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## Retrospective Study

# Prediction of microvascular invasion in solitary hepatocellular carcinoma $\leq 5$ cm based on computed tomography radiomics

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## Abstract

### BACKGROUND

Liver cancer is one of the most common malignant tumors, and ranks as the fourth leading cause of cancer death worldwide. Microvascular invasion (MVI) is considered one of the most important factors for recurrence and poor prognosis of liver cancer. Thus, accurately identifying MVI before surgery is of great importance in making treatment strategies and predicting the prognosis of patients with hepatocellular carcinoma (HCC). Radiomics as an emerging field, aims to utilize artificial intelligence software to develop methods that may contribute to cancer diagnosis, treatment improvement and evaluation, and better prediction.

### AIM

To investigate the predictive value of computed tomography radiomics for MVI in solitary HCC  $\leq 5$  cm.

### METHODS

A total of 185 HCC patients, including 122 MVI negative and 63 MVI positive patients, were retrospectively analyzed. All patients were randomly assigned to



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the training group ( $n = 124$ ) and validation group ( $n = 61$ ). A total of 1351 radiomic features were extracted based on three-dimensional images. The diagnostic performance of the radiomics model was verified in the validation group, and the Delong test was applied to compare the radiomics and MVI-related imaging features (two-trait predictor of venous invasion and radiogenic invasion).

## RESULTS

A total of ten radiomics features were finally obtained after screening 1531 features. According to the weighting coefficient that corresponded to the features, the radiomics score (RS) calculation formula was obtained, and the RS score of each patient was calculated. The radiomics model exhibited a better correction and identification ability in the training and validation groups [area under the curve: 0.72 (95% confidence interval: 0.58-0.86) and 0.74 (95% confidence interval: 0.66-0.83), respectively]. Its prediction performance was significantly higher than that of the image features ( $P < 0.05$ ).

## CONCLUSION

Computed tomography radiomics has certain predictive value for MVI in solitary HCC  $\leq 5$  cm, and the predictive ability is higher than that of image features.

**Key Words:** Hepatocellular carcinoma; Microvascular invasion; Radiomics; Image features; Computed tomography

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**Core Tip:** Microvascular invasion (MVI) is considered one of the most important factors for recurrence and poor prognosis of liver cancer. Thus, accurately identifying MVI before surgery is of great importance in making treatment strategies and predicting the prognosis of patients with hepatocellular carcinoma (HCC). This study showed that radiomics as an emerging method at present had a good diagnostic efficiency and exhibited better accuracy in predicting MVI than image features, indicating that radiomics is a more suitable method in predicting MVI in solitary HCC  $\leq 5$  cm.

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## INTRODUCTION

Liver cancer is one of the most common malignant tumors, and ranks as the fourth leading cause of cancer death worldwide[1]. Furthermore, more than half of liver cancers occur in China, where there is a high incidence of hepatitis B[2]. Moreover, the recurrence rate after hepatectomy is 70%[3], and microvascular invasion (MVI) is considered one of the most important factors for recurrence and poor prognosis of liver cancer[4]. MVI has a relatively high incidence in hepatocellular carcinoma (HCC), which ranges from 12.4% to 57.1%, and may occur even in solitary HCC  $\leq 2$  cm[5,6]. Thus, accurately identifying MVI before surgery is of great importance in making treatment strategies and predicting the prognosis of patients with HCC[7]. However, MVI can only be confirmed by histopathology *via* surgical resection at present. Therefore, the accurate prediction of MVI before surgery is desperately needed. Radiomics as an emerging field, which aims to utilize artificial intelligence software to develop methods that may contribute to cancer diagnosis, treatment improvement and evaluation, and better prediction[8]. At present, few studies have focused on the prediction of MVI in the early stage of HCC (which refers to solitary tumor with a size of  $\leq 5$  cm, without intrahepatic venous invasion[9]). The present study aimed to

investigate the predictive value of computed tomography (CT) radiomics for MVI in solitary HCC  $\leq 5$  cm.

## MATERIALS AND METHODS

### Patients

Patients were retrospectively collected from January 1, 2014 to November 15, 2018 (Hunan provincial People's Hospital). The inclusion criteria were: (1) Pathologically diagnosed hepatocellular carcinoma with MVI; (2) Solitary tumor with the maximum diameter of  $\leq 5$  cm; and (3) Enhanced CT scanning was performed before surgery. The exclusion criteria were: (1) Complicated with other malignant tumors, and multiple primary or recurrent liver cancer; (2) History of preoperative treatment; (3) CT revealed a vascular tumor thrombus or macrovascular invasion; or (4) The tumor boundary was difficult to determine. The flowchart for the screening of patients is presented in [Figure 1](#).

### Examination methods

The Philips (Brilliance iCT 256) and Neusoft (NeuViz 64EN) scanners were used, with a tube voltage of 100-200 kV, tube current of 171-313 mAs, scanning layer thickness of 5 mm, layer spacing of 5 mm, and matrix of  $1024 \times 1024$ . The contrast medium (iopromide injection, 300 mgI/mL) was injected using a high-pressure syringe through the anterior cubital vein at a rate of 3.5 mL/s and at a dose of 1.2 mL/kg. Dynamic contrast-enhanced imaging data acquisition was performed at fixed time points: For the arterial phase, acquisition occurred at approximately 25-33 s after administration; for the portal vein phase, it was 57-63 s, and for the delayed phase, 117-123 s.

### Observation of imaging features

The two-trait predictor of venous invasion (TTPVI) was defined as having two independent imaging characteristics at the same time, and the development of an internal tumor artery without the signs of low density at the tumor margin[10]. Radiogenomic invasion (RVI) comprised of three independent image features: Intratumoral artery, low-density ring, and tumor-liver difference[11]. If there was an intratumoral artery, but there was no low-density ring or tumor-liver difference, the tumor was considered to have RVI. The imaging features (TTPVI and RVI) were evaluated double-blindly by two radiologists (with three years and seven years of experience in abdominal radiology, respectively). If these radiologists had inconsistent evaluation results, a third senior radiologist (with 13 years of experience in abdominal radiology) would make the further confirmation. The detailed description is presented in [Figure 2](#).

### Drawing, image feature extraction, selection, and construction of region of interest

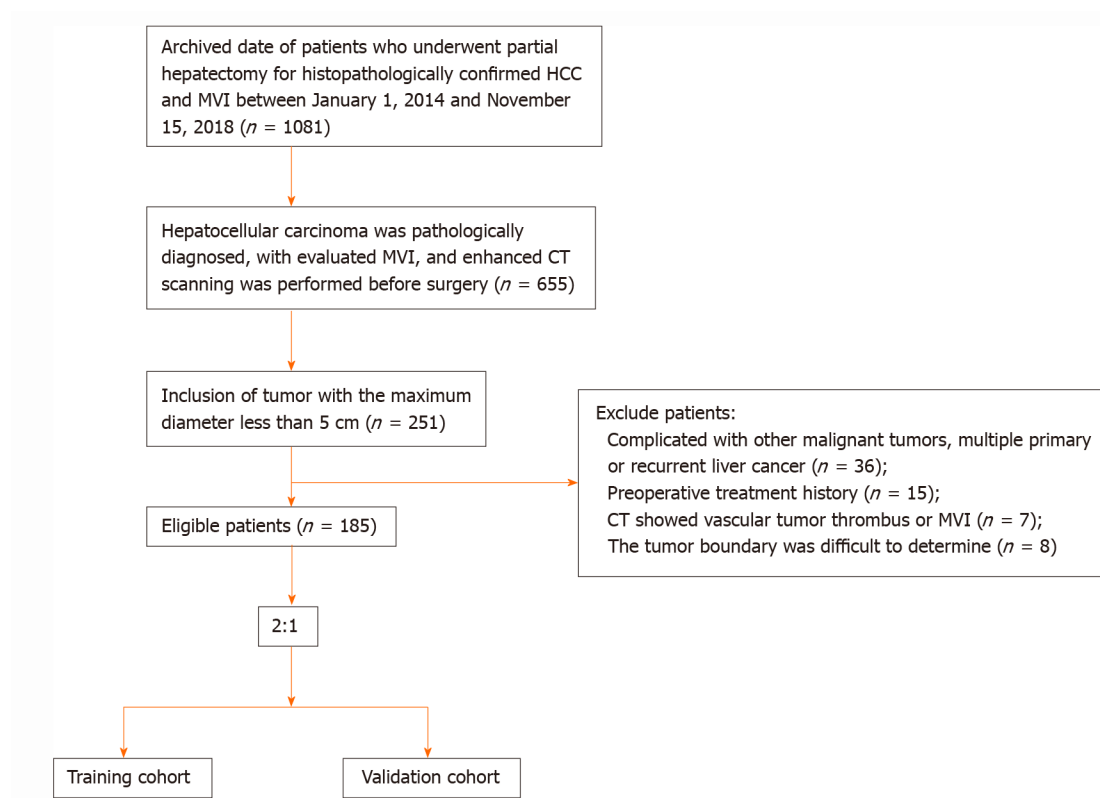
**Region of interest selection:** The CT images of the patients in the arterial phase were exported in DICOM format. Without knowing the pathological results, the radiologist with three years of experience in abdominal radiology used the 3D-Slice software ([www.slice.org](http://www.slice.org)) to delineate the region of interest (ROI) in each layer that contained the tumor, and finally formed the three-dimensional segmented image.

**Extraction of radiomic features:** After segmenting the images, the plug-in radiomics in the 3D-Slice software was used to analyze the original image data in the ROI, and 1351 candidate texture parameters were extracted, including the histogram features, morphological features, original features, and texture features.

**Selection of features correlated to the MVI status and construction of the radiomics tags:** First, the extracted intraclass features were evaluated by intraclass correlation coefficient (ICC), and features with  $ICC < 0.75$  were removed. Then, the least absolute shrinkage and selection operator (LASSO) algorithm was used for dimension reduction and feature construction, and the least feature variables were selected by 10-fold cross-validation with the minimal value. The selected features were modeled by Logistic regression, in order to generate the formula of the radiomics score (rad-score, RS) and calculate the RS score of each patient.

### Statistical analysis

The R language (version 3.4.0, <https://www.r-project.org>) was used for the statistical analyses, and  $P < 0.05$  was considered statistically significant. The chi-square test for



**Figure 1** Screening process for patients with liver cancer. HCC: Hepatocellular carcinoma; CT: Computed tomography; MVI: Microvascular invasion.

two independent samples was used for categorical variables, and Mann-Whitney *U*-test for two independent samples was used for continuous variables, in order to analyze the difference between the training group and verification group. The receiver operator characteristic curve and the area under the curve (AUC) were used to evaluate the prediction efficiency of the radiomic and image features, and the Delong test was used to determine whether there was a statistical difference between the two methods.

## RESULTS

### Patient clinicopathologic features

Finally, the study consisted of 185 patients. Tumor size ranged from 10 mm to 50 mm. The clinicopathological and CT features in the training group and verification group are presented in Table 1. There was no significant difference in scores for MVI, other clinical data, imaging features (TTPVI and RVI), or RS between the two groups ( $P > 0.05$ ), indicating the reasonable arrangement of the training group and verification group.

### Construction of radiomics tags

After removal by high correlation, 185 of the 1531 features remained. Then, ten features were selected by dimension reduction using the LASSO algorithm, as shown in Figure 3.

According to the weighting coefficient corresponding to the feature (Figure 3), the radiomics formula was obtained and used to calculate the histological score of each lesion in the training group and verification group. The formula is as follows: Rad-score =  $-0.66692761 - 0.02491645 \times \text{originalshapeFlatness} + 0.10564798 \times \log.\text{sigma}.1.0.\text{mm}.3\text{DglcmContrast} + 0.08789965 \times \log.\text{sigma}.2.0.\text{mm}.3\text{DglcmDependenceVariance} - 0.06847308 \times \log.\text{sigma}.2.0.\text{mm}.3\text{DglcmSizeZoneNonUniformityNormalized} - 0.01878760 \times \log.\text{sigma}.2.0.\text{mm}.3\text{DglcmSizeZoneNonUniformity} + 0.14275995 \times \log.\text{sigma}.3.0.\text{mm}.3\text{Dfirstorder90Percentile} + 0.04978394 \times \log.\text{sigma}.4.0.\text{mm}.3\text{DglcmSmallAreaEmphasis} - 0.02095096 \times \log.\text{sigma}.4.0.\text{mm}.3\text{DngtdmBusyness} + 0.06796763 \times \text{wavelet.HLLglcmSumSquares} + 0.0767876 \times 1\text{wavelet.HLLglcmLargeAreaLow}$



**Table 1 Clinical and imaging features of patients in the training group and verification group**

Feature	Training group (n = 124)	Verification group (n = 61)	Z value/ $\chi^2$ value	P value
Age (yr), median (quartile)	54 (47; 63)	52 (46; 62)	-0.900	0.368
Gender/cases				
Male	102	53	0.349	0.555
Female	22	8		
Hepatitis B	104	56	1.575	0.210
Liver cirrhosis	84	43	0.044	0.833
AFP (ng/mL)				
≤ 20	73	30	1.188	0.276
> 20	51	31		
MVI				
Negative	82	40	0.000	1.000
Positive	42	21		
Tumor size (mm), median (quartile)	36 (28; 44)	34 (27.5; 41)	-0.746	0.456
TTPVI				
Negative	56	25	0.145	0.703
Positive	68	36		
RVI				
Negative	93	40	1.362	0.243
Positive	31	21		
Rad-score, median (quartile)	-0.669 (-0.831; -0.546)	-0.640 (-0.780; -0.494)	-0.917	0.359

AFP:  $\alpha$ -fetoprotein; MVI: Microvascular invasion; TTPVI: Two-trait predictor of venous invasion; RVI: Radiogenomic invasion; Rad-score: Radiomics score.

GrayLevelEmphasis.

### Predictive performance of radiomics tags

The median RS [quartile interval] of MVI positive patients [training group: -0.574 (-0.695, -0.412); verification group: -0.495 (-0.644, -0.429)] was significantly higher than that of MVI negative patients [training group: -0.710 (-0.899, -0.610); verification group: -0.709 (-0.805, -0.589)] in both the training and verification groups, and the difference was statistically significant ( $P < 0.05$ ). The radiomics tags exhibited better diagnostic efficacy in both the training and verification groups, as shown in [Table 2](#).

### Comparison of predictive performance between radiomics tags and MVI-related image features

As shown in [Figure 4](#) and [Table 2](#), the diagnostic efficacy of the radiomics score was higher than that of the image features in the training group and verification group, and the difference was statistically significant ( $P < 0.05$ ).

## DISCUSSION

The present study revealed that radiomics, as an emerging method at present, exhibited good diagnostic efficiency and better accuracy in predicting MVI, when compared to image features, indicating that radiomics is a more suitable method for predicting MVI in solitary HCC  $\leq 5$  cm.

A number of studies have shown that tumor size and imaging features can predict MVI, in which TTPVI and RVI are good predictors with an ideal sensitivity and specificity[11]. However, the diagnostic performance of TTPVI and RVI in the present study was significantly lower than that in previous studies, which might be attributed

**Table 2 Comparison of predictive performance between radiomics tags and image features**

	Training group		Verification group	
	AUC (95%CI)	P value	AUC (95%CI)	P value
Rad-score	0.724 (0.584-0.863)		0.745 (0.655-0.834)	
TTPVI	0.590 (0.500-0.679)		0.522 (0.393-0.651)	
RVI	0.545 (0.462-0.628)		0.528 (0.401-0.655)	
Rad vs TTPVI		0.018		0.043
Rad vs RVI		0.002		0.048

AUC: Area under the curve; CI: Confidence interval; TTPVI: Two-trait predictor of venous invasion; RVI: Radiogenomic invasion; Rad-score: Radiomics score.

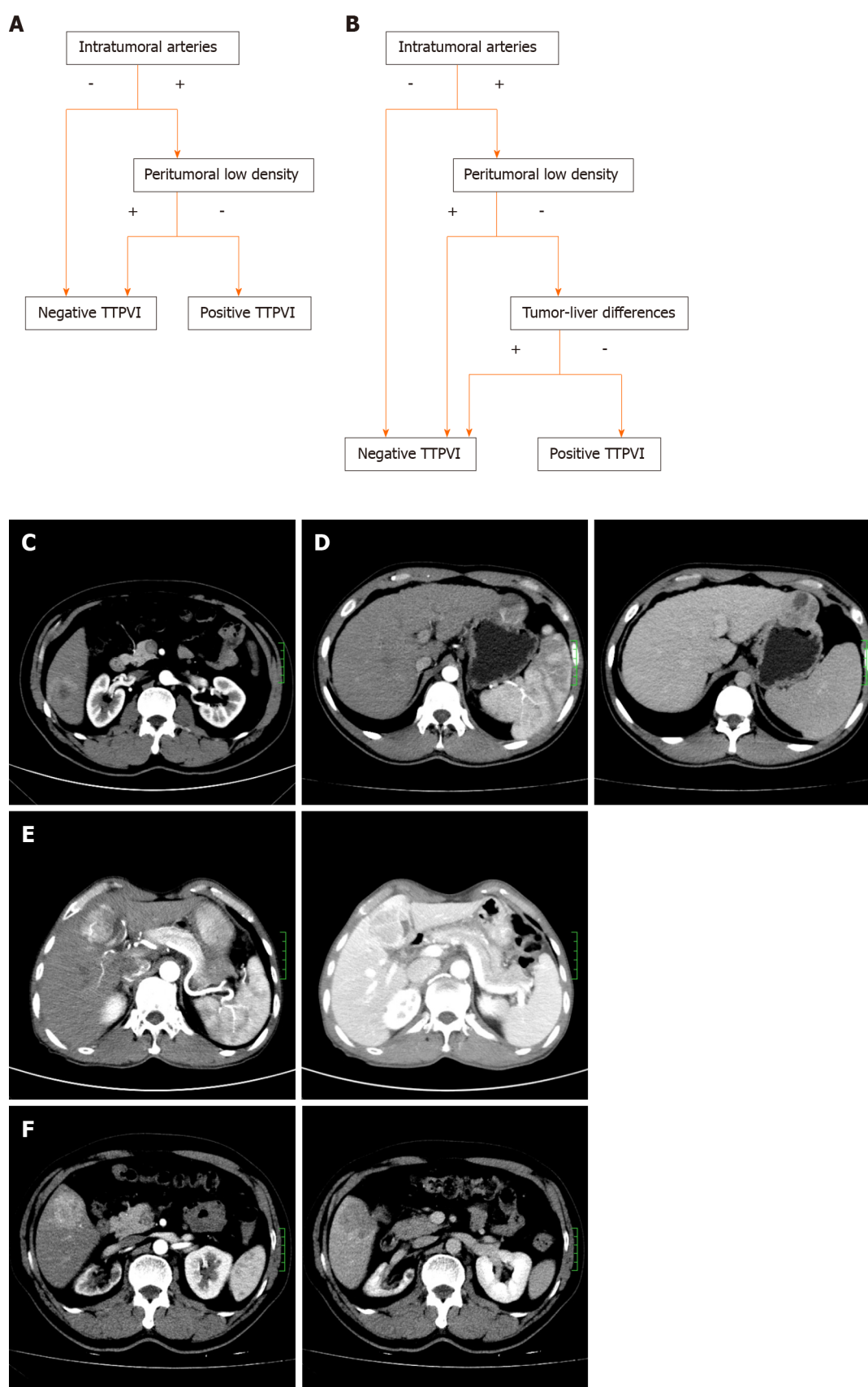
to the difference in tumor diameter of the study samples. Furthermore, previous studies did not define the tumor size, while the present study merely included patients with a tumor diameter of  $\leq 5$  cm. In addition, the imaging predictors, such as internal arteries and low-density shadow, were not commonly observed in cases with small tumors, which significantly reduced the positive rate of the MVI-related image features in the present study.

With the recent increase in development of radiomics, numerous studies have indicated that radiomics can reveal the pathological grade, prognosis, MVI, and treatment response of liver cancer. The nomogram of MVI in HCC based on CT radiomics established by Peng *et al*[12] exhibited good decision-making efficiency (AUC = 0.84), which was slightly higher than the results of the present study, but the model did not involve the tumor diameter. In another study[13], the diagnostic efficacy of predicting MVI in a tumor diameter of  $\leq 5$  cm based on radiomics (the AUC for the verification and verification group was 0.637 and 0.583, respectively) was slightly lower than that of the present study. Compared to that study[13], the present study adopted more stringent inclusion and exclusion criteria. Moreover, the present study only included solitary liver cancer with a diameter of  $\leq 5$  cm. Partial hepatectomy is the first choice for patients with solitary liver cancer  $\leq 5$  cm and good liver function[9]. However, if the lesion is not fully resected, the residual MVI near the surgical margin may be an important cause of recurrence in patients with HCC[14], and some studies have demonstrated that extended resection can reduce the early recurrence rate of patients with liver cancer complicated with MVI[15]. Therefore, the present study has certain reference value for surgery choice in patients with liver cancer. In addition, the present study excluded patients with a visible thrombus or the invasion of large blood vessels, rupture and bleeding of liver cancer, and intangible tumor boundary due to other reasons, because the ROI of these patients was difficult to delineate. Hence, measurement errors were hard to avoid. Furthermore, the present study employed the three-dimensional ROI of tumors to the extract radiomics features, which can better reflect the whole outline of the tumor, and allow for the extraction of more tumor information, when compared to two-dimensional ROI. Some studies have revealed that the feature extraction of the maximum cross-sectional area cannot represent the whole tumor[16]. Hence, the present study has obtained more objective prediction results.

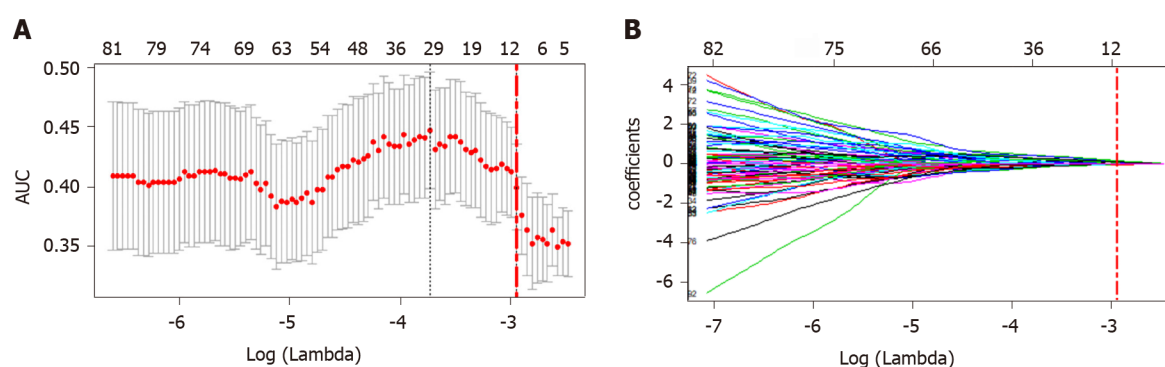
The present study had some limitations. First, the present retrospective and single-center study may have selection bias. Second, the present study only used arterial phase images. Multi-phase images may be utilized to obtain more tumor information and improve the diagnostic efficiency. Therefore, multi-center studies with large samples and multi-phase images would become our future research content.

## CONCLUSION

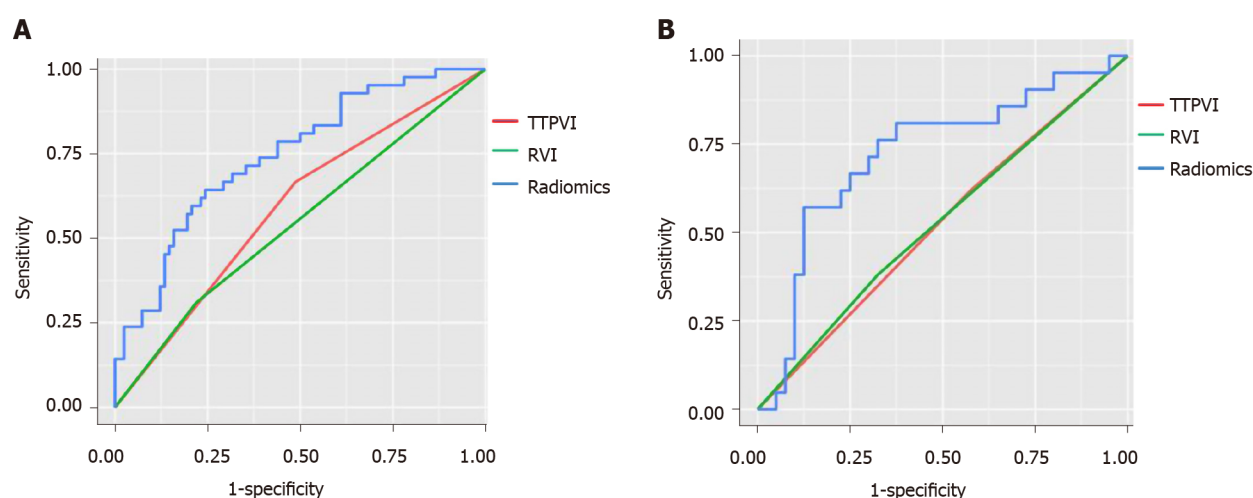
In conclusion, CT radiomics has certain predictive value for MVI in solitary HCC  $\leq 5$  cm. Compared to imaging features, the predictive ability of radiomics tags is significantly higher. The radiomics model of MVI would facilitate clinicians in choosing the appropriate treatment.



**Figure 2 Specific performance of two-trait predictor of venous invasion and radiogenomic invasion.** A and B: The discriminant process of two-trait predictor of venous invasion (TTPVI) (A) and radiogenomic invasion (RVI) (B); C: Negative intratumoral arteries: Negative TTPVI and RVI; D: Positive intratumoral arteries and peritumoral low density: Negative TTPVI and PVI; E: Positive intratumoral arteries, negative peritumoral low-density shadow, and positive tumor-liver differences: Positive TTPVI and negative RVI; F: Positive intratumoral arteries, negative peritumoral low density, and negative tumor-liver differences: Positive TTPVI and RVI. TTPVI: Two-trait predictor of venous invasion; RVI: Radiogenomic invasion.



**Figure 3** Selection of radiomic features using the least absolute shrinkage and selection operator-logistic regression model. A: Coefficient profile of 158 radiomics features against the area under the curve; B: Cross-validation curve. Red dotted vertical lines are drawn at the optimal log (Lambda) by using 10-fold cross-validation and the 1-SE criteria. Ten nonzero coefficients are chosen. AUC: Area under the curve.



**Figure 4** The two-trait predictor of venous invasion (green curve), radiogenomic invasion (red curve), and receiver operator characteristic of radiomics tag (blue curve) from the two groups. A: Training group; B: Verification group. TTPVI: Two-trait predictor of venous invasion; RVI: Radiogenomic invasion.

## ARTICLE HIGHLIGHTS

### Research background

Liver cancer is one of the most common malignant tumors, and ranks as the fourth leading cause of cancer death worldwide. Microvascular invasion (MVI) is considered one of the most important factors for recurrence and poor prognosis of liver cancer. Radiomics as an emerging field, aims to utilize artificial intelligence software to develop methods that may contribute to cancer diagnosis, treatment improvement, and evaluation and better prediction.

### Research motivation

At present, few studies have focused on the prediction of MVI in the early stage of hepatocellular carcinoma (HCC) (which refers to solitary tumor with a size of  $\leq 5$  cm, without MVI). Our study aimed to investigate the predictive value of computed tomography (CT) radiomics for MVI in solitary HCC  $\leq 5$  cm.

### Research objectives

This study aimed to investigate the predictive value of radiomics for MVI in solitary HCC  $\leq 5$  cm.

### Research methods

A total of 185 HCC patients, including 122 MVI negative and 63 MVI positive patients, were retrospectively analyzed. All patients were randomly assigned to the training

group ( $n = 124$ ) and validation group ( $n = 61$ ), at a ratio of 2:1. A total of 1351 radiomic features were extracted based on three-dimensional images. In the training group, the least absolute shrinkage and selection operator feature selection algorithm was used to reduce the dimensions, and the most relevant radiomic features of MVI were selected to calculate the image score (Rad-score, RS) of each patient. The diagnostic performance of the radiomics model was verified in the validation group, and the Delong test was applied to compare the radiomics and MVI-related imaging features (two-trait predictor of venous invasion and radiogenomic invasion).

### Research results

A total of ten radiomics features were finally obtained after screening 1531 features. According to the weighting coefficient that corresponded to the features, the RS calculation formula was obtained, and the RS score of each patient was calculated. The radiomics model exhibited a better correction and identification ability in the training and validation groups [area under the curve: 0.72 (95% confidence interval: 0.58-0.86) and 0.74 (95% confidence interval: 0.66-0.83), respectively]. Its prediction performance was significantly higher than that of the image features ( $P < 0.05$ ).

### Research conclusions

CT radiomics has certain predictive value for MVI in solitary HCC  $\leq 5$  cm, and the predictive ability is higher than that of image features.

### Research perspectives

The accurate prediction of MVI before surgery is desperately needed. Radiomics as an emerging field, aims to utilize artificial intelligence software to develop methods that may contribute to cancer diagnosis, treatment improvement and evaluation, and better prediction. At present, few studies have focused on the prediction of MVI in the early stage of HCC (which refers to solitary tumor with a size of  $\leq 5$  cm, without MVI). The present study aimed to investigate the predictive value of CT radiomics for MVI in solitary HCC  $\leq 5$  cm.

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