.83   Tumor location Left Lobe Reference Reference Reference Reference   Tumor location Left Lobe Reference Reference Reference Reference   Tumor location Left Lobe Reference Reference Reference Reference   Tumor Number Solitary/Multiple 2.165 (1.323–3.544) 0.002* 1.829 (1.223–2.736) 0.000   Tumor Size (cm) Solitary/Multiple 2.165 (1.412–3.391) <0.001* 1.370 (0.984–1.907) 0.000   Operation LT Reference Reference Reference Reference 0.001* 1.370 (0.984–1.907) 0.000   MELD score grade Sq Sq.905 (2.068–7.372) <0.001* 1.370 (0.984–1.907) <0.001   MELD score grade Sq Sq.905 (2.068–7.372) <0.001* 12.089 (7.417–19.703) <0.000   MELD score gr
Cirrhosis No/Yes 0.705 (0.393–1.265) 0.241 0.954 (0.613–1.483) 0.85   Tumor location Left Lobe Reference
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$
Right Lobe 0.967 (0.598–1.565) 0.893 0.859 (0.588–1.253) 0.44   Both Lobe 0.750 (0.243–2.315) 0.617 1.271 (0.568–2.845) 0.555   Tumor Number Solitary/Multiple 2.165 (1.323–3.544) 0.002* 1.829 (1.223–2.736) 0.000   Tumor Size (cm) ≤3/3 < R ≤ 5
Both Lobe 0.750 (0.243–2.315) 0.617 1.271 (0.568–2.845) 0.555   Tumor Number Solitary/Multiple 2.165 (1.323–3.544) 0.002* 1.829 (1.223–2.736) 0.000   Tumor Size (cm) ≤3/3 < R ≤ 5
Tumor Number Solitary/Multiple 2.165 (1.323–3.544) 0.002* 1.829 (1.223–2.736) 0.00   Tumor Size (cm) ≤3/3 < R ≤ 5
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$
Operation LT Reference Reference   LR 3.905 (2.068–7.372) <0.001*
LR 3.905 (2.068–7.372) <0.001* 6.019 (3.588–10.098) <0.00   RFA or MWA 7.135 (3.906–13.033) <0.001*
RFA or MWA 7.135 (3.906–13.033) <0.001* 12.089 (7.417–19.703) <0.00   MELD score grade <9
MELD score grade <9 Reference Reference   5-9 0.997 (0.552-1.799) 0.991 1.220 (0.802-1.870) 0.34   >15 0.843 (0.291-2.444) 0.753 1.286 (0.572-2.895) 0.54   Child-Pugh Classification A/B 1.090 (0.585-2.032) 0.786 0.973 (0.599-1.582) 0.91   ALBI grade I Reference Reference Reference Reference III 0.898 (0.543-1.486) 0.676 1.141 (0.787-1.655) 0.482   III 1.249 (0.485-3.215) 0.645 1.443 (0.648-3.214) 0.362
5-9 0.997 (0.552-1.799) 0.991 1.220 (0.802-1.870) 0.34   >15 0.843 (0.291-2.444) 0.753 1.286 (0.572-2.895) 0.54   Child-Pugh Classification A/B 1.090 (0.585-2.032) 0.786 0.973 (0.599-1.582) 0.91   ALBI grade I Reference Reference Reference 1.141 (0.787-1.655) 0.44   III 1.249 (0.485-3.215) 0.645 1.443 (0.648-3.214) 0.36
>15 0.843 (0.291–2.444) 0.753 1.286 (0.572–2.895) 0.54   Child–Pugh Classification A/B 1.090 (0.585–2.032) 0.786 0.973 (0.599–1.582) 0.91   ALBI grade I Reference Reference Reference 1.141 (0.787–1.655) 0.48   III 1.249 (0.485–3.215) 0.645 1.443 (0.648–3.214) 0.36
Child–Pugh Classification A/B 1.090 (0.585–2.032) 0.786 0.973 (0.599–1.582) 0.973   ALBI grade I Reference Reference Reference 1.141 (0.787–1.655) 0.488   III 1.249 (0.485–3.215) 0.645 1.443 (0.648–3.214) 0.368
ALBI grade I Reference Reference   II 0.898 (0.543-1.486) 0.676 1.141 (0.787-1.655) 0.48   III 1.249 (0.485-3.215) 0.645 1.443 (0.648-3.214) 0.36
II 0.898 (0.543–1.486) 0.676 1.141 (0.787–1.655) 0.48   III 1.249 (0.485–3.215) 0.645 1.443 (0.648–3.214) 0.365
III 1.249 (0.485–3.215) 0.645 1.443 (0.648–3.214) 0.36
Laboratory examination
GGT (U/L) <45/≥45 1.396 (0.934–2.088) 0.104 1.650 (1.199–2.269) 0.00
ALB (g/L) <35/235 0.828 (0.461–1.487) 0.527 0.971 (0.621–1.519) 0.90
PT (S) ≤13/>13 1.191 (0.722−1.964) 0.495 0.885 (0.604−1.297) 0.53
AST (U/L) <40/>40 1.448 (0.885–2.367) 0.140 0.968 (0.678–1.381) 0.85
TBL (µmol/L) <20/≥20 0.919 (0.556–1.519) 0.741 0.823 (0.565–1.198) 0.30
AFP (ng/ml) <400/≥400 1.159 (0.682–1.971) 0.586 1.449 (0.963–2.179) 0.07
HBV No/Yes 0.737 (0.445-1.220) 0.235 0.741 (0.481-1.142) 0.17
HCV No/Yes 0.511 (0.223–1.170) 0.112 0.430 (0.213–0.870) 0.01

\*OS, overall survival; RFS, recurrence-free survival; HCC, hepatocellular carcinoma; BMI, body mass index; MELD, model for end-stage liver disease; ALBI, albumin-bilirubin; GGT, gamma-glutamyl transpeptidase; ALB, albumin; PT, prothrombin time; AST, aspartate aminotransferase; TBIL, total bilirubin; AFP, alpha fetoprotein; HBV, hepatitis B virus; HCV, hepatitis C virus.



Furthermore, the K–M survival curve revealed that high-risk individuals have a worse prognosis than low-risk patients (**Figure 5**). In both the OS and RFS analyses, the nomogram models outperformed the other factors, with C-indices of 0.750 (OS, 95% CI: 0.713–0.787) and 0.746 (RFS, 95% CI: 0.715–0.777) in the training cohort and 0.794 (OS, 95% CI: 0.739–0.849) and 0.757 (RFS, 95% CI: 0.708–0.806) in the validation cohort (**Table 3**).

prognosis of patients with HCC with different invasive treatments, including LT, LR, and minimally invasive approach (RFA or MWA). Although our nomograms had good performance in predicting survival at 3 and 5 years in patients with HCC, it might be further improved by integrating more types of data, such as pathological images and multi-omics data (17, 21), and by applying more advanced classification algorithms as used in cancer diagnosis and other biological problems (15, 22). In the future, we will explore these directions.

# DISCUSSION

To our knowledge, this is the first attempt to construct prognostic nomograms for OS and RFS to predict the

Multivariate analyses revealed that the choice of different invasive treatments (including LT, LR, RFA, and MWA) may be an important independent prognostic factor in the patients with HCC. We have further shown that minimally invasive approach



the validation cohort [(G) 3 years; (H) 5 years].



FIGURE 3 | The overall survival calibration curve for predicting patient survival at 3 years (A) and 5 years (B) in the training cohort and 3 years (C) and 5 years (D) in the validation cohort. The recurrence-free survival calibration curve for predicting patient survival at 3 years (E) and 5 years (F) in the training cohort and 3 years (G) and 5 years (G) and 5 years (H) in the validation cohort.

(RFA or MWA) was the strongest predictor, followed by LR and LT, but some studies reported retrospective studies contrary to our study. Resection is the preferred option for patients with early-stage liver cancer (BCLC 0/A) and confer 5-year OS rates of 64.2% (4, 23, 24). Studies by other investigators suggest that LT is the best treatment option for patients with early HCC (25, 26). Minimally invasive surgery has undeniably played a significant role in HCC treatment in recent years. RFA or MWA was a common type of minimally invasive surgery, they showed comparable outcome and similar survival rates with LR (27–29). Compared with our previous nomogram studies, we were

fortunate to have access to treatment-independent prognostic factors, including LT, LR, RFA, and MWA (30). As a result, we created predictive nomograms to predict OS and RFS in patients with HCC undergone various invasive therapies. The nomograms were validated as an effective tool for predicting long-term outcomes. The current findings will need to be confirmed by larger prospective investigations into why different invasive treatments have different outcomes in the future. However, it is worthy noticing that we only considered single factors in the current analyses and ignore the relationship among these factors, for example, correlation and collinearity.







cohort (D).

Theoretically, integrating the correlation (collinearity) among the factors into multivariate analyses will increase the size of feature space and should increase the performance of our model. However, it will also make the model more complicated, and thus, we consider it as a future work.

In our study, tumor number (solitary versus multiple) was shown to play a role in the OS nomogram as well as in the RFS nomogram. Several studies have identified the presence of multiple tumors as a crucial risk factor for recurrence, which is consistent with the findings of our study (31, 32). Patients with HCC have a poor prognosis due to metastasis and recurrence. There is a strong association between tumor number > 1 and 3year and 5-year OS, according to Xiao et al. (33). Compared with single tumor, multiple tumors are more prone to microvascular invasion (MVI) which will lead to increased tumor recurrence after surgery (34, 35). In the OS nomogram, gender, BMI, and tumor size were also independent prognostic risk factors of patients with HCC. Gender was a prognostic factor also find in the nomogram for predicting the prognosis of patients with HCC with pulmonary metastases (36). In Global Cancer Statistics 2020, the incidence and mortality rates of liver cancer are two to three times greater in men than in women (1). Women are generally at lower risk for the development of HCC compared with men, and this may be due, in part, to the beneficial effects of sex hormones (37, 38). Sex hormone therapy is one of the potential development avenues of HCC treatment as part of multimodal liver cancer treatment.

In patients undergoing LR for HCC, preoperative bodyweight is linked to long-term prognosis (39). Furthermore, BMI  $\ge 25$  kg/m<sup>2</sup> negatively affected the surgical outcomes of patients with HBV-related HCC (BMI < 25 kg/m<sup>2</sup> group: 3-, 5-, and 8-year survival rates of 88.3%, 81.6%, and 73.9%, respectively, versus

TABLE 3 | Ranking of clinical staging system using C-index for OS and RFS in the training and validation cohorts.

Variables		Traini	ng Cohort	Validation Cohort		
		c-index	95% CI	c-index	95% CI	
os	Nomogram	0.750	0.713-0.787	0.794	0.739–0.849	
	Operation	0.648	0.605-0.691	0.685	0.624-0.746	
	Tumor Size	0.571	0.528-0.614	0.508	0.443-0.573	
	BMI	0.550	0.507-0.593	0.544	0.483-0.605	
	Tumor Number	0.530	0.491-0.569	0.511	0.454-0.568	
	Gender	0.526	0.493-0.559	0.524	0.481-0.567	
RFS	Nomogram	0.746	0.715-0.777	0.757	0.708-0.806	
	Operation	0.684	0.653-0.715	0.678	0.625-0.731	
	GGT	0.540	0.507-0.573	0.525	0.474-0.576	
	HCV	0.520	0.502-0.538	0.506	0.477-0.535	
	Tumor Number	0.512	0.483-0.541	0.513	0.470-0.556	

OS, overall survival; RFS, recurrence-free survival; BMI, body mass index; GGT, gamma-glutamyl transpeptidase; HCV, hepatitis C virus.

BMI  $\ge 25$  kg/m<sup>2</sup> group 85.8%, 61.0%, and 48.1%, respectively) (40). Previous study had revealed that the beneficial BMI level for patients with HCC following MWA is 21.5 to 23.1 kg/m<sup>2</sup> and can therefore achieve a longer survival time (41). Hence, it is critical for patients with HCC with weight concerns to confirm the beneficial BMI levels and the need for further research for different treatments.

Tumor size was not associated with RFS, and increasing trends toward the mortality of all patients with OS were observed for patients with a tumor size of  $\leq 3$  cm (21.9%) compared with patients with a tumor size of  $3 < R \le 5$  cm (33.2%), which was consistent with previous work. However, in a large international study, large tumor size was the key parameter related to early HCC recurrence after LR, and they built a preoperative model for RFS in the entire cohort (low risk: 2year RFS 64.8%; intermediate risk: 2-year RFS 42.5%; and high risk: 2-year RFS 20.7%) (42). A previous study has also reported that tumor size was not an independent prognostic factor of OS or RFS after curative resection and did not influence survival in patients with HCC without vascular invasion (43). We pointed out that tumor size is an important risk factor for OS and RFS, but different invasive treatments can obtain good clinical results and effectively reduce the recurrence rate. Therefore, tumor size is not a prognostic risk factor for the RFS nomogram.

In the RFS nomogram, there were prognostic risk factors also include GGT and HCV. A 384-patient study has shown that GGT > 50 U/L and indocyanine green retention of 15 min (ICG-R15) > 10% were identified as preoperative independent risk factors affecting 1-, 3-, and 5-year RFS (72.8%, 43.3%, and 27%, respectively) (44). A meta-analysis shows that high pretreatment serum GGT level is significantly correlated with poor survival and unfavorable clinicopathological features in patients with HCC, suggesting that pretreatment serum GGT may be an economical and effective prognostic biomarker for patients with HCC (45). Surgical patients who received HCV treatment had improved RFS compared with those who did not (91 vs. 80 months, p = 0.03) (46). Our findings are consistent with the few prior studies that found patients with HCC with HCV infection as a protective prognostic factor for patients with HCC in different invasive approaches to treatments. Patients with nonviral HCC have poorer prognosis than those with HCV-HCC (47). The reason why patients with HCV-HCC improve survival is that, possibly, antiviral therapy and virus replication reduced

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cancerous HCV-HCC tissues (46, 48–50). As a result, more research studies into the impact of antiviral medication on the outcomes of patients with HCC following surgery are needed.

# CONCLUSIONS

Our study identified prognostic risk factors for OS and RFS in patients with early-stage HCC treated with different invasive treatments (including LT, LR, RFA, and MWA), and we established and validation two prognostic nomograms. Two nomograms will be clinical settings for customized risk assessment and surgical decision-making. Furthermore, developing personalized treatment regimens for patients with different prognoses is beneficial.

## DATA AVAILABILITY STATEMENT

The original contributions presented in the study are included in the article/supplementary material, further inquiries can be directed to the corresponding author.

# **ETHICS STATEMENT**

The studies involving human participants were reviewed and approved by the Ethics Reviewer Board of Tianjin Medical University Cancer Institute and Hospital (bc2019082). The patients/participants provided their written informed consent to participate in this study.

# **AUTHOR CONTRIBUTIONS**

Study concept: WL and N-NZ. Study design: WL, N-NZ, S-WZ, and W-WZ. Data collection: S-WZ, W-WZ, J-YL, W-TJ, Y-MZ, T-QS, LZ, YX, and Y-HZ. Analysis and interpretation of data: S-WZ, TL, and W-WZ. Manuscript drafting: S-WZ, W-WZ, TL, and J-YL. Revising the manuscript: N-NZ, and WL. Acquisition and review of data, provision of critical comments or suggestions: all authors. Manuscript final version approval: all authors.

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