

# Development and Validation of Prognostic Nomograms for Hepatocellular Carcinoma After Hepatectomy Based on Inflammatory Markers

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**Background:** The value of lactate dehydrogenase (LDH) compared with other inflammation-based scores in predicting the outcomes of hepatitis B virus (HBV)-related hepatocellular carcinoma (HCC) patients after curative resection remains unknown. This study aims to evaluate the predictive value of LDH and develop novel nomograms to predict postoperative recurrence and survival in these patients.

**Methods:** This study retrospectively collected 1560 patients with HBV-related HCC who underwent curative resection from four institutions in China. In total, 924 patients were recruited from our center and randomly divided into the training cohort (n = 616) and internal validation (n = 308) cohorts. Additionally, 636 patients were selected from three other centers as the external validation cohort. The C index of inflammation-based scores was calculated and compared in the training cohort. Novel models were developed according to multivariable Cox regression analysis in the training cohort and validated in the internal and external validation cohorts.

**Results:** LDH showed a higher C-index than other inflammation-based scores for recurrence survival (RFS, 0.60, 95% CI, 0.58–0.61) and overall survival (OS, 0.65, 95% CI, 0.63–0.68). The nomograms of RFS and OS were developed based on tumor diameter, macrovascular invasion, AFP, operative hemorrhage, tumor differentiation, tumor number and LDH and achieved a high C-index (0.78, 95% CI, 0.76–0.79 and 0.81, 95% CI, 0.79–0.83), which were remarkably higher than the C-indexes of the five conventional HCC staging systems (0.52–0.62 for RFS and 0.53–0.67 for OS). The nomograms were validated in the internal validation cohort (0.77 for RFS, 0.78 for OS) and external validation cohort (0.80 for RFS, 0.81 for OS) and performed well-fitted calibration curves.

**Conclusion:** The two nomograms based on inflammatory markers achieved optimal prediction for RFS and OS of patients with HBV-related HCC after hepatectomy.

**Keywords:** hepatocellular carcinoma, hepatectomy, inflammation-based scores, lactate dehydrogenase, nomogram

## Introduction

Hepatocellular carcinoma (HCC) is the fifth most frequent cancer and the third leading cause of cancer-related mortality globally.<sup>1–3</sup> Many staging systems have been developed to predict the overall survival (OS) of HCC patients, including the American Joint Committee on Cancer (AJCC) seventh edition,<sup>4</sup> Barcelona Clinic Liver Cancer (BCLC),<sup>5</sup> Okuda staging system,<sup>6</sup> Japan Integrated Staging Score (JIS),<sup>7</sup> Cancer of the Liver Italian Program (CLIP)<sup>8</sup> etc. However, none

of the staging systems can predict the OS of HCC patients accurately and are popularized worldwide. A powerful and universal predicting system based on objective measures is urgently needed.

Cancer-related inflammation is recognized as the seventh hallmark of cancer.<sup>9,10</sup> In addition, hepatitis B virus (HBV) is a major cause of HCC, especially in the Asia-Pacific region, including China.<sup>11,12</sup> Consequently, inflammation plays a vital role in HBV-related HCC. Lactate dehydrogenase (LDH) is a well-known serum inflammatory marker. It has been reported that the release of LDH from hepatocytes increases after HBV infection<sup>13,14</sup> and LDH is a well-identified prognostic factor in HCC.<sup>15–17</sup> However, there is no research to evaluate the predictive competence of LDH compared with other inflammation-based scores in HBV-related HCC patients after hepatectomy.

Herein, we aimed to evaluate the prognostic role of LDH in HBV-related HCC patients after hepatectomy compared with other inflammation-based scores and establish novel prediction systems with nomograms based on LDH, which give rise to a satisfying prognostic indication for HBV-related HCC patients after hepatectomy.

## Methods

### Study Populations and Design

We retrospectively collected consecutive patients diagnosed with HBV-related HCC who underwent liver resection as initial treatment at Sun Yat-sen University Cancer Center (SYSUCC) from June 2011 to September 2019.

Patients were included if they met the following eligibility criteria: (1) age 18 to 75 years; (2) primary resectable HBV-related HCC; (3) histological confirmation of HCC; (4) liver function at Child-Pugh class A; (5) an Eastern Cooperative Oncology Group (ECOG) performance status of 0. Patients were excluded if they met any of the following criteria: (1) coinfection with hepatitis virus C; (2) received preoperative treatment, including interventional therapy, radiofrequency ablation, etc.; (3) metastasis to extrahepatic sites, including lymph nodes, lung, etc.; (4) patients with incomplete clinical data; (5) patients who were lost to follow-up within 3 months after hepatectomy; (6) history of other malignancies; (7) non-R0 liver resection. In our center, a total of 924 consecutive patients were included and randomly distributed to a training cohort ( $n = 616$ ) and an internal validation cohort ( $n = 308$ ).

In addition, a total of 636 patients from three other institutions served as an external validation cohort: The First Affiliated Hospital of Sun Yat-sen University (Jan 2009 to Jun 2015,  $n = 312$ ), Hunan Provincial People's Hospital (Mar 2008 to Dec 2016,  $n = 222$ ) and Shunde Hospital of Southern Medical University (Jan 2009 to Dec 2016,  $n = 102$ ). The flowchart of patient selection is shown in [Figure 1S](#). This study was conducted in accordance with the Declaration of Helsinki and approved by the Ethics Committee Board of the four institutions.

### Inflammation-Based Prognostic Scores

Blood samples were obtained within 1 week anterior to the liver resection. The inflammatory-based scores were calculated and are described in detail in [Table 1S](#).

### Following Up

In this study, all patients were observed in the first month after surgery, every 3 months within 2 years and then every 6 months thereafter. Laboratory tests (including serum AFP level, liver function tests, and blood tests) and magnetic resonance imaging (MRI) were conducted in follow-up examinations.

### Definitions

Overall survival (OS) was defined as the interval between curative resection and death from any cause or the date of last follow-up. Recurrence-free survival (RFS) was defined as the time from the date of curative resection to the date at which HCC recurred. HBV-related HCC was defined as the persistence of serum hepatitis B surface antigen (HBsAg) positivity for more than 6 months before the diagnosis of HCC. We divided the histologic grade of tumor differentiation into three levels based on the Edmondson–Steiner (ES) classification:<sup>18</sup> ES stage I was defined as high differentiation, ES stage II was defined as medium differentiation, and ES stages III and IV were defined as low differentiation. Cirrhosis was defined histologically according to the pathology of resected liver specimens.

## Statistical Analysis

All statistical analyses were performed with SAS (version 26.0, SAS Institute, Cary, NC) and R 3.63 (R Foundation for Statistical Computing, Vienna, Austria, <https://www.R-project.org/>). Categorical variables are presented as frequencies and percentages and were compared by the chi-square test between two groups. Continuous variables are presented as the median with interquartile range and were compared by Student's *t* test. The scores and cut points of the GPS, mGPS, PI, PNI, NLR, and PLR were defined and calculated as described in previous studies,<sup>19,20</sup> which were generally recognized. In view of the lack of widely accepted cut points for the LDH and SII scores, we defined their optimal cut points (LDH = 219.2; SII = 293.6 for RFS and 366.6 for OS) using the “maxstat” R package. We calculated the C-index and area under the ROC curve (AUC) by the “Hmisc” and “time ROC” R packages to estimate the predictive value of the inflammation-based scoring system.<sup>21</sup>

Survival curves were generated by the Kaplan–Meier method and compared by log-rank. Univariable and multivariable Cox proportional hazards models were used to assess the risk factors for tumor recurrence and patient overall survival. The nomogram was formulated based on the risk factors in multivariate analysis and performed by the “rms” R package. The calibration curve was generated based on regression analysis. We used the “Hmisc” R package to compare the C-index of the nomogram and other staging systems. We used X-tile to determine the optimal cut off values to stratify the risk of patients based on nomogram scores. We applied the nomogram in the validation groups to confirm the predictive value by using the same statistical methods.

## Results

### Patient Characteristics

This study collected a total of 1560 HBV-related HCC patients receiving hepatectomy from four different institutions. The median follow-up time was 43.0 (95% CI 40.7–45.3) months. The total population consisted of 1315 (84%) males and 245 (16%) females. 1223 (78%) patients were under 60 years old. All patients were infected with HBV. The median tumor size was 5.6 (interquartile range: 3.98–8.1) cm. The median LDH level was 182.8 (interquartile range: 160.4–215.7) U/L. The other clinical and pathological characteristics of the patients are presented in Table 1.

There were no significant differences in baseline characteristics among the 3 cohorts except that there were more female patients and fewer cirrhosis patients in the external validation cohort than in the training and internal validation

**Table 1** Baseline Clinicopathological Characteristics of the Patients

| Variable                                  | Training Cohort<br>(n=616) | Internal Validation<br>Cohort (n=308) | External Validation<br>Cohort (n=636) | <i>p</i> |
|---|----------------------------|---------------------------------------|---------------------------------------|----------|
| Age, years (≤60:>60)                      | 487/129 (79/21)            | 237/71 (77/23)                        | 499/137 (78/22)                       | 0.762    |
| Sex (female: male)                        | 81/535 (13/87)             | 44/264 (14/86)                        | 120/516 (19/81)                       | 0.016    |
| Tumor size, cm                            | 6 (4, 8.6)                 | 5.7 (3.88, 8)                         | 5 (3.6, 8)                            | 0.151    |
| Tumor number (solitary: multiple)         | 504/112 (82/18)            | 263/45 (85/15)                        | 528/108 (83/17)                       | 0.395    |
| Tumor differentiation (high, medium: low) | 372/244 (60/40)            | 180/128 (58/42)                       | 401/235 (63/37)                       | 0.356    |
| Cirrhosis (no: yes)                       | 292/324 (47/53)            | 154/154 (50/50)                       | 82/554 (13/87)                        | <0.001   |
| Microvascular invasion (no: yes)          | 413/203 (67/33)            | 221/97 (69/31)                        | 458/178 (72/28)                       | 0.162    |
| Operative hemorrhage, mL (≤400:>400)      | 449/167 (73/27)            | 235/73 (76/24)                        | 450/186 (71/29)                       | 0.199    |
| Platelet, ×10 <sup>9</sup> /L             | 185 (147, 240)             | 181 (141.38, 227.5)                   | 174 (127, 225)                        | 0.092    |
| AFP, ng/mL (≤200:>200)                    | 405/211 (66/34)            | 194/114 (63/37)                       | 383/253 (60/40)                       | 0.129    |
| ALT, U/L                                  | 34.75 (25.8, 49.62)        | 37.35 (24.87, 52.18)                  | 36 (24, 53.02)                        | 0.736    |
| AST, U/L                                  | 33.6 (25.4, 47.58)         | 34.2 (27.28, 51.2)                    | 35 (23, 52.24)                        | 0.388    |
| ALB, g/L                                  | 43.1 (40.48, 45.1)         | 42.8 (40.3, 45.2)                     | 40.4 (37.2, 42.92)                    | 0.172    |
| TBIL, μmol/L                              | 12.15 (9.4, 16.3)          | 12.65 (9.8, 17.02)                    | 13.8 (11, 18.12)                      | 0.253    |
| PT, second                                | 11.8 (11.3, 12.4)          | 11.8 (11.28, 12.6)                    | 12.65 (11.9, 13.6)                    | 0.325    |
| LDH, U/L                                  | 184.2 (158.6217, 12)       | 185.05 (165.57, 215.55)               | 181.5 (159.2, 213.12)                 | 0.333    |

**Notes:** Categorical variables are described as frequencies and percentages. Continuous variables are described as mean ± standard deviation (SD) and median with interquartile range for parametric and non-parametric variables, respectively.

**Abbreviations:** AFP, alpha fetoprotein; ALT, alanine aminotransferase; AST, aspartate aminotransferase; ALB, albumin; TBIL, total bilirubin; PT, prothrombin time; LDH, lactate dehydrogenase.

cohorts from SYSUCC. The judgment of cirrhosis by pathologists from different institutions is diverse, which may lead to the different cirrhosis rates among the four institutions.

## Independent Prognostic Factors for RFS and OS

Our univariable analysis revealed that male, larger tumor size, lower tumor differentiation, more tumor number, microvascular invasion, cirrhosis, more operative hemorrhage, shorter resection margin, higher serum levels of AFP, AST, ALB and longer PT were significantly associated with tumor recurrence. Five of eight inflammation-based scores (LDH, GPS, mGPS, PI, SII) were dependent prognostic factors for RFS. Multivariate analysis identified that tumor size ( $p < 0.001$ ), tumor number ( $p < 0.001$ ), tumor differentiation ( $p = 0.005$ ), microvascular invasion ( $p = 0.003$ ) and LDH ( $p < 0.001$ ) were independent prognostic factors for RFS (Table 2).

Our univariable analysis revealed that larger tumor size, lower tumor differentiation, microvascular invasion, more operative hemorrhage, higher serum levels of AFP, AST, ALB and longer PT were significantly associated with unfavorable overall survival. All eight inflammation-based scores were dependent prognostic factors for OS, which indicated that inflammation plays a vital role in HCC. Multivariate analysis identified that tumor size ( $p = 0.001$ ), microvascular invasion ( $p < 0.001$ ), operative hemorrhage ( $p = 0.033$ ), AFP ( $p = 0.020$ ), LDH ( $p = 0.030$ ), mGPS ( $p < 0.001$ ) and SII ( $p = 0.001$ ) were independent prognostic factors for OS (Table 2).

## Comparison of LDH and the Current Commonly Used Inflammation-Based Prognostic Systems

Plots of the time-dependent AUC for RFS and OS are shown in Figure 1, LDH performed better in predicting RFS and OS. The details of the AUC and C-indexes of the eight inflammation-based scores for RFS and OS are provided in Tables 3 and 4. LDH had a higher C-index value than all of the other scoring systems for both RFS (0.60, 95% CI, 0.58–0.61) and OS (0.65, 95% CI, 0.63–0.68).

## Development of Inflammation-Based Nomograms for RFS and OS in the Training Cohort

According to the multivariate analysis, tumor size, tumor number, microvascular invasion and tumor differentiation were integrated to build the nomogram of RFS; tumor size, microvascular invasion, operative hemorrhage and AFP were integrated to build the nomogram of OS. Regarding inflammation-based prognostic scores, LDH was an independent prognostic factor for both RFS and OS, and mGPS and SII were independent prognostic factors for OS as well. In view of better predictive ability compared with other inflammation-based prognostic scores, LDH was selected to build the nomograms of both RFS and OS. The details of the nomograms are shown in Figure 2.

## Calibration and Validation of Novel Nomograms

The C-indexes of the nomograms for RFS and OS prediction were 0.78 (95% CI, 0.76–0.79) and 0.81 (95% CI, 0.79–0.83), respectively, in the training cohort. In the internal validation cohort, the C-indexes for RFS and OS prediction were 0.77 (95% CI, 0.75–0.79) and 0.78 (95% CI, 0.75–0.81), respectively. In the external validation cohort, the C-indexes for RFS and OS prediction were 0.80 (95% CI, 0.78–0.81) and 0.81 (95% CI, 0.79–0.83), respectively. The calibration plots for the probability of recurrence and survival at 1, 3 or 5 years showed fair agreement between the prediction by the nomogram and actual observation in the training cohort (Figure 3A and B), internal validation cohort (Figure 3C and D), and external validation cohort (Figure 3E and F).

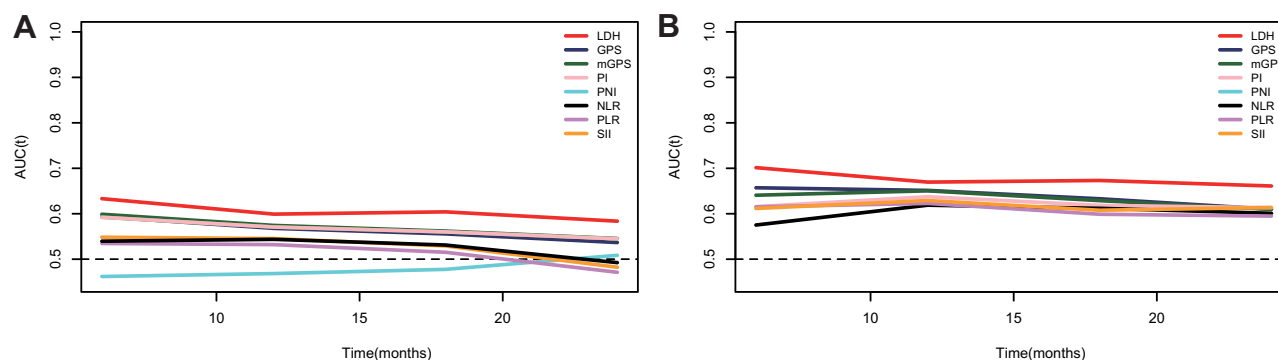
## Stratifying the Risk of Patients Based on Nomogram Scores

We grouped patients in the training cohort into three subgroups according to the optimal cut-off values determined by total nomogram scores. We stratified patients into low risk (score  $\leq 76$ ), medium risk (score 76–117) and high risk (score  $> 117$ ) of recurrence based on nomogram scores of RFS, and each group represented a distinct prognosis (Figure 4A).

**Table 2** Univariate and Multivariate Analysis of Risk Factors for Overall Survival and Recurrence-Free Survival

| Variables                                 | Overall Survival    |                |        |                       |               |         | Recurrence-Free Survival |               |        |                       |               |        |
|---|---------------------|----------------|--------|-----------------------|---------------|---------|--------------------------|---------------|--------|-----------------------|---------------|--------|
|   | Univariate Analysis |                |        | Multivariate Analysis |               |         | Univariate Analysis      |               |        | Multivariate Analysis |               |        |
|   | HR                  | 95% CI         | P      | HR                    | 95% CI        | P value | HR                       | 95% CI        | P      | HR                    | 95% CI        | P      |
| Age (≤60:>60)                             | 1.242               | (0.902–1.710)  | 0.184  |                       |               |         | 0.96                     | (0.769–1.199) | 0.718  |                       |               |        |
| Sex (female: male)                        | 0.898               | (0.605–1.333)  | 0.594  |                       |               |         | 1.447                    | (1.076–1.947) | 0.015  |                       |               |        |
| Tumor size, cm (≤5:>5)                    | 3.712               | (2.469–5.580)  | <0.001 | 2.170                 | (1.401–3.359) | 0.001   | 2.368                    | (1.913–2.932) | <0.001 | 2.073                 | (1.662–2.586) | <0.001 |
| Tumor number (solitary: multiple)         | 0.930               | (0.618–1.398)  | 0.726  |                       |               |         | 1.874                    | (1.557–2.255) | <0.001 | 1.710                 | (1.417–2.062) | <0.001 |
| Tumor differentiation (high, medium: low) | 1.811               | (1.363–2.405)  | <0.001 |                       |               |         | 1.577                    | (1.312–1.897) | <0.001 | 1.318                 | (1.089–1.596) | 0.005  |
| Cirrhosis (no: yes)                       | 0.825               | (0.638–1.068)  | 0.145  |                       |               |         | 0.713                    | (0.601–0.846) | <0.001 |                       |               |        |
| Microvascular invasion (no: yes)          | 2.899               | (2.181–3.853)  | <0.001 | 2.140                 | (1.592–2.876) | <0.001  | 1.946                    | (1.615–2.345) | <0.001 | 1.572                 | (1.168–2.117) | 0.003  |
| Operative hemorrhage, mL (400:>400)       | 2.150               | (1.618–2.858)  | <0.001 | 1.384                 | (1.027–1.866) | 0.033   | 1.562                    | (1.283–1.902) | <0.001 |                       |               |        |
| Resection margin, cm (≥1:<1)              | 1.162               | (0.877–1.541)  | 0.295  |                       |               |         | 1.271                    | (1.057–1.528) | 0.011  |                       |               |        |
| HBV DNA, U/mL (≤2000:>2000)               | 1.009               | (0.743–1.370)  | 0.956  |                       |               |         | 1.201                    | (0.991–1.456) | 0.062  |                       |               |        |
| WBC, ×10 <sup>9</sup> /L (≥4:<4)          | 0.723               | (0.394–1.330)  | 0.297  |                       |               |         | 0.848                    | (0.582–1.236) | 0.391  |                       |               |        |
| Platelet, ×10 <sup>9</sup> /L (≥100:<100) | 0.922               | (0.488–1.744)  | 0.804  |                       |               |         | 1.074                    | (0.723–1.595) | 0.725  |                       |               |        |
| AFP, ng/mL (≤200:>200)                    | 1.823               | (1.375–2.417)  | <0.001 | 1.413                 | (1.057–1.888) | 0.020   | 1.394                    | (1.153–1.684) | <0.001 |                       |               |        |
| ALT, U/L (≤50:>50)                        | 1.115               | (0.815–1.527)  | 0.496  |                       |               |         | 1.193                    | (0.972–1.466) | 0.092  |                       |               |        |
| AST, U/L (≤40:>40)                        | 1.515               | (1.143–2.009)  | 0.004  |                       |               |         | 1.608                    | (1.336–1.936) | <0.001 |                       |               |        |
| ALB, g/L (≥35:<35)                        | 3.309               | (2.010–5.447)  | <0.001 |                       |               |         | 1.489                    | 9(0.97–2.285) | 0.069  |                       |               |        |
| TBIL, μmol/L (≤17.1:>17.1)                | 1.230               | (0.887–1.704)  | 0.214  |                       |               |         | 1.030                    | (0.824–1.287) | 0.797  |                       |               |        |
| PT, s (≤13.5:>13.5)                       | 2.641               | (1.708–4.084)  | <0.001 |                       |               |         | 1.502                    | (1.048–2.152) | 0.027  |                       |               |        |
| LDH (0:1)                                 | 2.371               | (1.780–3.158)  | <0.001 | 1.413                 | (1.057–1.888) | 0.030   | 1.886                    | (1.548–2.297) | <0.001 | 1.500                 | (1.216–2.873) | <0.001 |
| GPS                                       |                     |                |        |                       |               |         |                          |               |        |                       |               |        |
| 0   |                     |                |        |                       |               |         |                          |               |        |                       |               |        |
| 1   | 1.891               | (1.416–2.514)  | <0.001 |                       |               |         | 1.573                    | (1.271–1.946) | <0.001 |                       |               |        |
| 2   | 8.572               | (5.082–14.457) | <0.001 |                       |               |         | 2.935                    | (1.801–4.784) | <0.001 |                       |               |        |
| mGPS                                      |                     |                |        |                       |               |         |                          |               |        |                       |               |        |
| 0   |                     |                |        |                       |               |         |                          |               |        |                       |               |        |
| 1   | 2.007               | (1.500–2.685)  | <0.001 | 1.011                 | (0.712–1.434) | 0.952   | 1.688                    | (1.360–2.094) | <0.001 |                       |               |        |
| 2   | 8.626               | (5.117–14.543) | <0.001 | 4.623                 | (2.607–8.196) | <0.001  | 2.958                    | (1.815–4.821) | <0.001 |                       |               |        |
| PI  |                     |                |        |                       |               |         |                          |               |        |                       |               |        |
| 0   |                     |                |        |                       |               |         |                          |               |        |                       |               |        |
| 1   | 2.123               | (1.601–2.815)  | <0.001 |                       |               |         | 1.652                    | (1.336–2.043) | <0.001 |                       |               |        |
| 2   | 2.834               | (1.603–5.012)  | <0.001 |                       |               |         | 2.379                    | (1.582–3.577) | <0.001 |                       |               |        |
| PNI (0:1)                                 | 1.695               | (1.239–2.320)  | <0.001 |                       |               |         | 0.998                    | (0.787–1.265) | 0.986  |                       |               |        |
| NLR (0:1)                                 | 1.455               | (1.072–1.975)  | 0.016  |                       |               |         | 0.928                    | (0.745–1.156) | 0.507  |                       |               |        |
| PLR (0:1)                                 | 1.771               | (1.334–2.350)  | <0.001 |                       |               |         | 1.076                    | (0.889–1.301) | 0.453  |                       |               |        |
| SII (0:1)                                 | 2.110               | (1.523–2.925)  | <0.001 | 1.801                 | (1.292–2.509) | 0.001   | 1.444                    | (1.161–1.795) | <0.001 |                       |               |        |

**Abbreviations:** HR, hazard rate; CI, confidence interval; HBV, hepatitis B virus; AFP, alpha fetoprotein; ALT, alanine aminotransferase; AST, aspartate aminotransferase; ALB, albumin; TBIL, total bilirubin; PT, prothrombin time; LDH, lactate dehydrogenase.



**Figure 1** Time-dependent AUC (area under ROC curve) plot for recurrence-free survival (A) and overall survival (B) prediction of inflammation-based scores.

Similar results were obtained from the internal validation cohort (Figure 4B) and the external validation cohort (Figure 4C).

Similarly, the nomogram of OS could also stratify patients into low risk (score  $\leq 126$ ), medium risk (score 126 to 193) and high risk (score  $>193$ ) of survival and each group represented a distinct group prognosis (Figure 4D). Similar results were observed in the internal validation cohort (Figure 4E) and in the external validation cohort (Figure 4F).

## Comparative Performance of Conventional Staging Systems

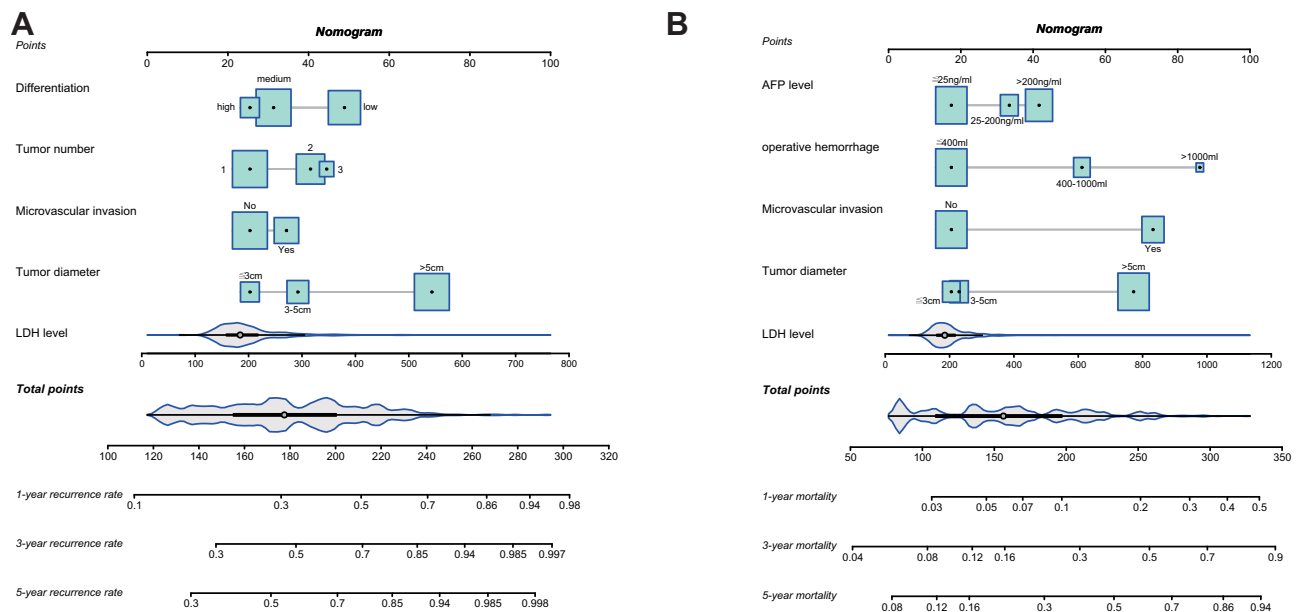
The C-index of the nomogram for RFS was 0.78 (95% CI, 0.76–0.79) in the training cohort, which was markedly higher than the C-indexes of five conventional staging systems, including AJCC (0.62, 95% CI, 0.60–0.63,  $P<0.001$ ), Okuda (0.60, 95% CI, 0.59–0.51,  $P<0.001$ ), JIS (0.60, 95% CI, 0.58–0.61,  $P<0.001$ ), CLIP (0.61, 95% CI, 0.59–0.62,  $P<0.001$ )

**Table 3** Comparison of the Performance and Discriminative Ability for Recurrence-Free Between the Inflammation-Based Scores

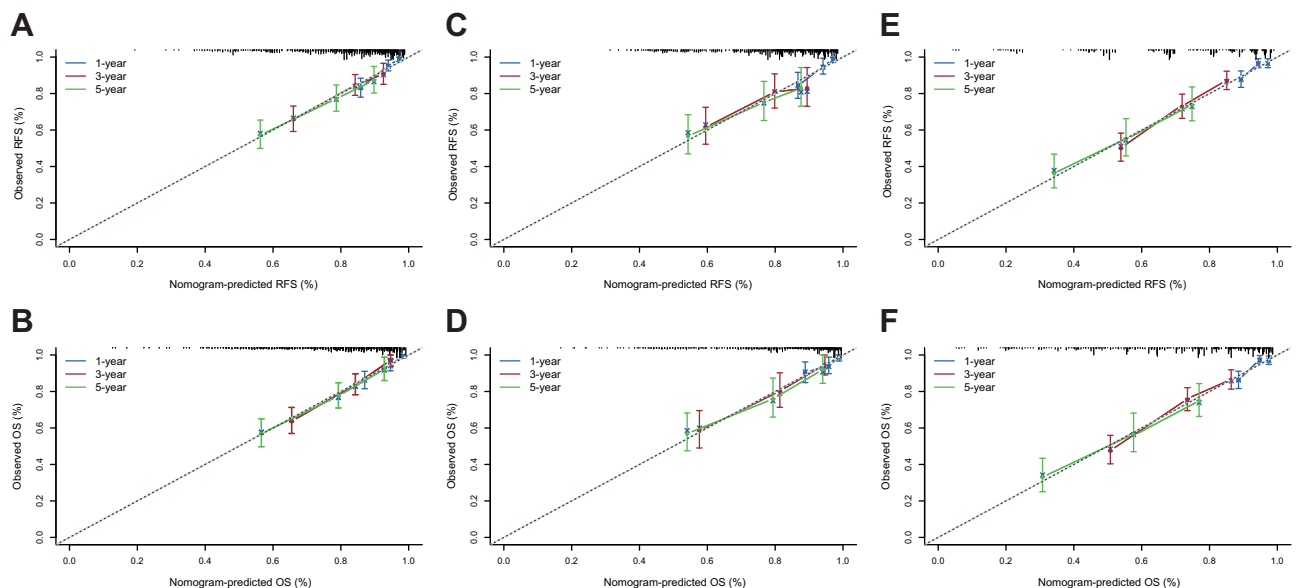
| Score | 6-Month AUROC (95% CI) | 12-Month AUROC (95% CI) | 18-Month AUROC (95% CI) | 24-Month AUROC (95% CI) | C-Index (95% CI) | P        |
|-------|------------------------|-------------------------|-------------------------|-------------------------|------------------|----------|
| LDH   | 0.63 (0.60–0.65)       | 0.59 (0.57–0.61)        | 0.60 (0.58–0.62)        | 0.58 (0.56–0.60)        | 0.60 (0.58–0.61) | —        |
| GPS   | 0.59 (0.57–0.61)       | 0.56 (0.55–0.58)        | 0.55 (0.54–0.57)        | 0.53 (0.52–0.55)        | 0.56 (0.55–0.57) | 0.016    |
| mGPS  | 0.59 (0.57–0.61)       | 0.57 (0.55–0.58)        | 0.56 (0.54–0.57)        | 0.54 (0.52–0.56)        | 0.56 (0.55–0.57) | 0.031    |
| PI    | 0.59 (0.57–0.61)       | 0.57 (0.55–0.58)        | 0.56 (0.54–0.57)        | 0.54 (0.52–0.56)        | 0.56 (0.55–0.57) | 0.027    |
| PNI   | 0.5 (0.48–0.51)        | 0.50 (0.48–0.51)        | 0.49 (0.47–0.50)        | 0.47 (0.45–0.48)        | 0.50 (0.49–0.51) | $<0.001$ |
| NLR   | 0.50 (0.48–0.52)       | 0.50 (0.49–0.52)        | 0.48 (0.47–0.50)        | 0.46 (0.44–0.47)        | 0.50 (0.49–0.51) | $<0.001$ |
| PLR   | 0.52 (0.50–0.54)       | 0.52 (0.50–0.54)        | 0.51 (0.50–0.53)        | 0.48 (0.46–0.50)        | 0.52 (0.51–0.53) | $<0.001$ |
| SII   | 0.55 (0.54–0.57)       | 0.54 (0.52–0.56)        | 0.54 (0.52–0.56)        | 0.52 (0.50–0.53)        | 0.47 (0.45–0.48) | 0.002    |

**Table 4** Comparison of the Performance and Discriminative Ability for Overall Survival Between the Inflammation-Based Scores

| Score | 6-Month AUROC (95% CI) | 12-Month AUROC (95% CI) | 18-Month AUROC (95% CI) | 24-Month AUROC (95% CI) | C-Index (95% CI) | P     |
|-------|------------------------|-------------------------|-------------------------|-------------------------|------------------|-------|
| LDH   | 0.69 (0.62–0.75)       | 0.66 (0.62–0.70)        | 0.66 (0.62–0.69)        | 0.65 (0.63–0.68)        | 0.65 (0.63–0.68) | —     |
| GPS   | 0.65 (0.59–0.71)       | 0.63 (0.60–0.66)        | 0.63 (0.60–0.66)        | 0.61 (0.58–0.63)        | 0.60 (0.58–0.62) | 0.037 |
| mGPS  | 0.64 (0.58–0.69)       | 0.65 (0.61–0.68)        | 0.62 (0.60–0.65)        | 0.60 (0.58–0.63)        | 0.60 (0.58–0.62) | 0.031 |
| PI    | 0.61 (0.56–0.67)       | 0.63 (0.60–0.67)        | 0.61 (0.59–0.64)        | 0.61 (0.58–0.63)        | 0.60 (0.58–0.62) | 0.020 |
| PNI   | 0.61 (0.56–0.67)       | 0.58 (0.55–0.61)        | 0.56 (0.53–0.59)        | 0.51 (0.49–0.54)        | 0.55 (0.54–0.57) | 0.006 |
| NLR   | 0.57 (0.51–0.62)       | 0.56 (0.53–0.59)        | 0.53 (0.50–0.56)        | 0.53 (0.50–0.55)        | 0.54 (0.53–0.56) | 0.002 |
| PLR   | 0.57 (0.52–0.62)       | 0.58 (0.55–0.61)        | 0.57 (0.54–0.60)        | 0.56 (0.53–0.58)        | 0.57 (0.56–0.59) | 0.032 |
| SII   | 0.57 (0.52–0.62)       | 0.61 (0.58–0.64)        | 0.60 (0.58–0.63)        | 0.60 (0.58–0.62)        | 0.59 (0.58–0.61) | 0.015 |



**Figure 2** Nomograms for predicting the 1-, 3- and 5-year recurrence (A) and mortality (B) rates in patients with hepatitis B virus-related hepatocellular carcinoma.

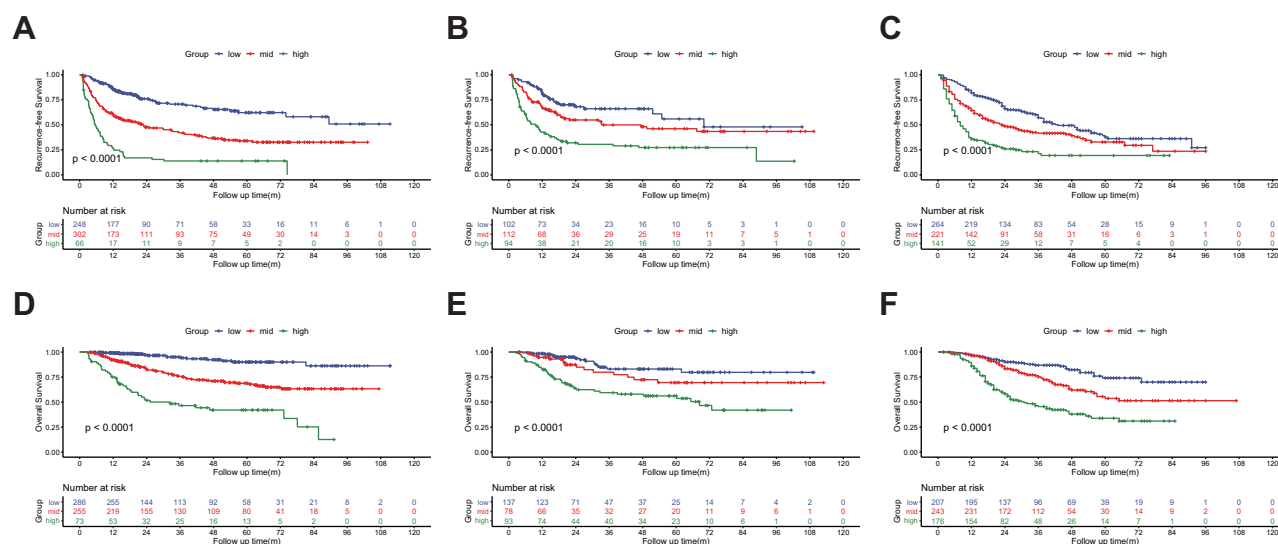


**Figure 3** The calibration curves for predicting the 1-, 3- and 5-year recurrence in the training cohort (A), internal validation cohort (C) and external validation cohorts (E) and the calibration curves for predicting the 1-, 3- and 5-year mortality in the training cohort (B), internal validation cohort (D) and external validation cohorts (F).

and BCLC (0.56, 95% CI, 0.55–0.57,  $P < 0.001$ ). We obtained similar results in the internal validation cohort and external validation cohort (Table 5).

The C-index of the nomogram for OS was 0.81 (95% CI, 0.79–0.83) in the training cohort, which was markedly higher than the C-indexes of five conventional staging systems, including AJCC (0.63, 95% CI, 0.61–0.65,  $P < 0.001$ ), Okuda (0.61, 95% CI, 0.60–0.62,  $P < 0.001$ ), JIS (0.65, 95% CI, 0.63–0.67,  $P < 0.001$ ), CLIP (0.66, 95% CI, 0.64–0.69,  $P < 0.001$ ) and BCLC (0.54, 95% CI, 0.52–0.56,  $P < 0.001$ ). We obtained similar results in the internal validation cohort and external validation cohort (Table 6).





## Discussion

In this study, we established two novel inflammation-based predictive systems with nomograms in a retrospective cohort of HBV-related HCC patients after hepatectomy. Among a series of inflammation-based scoring systems (LDH, GPS, mGPS, PI, PNI, NLR, PLR and SII), LDH was selected to construct the novel predictive system because of its better predictive ability. The other variables, including tumor diameter, macrovascular invasion, serum AFP level, operative

**Table 5** C-index and 95% CI of the Nomogram and Staging Systems (RFS)

| Score    | C-Index          |        |                            |        |                            |        |
|----------|------------------|--------|----------------------------|--------|----------------------------|--------|
|          | Training Cohort  | P      | Internal Validation Cohort | P      | External Validation Cohort | P      |
| nomogram | 0.78 (0.76–0.79) | —      | 0.77 (0.75–0.79)           | —      | 0.80 (0.78–0.81)           | —      |
| AJCC     | 0.62 (0.60–0.63) | <0.001 | 0.64 (0.62–0.66)           | <0.001 | 0.62 (0.60–0.63)           | <0.001 |
| Okuda    | 0.60 (0.59–0.61) | <0.001 | 0.60 (0.59–0.61)           | <0.001 | 0.58 (0.57–0.59)           | <0.001 |
| JIS      | 0.60 (0.58–0.61) | <0.001 | 0.61 (0.59–0.63)           | <0.001 | 0.61 (0.59–0.62)           | <0.001 |
| CLIP     | 0.61 (0.59–0.62) | <0.001 | 0.61 (0.59–0.63)           | <0.001 | 0.59 (0.57–0.61)           | <0.001 |
| BCLC     | 0.56 (0.55–0.57) | <0.001 | 0.53 (0.51–0.54)           | <0.001 | 0.52 (0.51–0.53)           | <0.001 |

**Abbreviations:** AJCC 7th, the American Joint Committee on Cancer seventh edition; BCLC, Barcelona Clinic Liver Cancer; CLIP, Modified Cancer of the Liver Italian Program; JIS, Japan Integrated Staging Score.

**Table 6** C-index and 95% CI of the Nomogram and Staging Systems (OS)

| Score    | C-Index          |        |                            |        |                            |        |
|----------|------------------|--------|----------------------------|--------|----------------------------|--------|
|          | Training Cohort  | P      | Internal Validation Cohort | P      | External Validation Cohort | P      |
| nomogram | 0.81 (0.79–0.83) | —      | 0.78 (0.75–0.81)           | —      | 0.81 (0.79–0.83)           | —      |
| AJCC     | 0.63 (0.61–0.65) | <0.001 | 0.62 (0.59–0.65)           | <0.001 | 0.67 (0.65–0.68)           | <0.001 |
| Okuda    | 0.61 (0.60–0.62) | <0.001 | 0.60 (0.58–0.62)           | <0.001 | 0.60 (0.59–0.61)           | <0.001 |
| JIS      | 0.65 (0.63–0.67) | <0.001 | 0.63 (0.60–0.66)           | <0.001 | 0.64 (0.62–0.66)           | <0.001 |
| CLIP     | 0.66 (0.64–0.69) | <0.001 | 0.66 (0.63–0.69)           | <0.001 | 0.64 (0.62–0.66)           | <0.001 |
| BCLC     | 0.54 (0.52–0.56) | <0.001 | 0.51 (0.49–0.53)           | <0.001 | 0.53 (0.53–0.54)           | <0.001 |

**Abbreviations:** AJCC 7th, the American Joint Committee on Cancer seventh edition; BCLC, Barcelona Clinic Liver Cancer; mCLIP, Modified Cancer of the Liver Italian Program; JIS, Japan Integrated Staging Score.



hemorrhage, tumor differentiation and tumor number were also selected according to multivariate analysis. The two nomograms showed a more accurate value in the prediction of postsurgical outcomes compared with the conventional HCC staging systems.

Serum LDH is a standardized and simple inflammatory marker, which is easy to use in the clinic, and is a well-identified prognostic marker in multiple malignancies, including HCC, colorectal cancer, breast cancer, lymphoma, melanoma, renal cell carcinoma, and germ cell tumors.<sup>15,22–29</sup> In addition, our study found that LDH displayed a better ability in predicting both RFS and OS of HBV-related HCC patients after hepatectomy compared with other inflammatory markers. Therefore, LDH was selected to establish the nomogram among a series of inflammatory markers.

Tumor diameter, AFP, MVI and tumor differentiation are well-known potential risk factors related to the postsurgical outcome of HCC.<sup>30–32</sup> Our study showed that these factors were also significantly associated with the prognosis of HBV-related HCC patients after hepatectomy. Several studies have reported that the incidence of MVI ranges from 15% to 57.1% in surgical specimens and is positively correlated with tumor diameter.<sup>33,34</sup> Interestingly, these two distinctive tumor pathological factors cooperatively affected the postsurgical prognosis of HBV-related HCC.

Operative hemorrhage was another surgical factor included in the OS nomogram model, whereas it was not a risk factor for RFS. It has been reported that operative hemorrhage is associated with poor postsurgical survival.<sup>35</sup> The reason may be that most HBV-related HCC patients have underlying cirrhosis, which causes an increased risk of haemorrhage due to abnormal liver function or decreased platelet counts.<sup>36</sup>

In 2016, Shen's research showed that HBeAg, AFP, resection margin, tumor number, tumor diameter, tumor capsule, MVI, and HBV-DNA level were independent risk factors for disease-free survival (DFS) or OS of HBV-related HCC patients. These independent risk factors were incorporated into the DFS and OS nomograms, respectively. Similar to Shen's research, our study showed that AFP, tumor number, tumor diameter and MVI were independent risk factors for RFS and OS. However, HBeAg and HBV-DNA levels were not independent risk factors in our study. Recent research reports that antiviral treatments can eliminate the adverse impacts of high baseline HBV-DNA levels on the survival of HBV-related HCC patients.<sup>37</sup> In recent years, effective antiviral drugs such as entecavir and tenofovir have been applied universally. Most patients in our study received NAs therapy and HBV-DNA levels were under good control, which could illustrate why HBeAg and baseline HBV-DNA levels were not independent risk factors in our study. In addition, inflammatory markers, which play a pivotal role in HBV-related HCC, were not considered in Shen's research.

Notably, patients were accurately stratified into three subgroups with significantly different risks of recurrence or mortality according to the proposed nomograms of RFS and OS. Therefore, it is helpful to guide clinicians to design follow-up regimens for patients with different risks of tumor recurrence after surgery and select proper candidates to receive effective adjuvant therapy. In addition, it is helpful to guide research to stratify patients when designing randomized controlled clinical trials of postsurgical adjuvant therapy.

Our study has several limitations. First, our nomograms are based on HBV-related HCC patients. Whether the nomograms can be used in HCC patients with aetiologies other than HBV infection needs further validation. Second, although our nomograms display a satisfactory C-index based on multicenter and large sample data, selection bias is inevitable as a retrospective study. Third, the enrolled patients in this study who underwent hepatectomy for HCCs within Milan criteria might be suitable for liver transplantation. Fourth, we only include inflammatory biomarkers in blood, inflammatory biomarkers in tumor tissue are not included, such as DNASE1L3.<sup>38</sup>

In conclusion, we developed and validated two nomograms based on inflammatory markers that could accurately and objectively predict RFS and OS in patients with HBV-related HCC after curative resection. For patients with high risk of recurrence after surgery, close surveillance regimens and adjuvant therapy should be considered.

## Data Sharing Statement

All data generated or analyzed during this study are included in this article. Further enquiries can be directed to the corresponding author.

## Ethics Approval Statement

This study was conducted according to the ethical guidelines of the 1975 Declaration of Helsinki. This research was approved by the institutional review board of four institutions.

## Informed Consent Statement

Informed consent was obtained from all subjects involved in the study.

## Author Contributions

All authors made a significant contribution to the work reported, whether that is in the conception, study design, execution, acquisition of data, analysis and interpretation, or in all these areas; took part in drafting, revising or critically reviewing the article; gave final approval of the version to be published; have agreed on the journal to which the article has been submitted; and agree to be accountable for all aspects of the work.

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## Disclosure

The authors have no conflicts of interest to declare in this work.

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