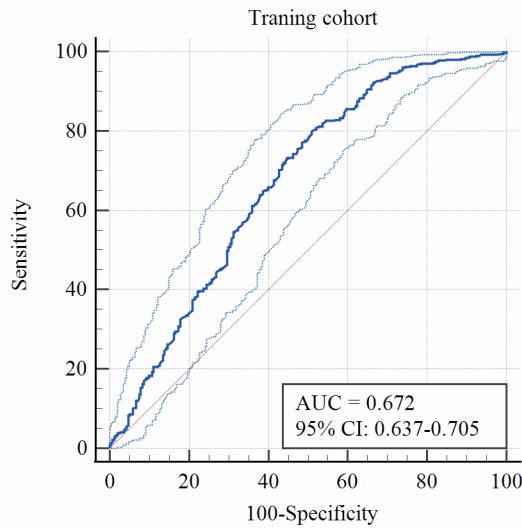


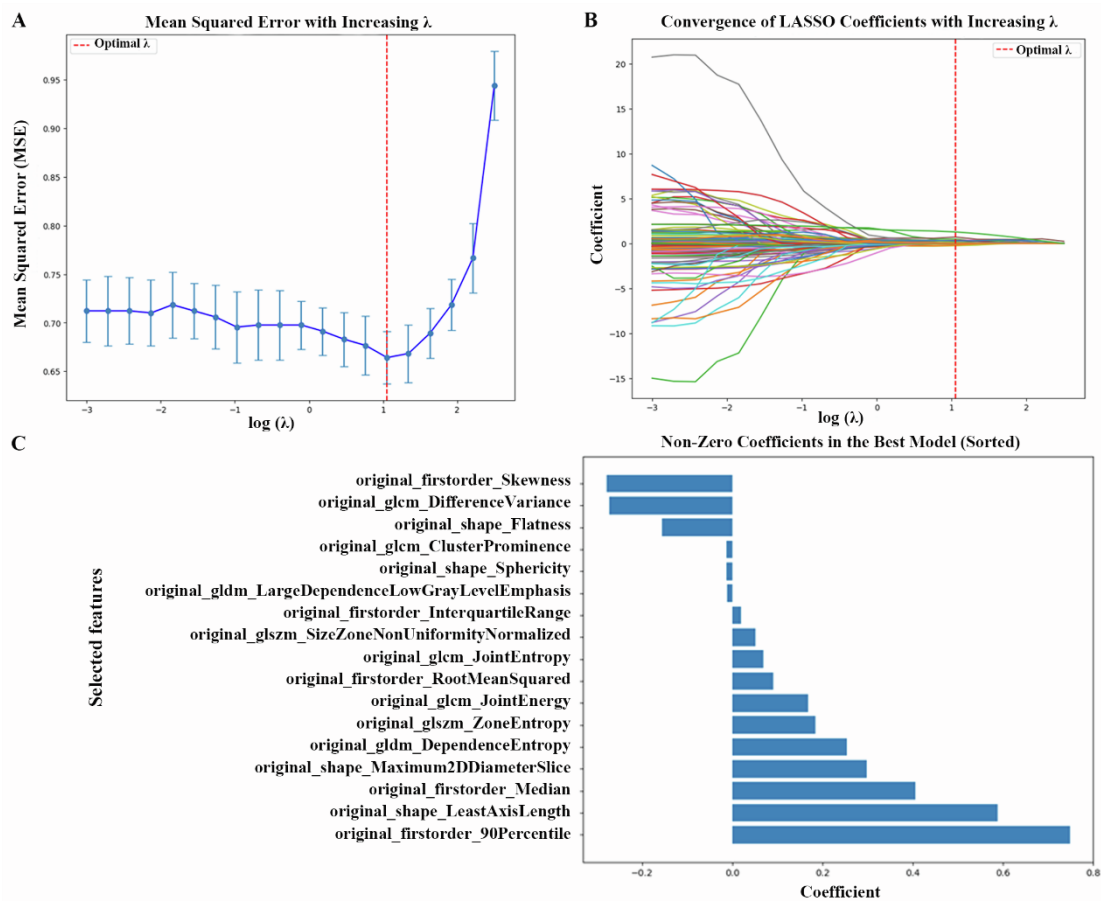
**Supplemental information**

**Multimodal integration to identify the invasion  
status of lung adenocarcinoma intraoperatively**

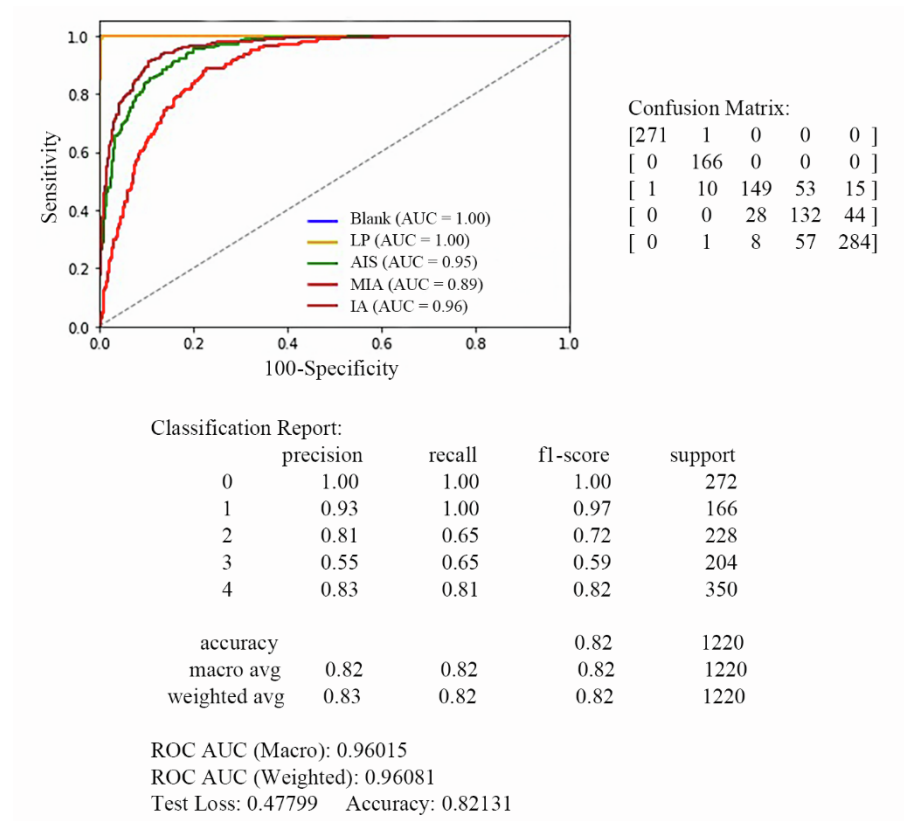
**Xueyun Tan, Feng Pan, Na Zhan, Sufei Wang, Zegang Dong, Yan Li, Guanghai Yang, Bo Huang, Yanran Duan, Hui Xia, Yaqi Cao, Min Zhou, Zhilei Lv, Qi Huang, Shan Tian, Liang Zhang, Mengmeng Zhou, Lian Yang, and Yang Jin**



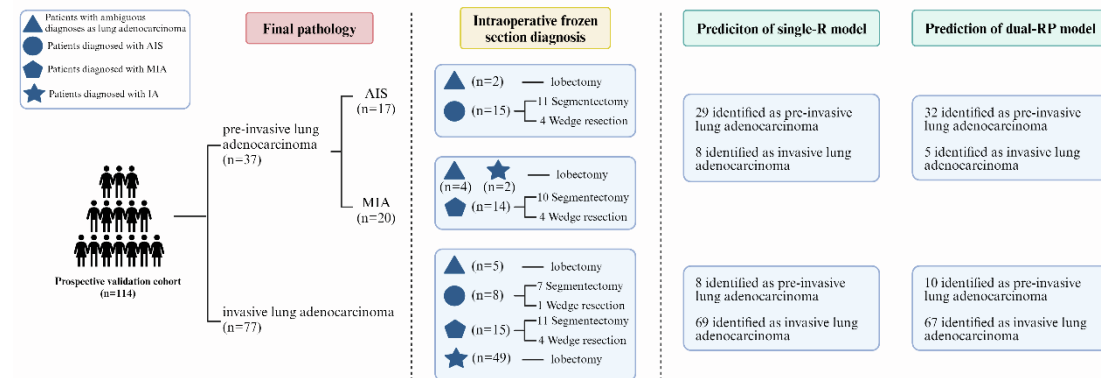
**Figure S1.** Performance of clinical indicators (age and eosinophil count) showed by ROC curve in the multimodal training cohort.



**Figure S2.** Radiomic feature selection using the least absolute shrinkage and the histogram of the Rad-score based on the selected features.



**Figure S3.** Predictive performance of EfficientNet B5 model to classify 1220 segmented patches into five pathologic classes, including “Blank”, “LP”, “AIS”, “MIA”, and “IA”.



**Figure S4.** Comparison of FP, intraoperative FS diagnosis, and predictions of single-R model and dual-RP model in the prospective validation cohort.

FP, final pathology; FS, frozen section; AIS, adenocarcinoma in situ; MIA, minimally invasive adenocarcinoma; Single-R model, single-modality radiomic model; Dual-RP model, radiomics + pathology dual-modality model.

**Table S1. Radiomic feature groups chosen in this study**

Feature Names	Feature Names
original_shape_Elongation	original_glcm_InverseVariance
original_shape_Flatness	original_glcm_MaximumProbability
original_shape_LeastAxisLength	original_glcm_SumEntropy
original_shape_MajorAxisLength	original_glcm_SumSquares
original_shape_Maximum2DDiameterColumn	original_glrIm_GrayLevelNonUniformity
original_shape_Maximum2DDiameterRow	original_glrIm_GrayLevelNonUniformityNormalized
original_shape_Maximum2DDiameterSlice	original_glrIm_GrayLevelVariance
original_shape_Maximum3DDiameter	original_glrIm_HighGrayLevelRunEmphasis
original_shape_MeshVolume	original_glrIm_LongRunEmphasis
original_shape_MinorAxisLength	original_glrIm_LongRunHighGrayLevelEmphasis
original_shape_Sphericity	original_glrIm_LongRunLowGrayLevelEmphasis
original_shape_SurfaceArea	original_glrIm_LowGrayLevelRunEmphasis
original_shape_SurfaceVolumeRatio	original_glrIm_RunEntropy
original_shape_VoxelVolume	original_glrIm_RunLengthNonUniformity
original_firstorder_10Percentile	original_glrIm_RunLengthNonUniformityNormalized
original_firstorder_90Percentile	original_glrIm_RunPercentage
original_firstorder_Energy	original_glrIm_RunVariance
original_firstorder_Entropy	original_glrIm_ShortRunEmphasis
original_firstorder_InterquartileRange	original_glrIm_ShortRunHighGrayLevelEmphasis
original_firstorder_Kurtosis	original_glrIm_ShortRunLowGrayLevelEmphasis
original_firstorder_Maximum	original_glszm_GrayLevelNonUniformity
original_firstorder_MeanAbsoluteDeviation	original_glszm_GrayLevelNonUniformityNormalized
original_firstorder_Mean	original_glszm_GrayLevelVariance
original_firstorder_Median	original_glszm_HighGrayLevelZoneEmphasis
original_firstorder_Minimum	original_glszm_LargeAreaEmphasis
original_firstorder_Range	original_glszm_LargeAreaHighGrayLevelEmphasis
original_firstorder_RobustMeanAbsoluteDeviation	original_glszm_LargeAreaLowGrayLevelEmphasis
original_firstorder_RootMeanSquared	original_glszm_LowGrayLevelZoneEmphasis
original_firstorder_Skewness	original_glszm_SizeZoneNonUniformity
original_firstorder_TotalEnergy	original_glszm_SizeZoneNonUniformityNormalized
original_firstorder_Uniformity	original_glszm_SmallAreaEmphasis
original_firstorder_Variance	original_glszm_SmallAreaHighGrayLevelEmphasis
original_glem_Autocorrelation	original_glszm_SmallAreaLowGrayLevelEmphasis
original_glcm_JointAverage	original_glszm_ZoneEntropy
original_glcm_ClusterProminence	original_glszm_ZonePercentage
original_glem_ClusterShade	original_glszm_ZoneVariance
original_glcm_ClusterTendency	original_gldm_DependenceEntropy
original_glcm_Contrast	original_gldm_DependenceNonUniformity
original_glem_Correlation	original_gldm_DependenceNonUniformityNormalized
original_glcm_DifferenceAverage	original_gldm_DependenceVariance
original_glcm_DifferenceEntropy	original_gldm_GrayLevelNonUniformity

original_glem_DifferenceVariance	original_gldm_GrayLevelVariance
original_glem_JointEnergy	original_gldm_HighGrayLevelEmphasis
original_glem_JointEntropy	original_gldm_LargeDependenceEmphasis
original_glem_Imc1	original_gldm_LargeDependenceHighGrayLevelEmphasis
original_glem_Imc2	original_gldm_LargeDependenceLowGrayLevelEmphasis
original_glem_Idm	original_gldm_LowGrayLevelEmphasis
original_glem_Idmn	original_gldm_SmallDependenceEmphasis
original_glem_Id	original_gldm_SmallDependenceHighGrayLevelEmphasis
original_glem_Idn	original_gldm_SmallDependenceLowGrayLevelEmphasis

**Table S2. Univariable and multivariable analyses in clinical indicators associated with invasiveness in multimodal training cohort.**

Characteristics	pre-invasive (254)	invasive (507)	Univariable analysis HR (95%CI)	P value	Multivariable analysis HR (95%CI)	P value
Age, years	53 (42-62)	60 (53-66)	1.06 (1.04-1.07)	<0.001	1.10 (1.04-1.17)	0.001
Sex			0.44 (0.32-0.62)	<0.001	0.73 (0.29-1.81)	0.493
Male	63 (24.8)	216 (42.6)				
Female	191 (75.2)	291 (57.4)				
Smoking history			1.26 (0.55-2.91)	0.583		
Yes	43 (16.9)	98 (19.3)				
No	211 (83.1)	409 (80.7)				
LDH, U/L	181 (160-201)	181 (162-201)	1.00 (0.99-1.01)	0.850		
Hematologic indicators, median (IQR)						
WBCs, ×10 <sup>9</sup> /L	5.27 (4.39-6.46)	5.27 (4.48-6.49)	0.99 (0.90-1.08)	0.779		
RBCs, ×10 <sup>9</sup> /L	4.21 (3.91-4.54)	4.29 (3.95-4.58)	1.08 (0.80-1.45)	0.636		
Hemoglobin, g/L	126 (119-138)	129 (120-140.75)	1.01 (1.00-1.02)	0.073		
HCT, %	38.40 (36.20-42.30)	39.10 (36.50-42.40)	1.02 (0.99-1.06)	0.216		
MCV, fl	92.10 (89.40-94.70)	92.60 (89.80-95.40)	1.02 (0.99-1.05)	0.222		
MCH, pg	30.30 (29.40-31.40)	30.70 (29.60-31.70)	1.09 (1.01-1.17)	0.031	1.12 (0.94-1.34)	0.216
MCHC, g/L	328 (322-336)	331 (323-338)	1.02 (1.00-1.03)	0.020	1.00 (0.97-1.03)	0.927
Platelets, ×10 <sup>9</sup> /L	213 (177-257)	206 (172-242)	1.00 (0.99-1.00)	0.061		
Neutrophils, ×10 <sup>9</sup> /L	2.91 (2.18-3.72)	2.99 (2.40-3.82)	1.00 (0.89-1.11)	0.965		
Lymphocytes, ×10 <sup>9</sup> /L	1.73 (1.47-2.16)	1.69 (1.35-2.10)	0.85 (0.65-1.11)	0.222		
NLR	1.65 (1.19-2.26)	1.76 (1.34-2.25)	1.06 (0.93-1.20)	0.390		
Monocytes, ×10 <sup>9</sup> /L	0.4 (0.3-0.5)	0.4 (0.3-0.5)	2.76 (0.97-7.81)	0.056		
Eosinophils, ×10 <sup>9</sup> /L	0.10 (0.06-0.16)	0.10 (0.07-0.19)	5.62 (1.29-24.49)	0.021	20.96 (1.18-373.04)	0.038
Basophils, ×10 <sup>9</sup> /L	0.03 (0.02-0.04)	0.03 (0.02-0.04)	0.35 (0-368.03)	0.765		
PDW, fl	12.70 (11.30-14.35)	13.20 (11.70-15.70)	1.09 (1.02-1.16)	0.016	1.02 (0.92-1.12)	0.753
MPV, fl	10.50 (9.70-11.20)	10.30 (9.30-11.20)	1.01 (0.96-1.06)	0.760		
Liver function indicators, median (IQR)						
AST, U/L	20 (17-23)	20 (17-24)	1.00 (0.98-1.01)	0.944		
ALT, U/L	16 (12-22)	17 (12-25)	1.01 (0.99-1.02)	0.319		

ALP, U/L	66.00 (51.25-77.75)	69.00 (56.00-82.25)	1.00 (1.00-1.01)	0.572		
γ-GT, U/L	17.00 (12.00-26.00)	19.00 (14.00-28.75)	1.00 (1.00-1.00)	0.559		
TBA, μmol/L	3.64 (2.40-6.44)	4.00 (2.50-6.18)	1.00 (0.97-1.02)	0.779		
TBIL, μmol/L	12.23 (9.30-15.57)	12.10 (9.30-15.40)	1.00 (0.97-1.03)	0.875		
DBIL, μmol/L	3.80 (2.90-5.00)	3.90 (2.80-5.10)	1.02 (0.94-1.11)	0.560		
Total protein, g/L	66.00 (63.00-70.18)	65.20 (61.90-68.70)	0.97 (0.94-1.00)	0.029	0.99 (0.94-1.04)	0.619
Albumin, g/L	41.40 (39.10-43.70)	40.65 (38.90-42.90)	0.94 (0.89-0.98)	0.006	0.94 (0.86-1.04)	0.216
Globin, g/L	25.00 (23.20-26.90)	24.20 (22.20-26.80)	0.96 (0.91-1.00)	0.055		
A/G	1.66 (1.55-1.80)	1.70 (1.51-1.86)	1.34 (0.71-2.53)	0.374		
Renal function indicators, median (IQR)						
BUN, mmol/L	5.05 (4.33-6.21)	5.42 (4.49-6.56)	1.15 (1.03-1.27)	0.010	0.94 (0.79-1.12)	0.475
Serum creatinine, μmol/L	59.00 (51.40-68.00)	64.00 (54.15-74.05)	1.02 (1.01-1.03)	0.003	1.03 (0.98-1.08)	0.220
UA, μmol/L	293.00 (245.00-345.00)	300.90 (247.00-361.50)	1.00 (1.00-1.00)	0.313		
Creatine kinase, U/L	76 (58-101)	72 (55-94)	1.00 (1.00-1.00)	0.776		
eGFR, mL/min per 1.73 m <sup>2</sup>	101.06 (92.67-111.64)	96.29 (89.39-103.14)	0.97 (0.95-0.98)	<0.001	1.06 (0.99-1.13)	0.106
Blood glucose and lipids, median (IQR)						
Glucose, mmol/L	4.85 (4.50-5.33)	5.00 (4.60-5.48)	1.20 (1.02-1.42)	0.032	1.04 (0.82-1.32)	0.752
TCh, mmol/L	4.52 (4.00-5.18)	4.56 (4.00-5.08)	1.01 (0.83-1.22)	0.920		
TG, mmol/L	1.05 (0.77-1.53)	1.20 (0.88-1.70)	1.28 (1.02-1.61)	0.032	1.16 (0.77-1.75)	0.485
HDLC, mmol/L	1.27 (1.05-1.53)	1.16 (0.99-1.44)	0.44 (0.26-0.72)	0.001	1.18 (0.50-2.81)	0.710
LDLC, mmol/L	2.64 (2.19-3.17)	2.68 (2.24-3.18)	1.02 (0.81-1.28)	0.861		
Coagulation function indicators, median (IQR)						
TT, seconds	16.90 (16.20-17.80)	17.30 (16.40-18.20)	1.18 (1.05-1.33)	0.005	1.07 (0.84-1.36)	0.581
FIB, g/l	2.64 (2.37-3.02)	2.78 (2.42-3.22)	1.44 (1.12-1.84)	0.004	1.33 (0.87-2.05)	0.190
APTT, seconds	27.90 (25.50-33.50)	31.30 (26.20-35.60)	1.05 (1.02-1.08)	0.002	0.98 (0.93-1.03)	0.437
INR	0.98 (0.95-1.06)	1.00 (0.95-1.06)	2.12 (0.38-12.00)	0.393		
PT, seconds	11.80 (11.10-12.80)	12.30 (11.40-12.90)	1.21 (1.06-1.39)	0.006	1.06 (0.83-1.36)	0.624

Data are presented as median (IQR) for continuous variables and n (%) for category variables.

HR, hazard ratio; CI, confidence interval; LDH, lactate dehydrogenase; WBC, white blood cell count; RBC, red blood cell count; HCT, hematocrit; MCV, mean corpuscular volume; MCH, mean corpuscular hemoglobin; MCHC, mean corpuscular hemoglobin concentration; NLR, neutrophil-to-lymphocyte ratio; PDW, platelet distribution width; MPV, mean platelet volume; AST, aspartate aminotransferase; ALT, alanine aminotransferase; ALP, alkaline phosphatase; γ-GT, γ-glutamyl transpeptidase; TBA, total bile acid; TBIL, total bilirubin; DBIL, direct bilirubin; A/G, albumin/globin; BUN, blood urea nitrogen; UA, urine acid; eGFR, estimated glomerular filtration rate; TCh, total cholesterol; TG, triglyceride; HDLC, high density lipoprotein cholesterol; LDLC, low density lipoprotein cholesterol; TT, thrombin time; FIB, fibrinogen; APTT, activated partial thromboplastin time; INR, international normalized ratio; PT: prothrombin time.

**Table S3. Characteristics of demographic information and clinical indicators of participants in radiology testing, pathology testing, and prospective validation cohorts.**

Characteristics	Radiology testing cohort (433)	Pathology testing cohort (230)	Prospective validation cohort (114)
Age, years	56 (48-64)	57.5 (50-65)	58 (52-65)
Sex			
Male	147 (33.9)	88 (38.3)	39 (34.2)
Female	286 (66.1)	142 (61.7)	75 (65.8)
Smoking history			
Yes	361 (83.4)	189 (82.2)	92 (80.7)
No	72 (16.6)	41 (17.8)	22 (19.3)
LDH, U/L	183 (163-205)	184 (162-203)	187 (169-204)
Hematologic indicators, median (IQR)			
WBCs, $\times 10^9/L$	5.15 (4.18-6.18)	5.27 (4.34-6.28)	5.14 (4.43-6.14)
RBCs, $\times 10^9/L$	4.16 (3.87-4.51)	4.22 (3.98-4.56)	4.20 (3.97-4.54)
Hemoglobin, g/L	126 (117-137)	128 (119-136)	127 (119-138)
HCT, %	37.50 (35.00-40.40)	0.42 (0.39-34.70)	38.00 (35.80-41.40)
MCV, fl	90.80 (88.60-93.40)	92.50 (89.90-95.00)	90.65 (88.50-94.30)
MCH, pg	30.70 (29.70-31.60)	30.40 (29.30-31.30)	30.30 (29.40-31.40)
MCHC, g/L	337 (331-342)	328 (321-335)	331.50 (326.00-339.00)
Platelets, $\times 10^9/L$	205 (170-243)	211 (173-250)	199 (173-239)
Neutrophils, $\times 10^9/L$	2.90 (2.20-3.50)	2.97 (2.24-3.63)	2.71 (2.20-3.44)
Lymphocytes, $\times 10^9/L$	1.68 (1.39-2.02)	1.70 (1.35-2.03)	1.71 (1.44-2.10)
NLR	1.71 (1.29-2.13)	1.75 (1.38-2.16)	1.48 (1.27-2.00)
Monocytes, $\times 10^9/L$	0.40 (0.30-0.50)	0.42 (0.32-0.51)	0.41 (0.35-0.50)
Eosinophils, $\times 10^9/L$	0.10 (0.07-0.18)	0.11 (0.07-0.19)	0.10 (0.08-0.18)
Basophils, $\times 10^9/L$	0.02 (0.00-0.03)	0.02 (0.01-0.04)	0.02 (0.01-0.04)
PDW, fl	13.00 (12.50-13.50)	12.90 (11.80-13.90)	12.75 (12.30-13.40)
MPV, fl	9.90 (8.70-10.80)	10.55 (9.80-11.50)	10.30 (9.70-11.00)
Liver function indicators, median (IQR)			
AST, U/L	21 (18-25)	19 (16-23)	21 (19-25)
ALT, U/L	18 (13-27)	15 (11-22)	20 (15-26)
ALP, U/L	62.00 (51.00-76.00)	67.00 (58.00-82.00)	69.00 (57.00-87.00)
$\gamma$ -GT, U/L	16.00 (12.00-23.00)	17.00 (13.00-23.00)	17.00 (14.00-27.00)
TBA, $\mu\text{mol/L}$	4.30 (2.70-6.70)	4.27 (2.31-6.28)	4.30 (2.60-6.80)
TBIL, $\mu\text{mol/L}$	12.10 (9.80-15.10)	12.40 (9.20-15.70)	11.50 (9.20-16.00)
DBIL, $\mu\text{mol/L}$	4.50 (3.50-5.60)	4.00 (2.90-5.30)	4.20 (3.30-6.30)
Total protein, g/L	64.10 (61.00-67.80)	64.70 (62.30-69.20)	64.35 (61.10-68.20)
Albumin, g/L	40.20 (38.40-42.50)	40.30 (37.80-42.80)	40.45 (38.30-42.60)
Globulin, g/L	23.80 (21.50-26.20)	25.10 (22.90-27.10)	24.10 (21.70-26.10)
A/G	1.70 (1.60-1.90)	1.62 (1.50-1.78)	1.70 (1.60-1.80)
Renal function indicators, median (IQR)			
BUN, mmol/L	5.14 (4.30-6.02)	5.39 (4.39-6.27)	5.22 (4.45-6.15)
Serum creatinine, $\mu\text{mol/L}$	60.20 (52.80-72.80)	62.00 (54.00-74.00)	60.80 (54.40-71.10)

UA, $\mu\text{mol/L}$	305.20 (258.00-359.00)	296.00 (246.00-352.80)	285.20 (234.95-352.20)
Creatine kinase, U/L	75 (56-101)	76 (55-105)	74 (59-108)
eGFR, mL/min per 1.73 m <sup>2</sup>	99.26 (90.16-107.83)	98.31 (91.00-104.01)	98.47 (89.88-103.77)
Blood glucose and lipids, median (IQR)			
Glucose, mmol/L	4.80 (4.50-5.30)	4.88 (4.54-5.26)	4.90 (4.60-5.30)
TCh, mmol/L	4.52 (3.95-5.10)	4.40 (3.72-4.80)	4.66 (4.05-5.32)
TG, mmol/L	1.10 (0.78-1.60)	1.17 (0.89-1.73)	1.19 (0.87-1.81)
HDLC, mmol/L	1.26 (1.05-1.54)	1.18 (0.94-1.36)	1.21 (1.06-1.53)
LDLC, mmol/L	2.55 (2.11-3.05)	2.46 (1.95-2.92)	2.67 (2.19-3.21)
Coagulation function indicators, median (IQR)			
TT, seconds	17.80 (17.10-18.50)	17.30 (16.60-18.15)	17.90 (17.25-18.55)
FIB, g/l	2.74 (2.46-3.11)	2.71 (2.31-3.07)	2.96 (2.58-3.34)
APTT, seconds	35.30 (33.30-37.50)	28.35 (25.50-34.15)	34.95 (33.70-37.55)
INR	1.09 (1.05-1.14)	1.03 (0.97-1.09)	0.95 (0.91-1.01)
PT, seconds	13.10 (12.70-13.50)	12.10 (11.30-12.90)	12.60 (12.25-12.90)

Data are presented as median (IQR) for continuous variables and n (%) for category variables. HR, hazard ratio; CI, confidence interval; LDH, lactate dehydrogenase; WBC, white blood cell count; RBC, red blood cell count; HCT, hematocrit; MCV, mean corpuscular volume; MCH, mean corpuscular hemoglobin; MCHC, mean corpuscular hemoglobin concentration; NLR, neutrophil-to-lymphocyte ratio; PDW, platelet distribution width; MPV, mean platelet volume; AST, aspartate aminotransferase; ALT, alanine aminotransferase; ALP, alkaline phosphatase;  $\gamma$ -GT,  $\gamma$ -glutamyl transpeptidase; TBA, total bile acid; TBIL, total bilirubin; DBIL, direct bilirubin; A/G, albumin/globin; BUN, blood urea nitrogen; UA, urine acid; eGFR, estimated glomerular filtration rate; TCh, total cholesterol; TG, triglyceride; HDLC, high density lipoprotein cholesterol; LDLC, low density lipoprotein cholesterol; TT, thrombin time; FIB, fibrinogen; APTT, activated partial thromboplastin time; INR, international normalized ratio; PT: prothrombin time.

**Table S4. Proportion of invasive classification for pathologic images in three cohorts identified by EfficientNet B5 model**

Cohort	Invasion status	Classification		
		AIS, %	MIA, %	IA, %
Multimodal training cohort (761)	AIS (84)	22.53 (15.80-34.70)	6.02 (3.29-9.00)	3.27 (1.47-8.04)
	MIA (170)	22.59 (14.64-34.46)	11.93 (6.38-19.82)	3.59 (0.81-9.05)
	IA (507)	18.72 (11.88-28.85)	17.63 (8.93-26.93)	20.39 (9.95-38.10)
Pathology testing cohort (230)	AIS (67)	23.82 (16.41-34.25)	4.60 (2.46-8.47)	2.30 (1.17-7.81)
	MIA (63)	29.28 (18.78-38.90)	9.30 (5.58-14.34)	3.52 (1.37-9.55)
	IA (100)	22.81 (12.65-34.74)	14.84 (9.52-22.98)	17.51 (6.48-32.92)
Prospective validation cohort (114)	AIS (17)	14.49 (10.16-21.45)	15.63 (5.16-21.79)	1.19 (0.20-2.91)
	MIA (20)	28.07 (17.69-33.75)	18.92 (12.57-33.26)	1.97 (0.74-5.17)
	IA (77)	14.90 (10.27-24.20)	21.40 (14.31-33.58)	19.93 (8.16-30.99)

Data are presented as median (IQR).

AIS, adenocarcinoma in situ; MIA, minimally invasive adenocarcinoma; IA, invasive adenocarcinoma.



**Table S5. NRI test for prediction improvements of dual-/multi-modality models compared to single-R or single-P models in multimodal training and prospective validation cohorts**

Initial model	Dual-/Multi-modality model	Multimodal training cohort		Prospective validation cohort	
		NRI (95% CI)	P value	NRI (95% CI)	P value
Single-R model	Dual-RC model	0.0277 (0.0002-0.0555)	0.0515	0.0270 (-0.0364-0.0905)	0.4038
	Dual-RP model	0.0652 (0.0258-0.1046)	0.0012	0.0530 (-0.0197-0.1257)	0.1532
	Multi-RPC model	0.0296 (-0.0096-0.0689)	0.1386	0.0660 (-0.0108-0.1428)	0.0921
Single-P model	Dual-PC model	0.1265 (-0.0237-0.2766)	0.0988	-0.5855 (-0.9571- -0.2139)	0.0020
	Dual-RP model	0.1403 (0.0892-0.1914)	<0.0001	0.1720 (0.0518-0.2922)	0.0050
	Multi-RPC model	0.1285 (0.0741-0.1828)	<0.0001	0.1580 (0.0281-0.2878)	0.0171

Data were metric value with 95% CI provided in parentheses.

NRI, net reclassification improvement; Single-R model, single-modality radiomic model; Single-P model, single-modality pathologic model; Dual-RC model, radiomics + clinical indicators dual-modality model; Dual-PC model, pathology + clinical indicators dual-modality model; Dual-RP model, radiomics + pathology dual-modality model; Multi-RPC model, radiomics + pathology + clinical indicators multi-modality model.

**Table S6. IDI test for prediction improvements of dual-RC, dual-RP and multi-RPC models compared to single-R model, and dual-PC, dual-RP and multi-RPC models compared to single-P model in multimodal training and prospective validation cohorts**

Initial model	Dual-/Multi-modality model	Multimodal training cohort		Prospective validation cohort	
		IDI (95% CI)	P value	IDI (95% CI)	P value
Single-R model	Dual-RC model	0.0073 (0.0016-0.0130)	0.0125	-0.0167 (-0.0345-0.0011)	0.0667
	Dual-RP model	0.0429 (0.0266-0.0593)	<0.0001	0.0402 (-0.0037-0.0841)	0.0724
	Multi-RPC model	0.0470 (0.0301-0.0639)	<0.0001	0.0238 (-0.0252-0.0728)	0.3415
Single-P model	Dual-PC model	0.0190 (0.0088-0.0292)	0.0003	-0.0194 (-0.0459-0.0070)	0.1502
	Dual-RP model	0.1402 (0.113-0.1674)	<0.0001	0.1947 (0.1215-0.2679)	<0.0001
	Multi-RPC model	0.1443 (0.1168-0.1717)	<0.0001	0.1782 (0.1070-0.2495)	<0.0001

Data were metric value with 95% CI provided in parentheses.

IDI, integrated discrimination improvement; Single-R model, single-modality radiomic model; Single-P model, single-modality pathologic model; Dual-RC model, radiomics + clinical indicators dual-modality model; Dual-PC model, pathology + clinical indicators dual-modality model; Dual-RP model, radiomics + pathology dual-modality model; Multi-RPC model, radiomics + pathology + clinical indicators multi-modality model.

**Table S7. AIC test for prediction improvements of multi-RPC model compared to single-/dual-modality models in the multimodal training cohort**

Models	AIC value
Single-P model	656.47
Single-R model	565.44
Dual-RC model	560.47
Dual-PC model	650.32
Dual-RP model	520.55
Multi-RPC model	517.85

Data were metric value with 95% CI provided in parentheses.

AIC, Akaike information criterion; Single-R model, single-modality radiomic model; Single-P model, single-modality pathologic model; Dual-RC model, radiomics + clinical indicators dual-modality model; Dual-PC model, pathology + clinical indicators dual-modality model; Dual-RP model, radiomics + pathology dual-modality model; Multi-RPC model, radiomics + pathology + clinical indicators multi-modality model.

**Table S8. Predictive performance of the models to distinguish AIS and MIA**

	Multimodal training cohort	Radiology testing cohort	Pathology testing cohort	Prospective validation cohort
Single-R model				
AUC	0.660 (0.598-0.718)	0.724 (0.668-0.775)	-	0.618 (0.444-0.772)
Accuracy	0.680 (0.618-0.737)	0.696 (0.639-0.749)	-	0.622 (0.448-0.775)
Sensitivity	0.544 (0.466-0.621)	0.567 (0.494-0.638)	-	0.800 (0.563-0.943)
Specificity	0.738 (0.631-0.828)	0.837 (0.745-0.906)	-	0.471 (0.230-0.722)
PPV	0.807 (0.740-0.860)	0.880 (0.820-0.922)	-	0.640 (0.519-0.745)
NPV	0.446 (0.395-0.498)	0.478 (0.433-0.524)	-	0.667 (0.421-0.846)
Single-P model				
AUC	0.658 (0.596-0.716)	-	0.680 (0.592-0.759)	0.685 (0.512-0.828)
Accuracy	0.684 (0.623-0.741)	-	0.600 (0.511-0.685)	0.622 (0.448-0.775)
Sensitivity	0.568 (0.490-0.644)	-	0.841 (0.727-0.921)	1.000 (0.832-1.000)
Specificity	0.750 (0.644-0.838)	-	0.478 (0.354-0.603)	0.353 (0.142-0.617)
PPV	0.821 (0.755-0.871)	-	0.602 (0.540-0.661)	0.645 (0.561-0.721)
NPV	0.463 (0.411-0.516)	-	0.762 (0.632-0.856)	1.000 (1.000-1.000)
Dual-RC model				
AUC	0.657 (0.595-0.716)	0.730 (0.675-0.781)	-	0.597 (0.423-0.755)
Accuracy	0.676 (0.614-0.733)	0.699 (0.643-0.752)	-	0.541 (0.369-0.705)
Sensitivity	0.444 (0.368-0.522)	0.649 (0.578-0.716)	-	1.000 (0.832-1.000)
Specificity	0.833 (0.736-0.906)	0.750 (0.649-0.834)	-	0.235 (0.068-0.499)
PPV	0.843 (0.763-0.899)	0.846 (0.791-0.888)	-	0.606 (0.542-0.667)
NPV	0.427 (0.387-0.468)	0.504 (0.448-0.560)	-	1.000 (1.000-1.000)
Dual-PC model				
AUC	0.647 (0.585-0.706)	-	0.679 (0.591-0.758)	0.641 (0.467-0.792)
Accuracy	0.644 (0.582-0.703)	-	0.615 (0.526-0.699)	0.622 (0.448-0.775)
Sensitivity	0.527 (0.449-0.604)	-	0.778 (0.655-0.873)	0.750 (0.509-0.913)

Specificity	0.786 (0.683-0.868)	-	0.522 (0.397-0.646)	0.588 (0.329-0.816)
PPV	0.832 (0.762-0.884)	-	0.605 (0.536-0.670)	0.682 (0.535-0.800)
NPV	0.452 (0.405-0.500)	-	0.714 (0.599-0.807)	0.667 (0.459-0.825)
Dual-RP model				
AUC	0.675 (0.613-0.732)	-	-	0.674 (0.500-0.818)
Accuracy	0.676 (0.614-0.733)	-	-	0.541 (0.369-0.705)
Sensitivity	0.527 (0.449-0.604)	-	-	0.600 (0.361-0.809)
Specificity	0.786 (0.683-0.868)	-	-	0.765 (0.501-0.932)
PPV	0.832 (0.762-0.884)	-	-	0.750 (0.542-0.884)
NPV	0.452 (0.405-0.500)	-	-	0.619 (0.472-0.747)
Multi-RPC model				
AUC	0.673 (0.612-0.731)	-	-	0.665 (0.491-0.811)
Accuracy	0.672 (0.610-0.729)	-	-	0.595 (0.421-0.753)
Sensitivity	0.544 (0.466-0.621)	-	-	0.600 (0.361-0.809)
Specificity	0.762 (0.657-0.848)	-	-	0.765 (0.501-0.932)
PPV	0.821 (0.754-0.874)	-	-	0.750 (0.542-0.884)
NPV	0.454 (0.404-0.505)	-	-	0.619 (0.472-0.747)

Data are presented as mean (95% CI).

AUC, area under the curve; NPV, negative predictive value; PPV, positive predictive value; Single-R model, single-modality radiomic model; Single-P model, single-modality pathologic model; Dual-RC model, radiomics + clinical indicators dual-modality model; Dual-PC model, pathology + clinical indicators dual-modality model; Dual-RP model, radiomics + pathology dual-modality model; Multi-RPC model, radiomics + pathology + clinical indicators multi-modality model.