## The development of a nomogram model for predicting left recurrent laryngeal nerve lymph node metastasis in esophageal cancer based on radiomics and clinical factors

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**Background:** The lymph node dissection for esophageal cancer is controversial. Some prediction models of lymph node metastasis (LNM) use the short diameter of lymph nodes measured by computed tomography (CT) examination as a predictor, but the size of that for judging metastasis is still controversial. However, radiomics can extract some features in tumors that cannot be obtained by naked eyes, which may have a higher value in predicting LNM. In this study, a nomogram was developed based on radiomics and clinical factors to predict left recurrent laryngeal nerve lymph node (RLNN) metastasis in patients with esophageal squamous cell carcinoma (ESCC).

**Methods:** There were 350 patients included in this retrospective study. And the postoperative pathological results determined whether there was left RLNN metastasis. A univariate analysis was conducted of the clinical data. The least absolute shrinkage and selection operator regression analysis was conducted to filter the radiomics features extracted from CT images. The multivariate logistic regression equation was used to construct a nomogram. The area under the curve (AUC) was used to evaluate the predictive ability. Due to the small sample size, we chose to perform internal validation after the model was established by 10-fold cross-validation, Harrell's concordance index (C-index), bootstrap validation and calibration.

**Results:** Ultimately, 3 indicators were screened out; that is, tumor location, surface volume ratio, and runlength non-uniformity. We then constructed the nomogram using these 3 indicators. The model had good accuracy and calibration performance. It has an AUC of 0.903 (95% confidence interval: 0.861–0.945), a sensitivity of 0.873, and a specificity of 0.756. Ten-fold cross-validation showed that the sensitivity and specificity of the training set were 88.08% and 75.81%, and the validation set had a sensitivity of 85.08% and a specificity of 75.49%. The Brier score was 0.074, and C-index was 0.904, which indicated good consistency between the actual and predicted results.

**Conclusions:** A nomogram constructed based on radiomics features and clinical factors can be used to predict the metastasis of left RLNN in patients with ESCC in a non-invasive way, which provided a reference for clinicians to formulate individualized lymph node dissection plans.

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#### Introduction

Esophageal cancer is a malignant digestive tract tumor. Progressive dysphagia is a typical symptom of esophageal cancer. Worldwide, esophageal cancer is the 7th most common cancer and ranks 6th in overall mortality (1). In China, esophageal cancer is the 6th most common cancer and the 4th leading cause of cancer-related death (2). According to the latest expert consensus, surgical resection is currently the most common and effective treatment for esophageal cancer (3), and lymph node metastasis (LNM) is the most critical factor affecting the prognosis of esophageal cancer patients. Thus, lymph node dissection is an essential part of radical surgery for esophageal cancer (4). However, the use of lymph node dissection in esophageal cancer is controversial (5,6).

Three-field lymph node dissection can effectively control local lesions, remove undetectable micro-metastases, help obtain accurate pathological staging, and improve the 5-year

#### **Highlight box**

#### Key findings

• The radiomics features combined with clinical factors can effectively predict the metastasis of left RLNN in ESCC and can be used to guide the dissection of lymph nodes before surgery.

#### What is known and what is new?

- The extent of lymph node dissection in esophageal cancer is controversial. The dissection of left RLNN brings some postoperative complications while obtaining certain benefits. However, there is still a lack of a better noninvasive way to predict left RLNN metastasis.
- Radiomics features combined with clinical factors are used to construct a nomogram to predict left RLNN metastasis in ESCC.

#### What is the implication, and what should change now?

 This nomogram provides clinicians with a non-invasive auxiliary method to predict left RLNN metastasis in esophageal cancer to guide the formulation of precise and personalized preoperative lymph node dissection plans for clinical patients. More accurate and effective radiomics feature extraction methods need to be further developed. overall survival (OS) rate of patients with esophageal cancer. However, the surgical trauma is greater than two-field lymph node dissection, and it increases the risk of related complications and the incidence of postoperative mortality (4).

Due to the lymphatic drainage system of esophageal cancer, LNM of esophageal cancer can be upward, downward, or skipped. Upper esophageal cancer tends to metastasize upward, lower esophageal cancer tends to metastasize downward, and LNM of middle esophageal cancer can spread either upward or downward (7). Recurrent laryngeal nerve lymph nodes (RLNNs) are the common sites for esophageal cancer LNM. Metastatic lymph nodes can spread along the recurrent laryngeal nerve (RLN), from the chest to the neck. The particularity of the anatomical location of the left RLNNs adjacent to the aorta and the RLN and the long surgical path increases the difficulty of lymph node dissection and increases postoperative complications, such as hoarseness and the pulmonary infection incidence rate. The selection of the appropriate lymph node for dissection according to preoperative pathological results or preoperative examinations can enable the trauma caused by dissection of the left RLNNs to be avoided, improve the cure rate of esophageal cancer, and reduce surgical trauma and thus improve patients' postoperative quality of life. Thus, the accurate preoperative determination of whether tumor metastasis occurs in the left RLNNs determines the extent of lymph node dissection in patients with esophageal cancer.

The gold standard for diagnosing LNM is pathological findings. A lymph node biopsy is an invasive procedure that can cause a certain degree of trauma to the patient, such as bleeding, pneumothorax, and even tumor dissemination. Invasive examinations, such as endoscopic ultrasonography and endoscopic ultrasonography fine needle aspiration place demanding requirements on the operator and carry certain risks, so it has limited application in the diagnose of LNMs in patients with esophageal cancer.

At present, the main non-invasive methods for preoperative lymph node evaluations in esophageal cancer patients include ultrasound, contrast-enhanced computed

tomography (CT), magnetic resonance imaging, and positron emission tomography-CT (PET-CT). In previous studies, contrast-enhanced CT and PET-CT have shown certain sensitivity and specificity in the clinical prediction of LNM in patients with esophageal cancer (8-10). However, there are still some limitations in predicting LNM in esophageal cancer.

A study has shown that CT has a lower accuracy in predicting lymph node involvement than endoscopic ultrasound (EUS) at 46–58%, compared to 72–80%. The sensitivity of 2-fluoro-2-deoxy-D-glucose positron emission tomography (FDG-PET) was 57% (range, 43–70%) and the specificity was 85% (range, 76–95%) (10). Some prediction models based on clinical data to predict LNM of esophageal squamous cell carcinoma (ESCC) mostly use the short diameter of lymph nodes as a predictor. Although they can achieve certain accuracy, the short diameter of lymph nodes as a threshold for predicting LNM is still controversial, and reactive lymph node enlargement easily causes false positives

CT is a traditional non-invasive method for obtaining tumor-related information. With continuous developments in medical imaging, Radiomics focuses on improvements of image analysis through the automatic analysis of a given imaging method. Medical imaging data can be quantitatively analyzed using automated or semi-automated software for a given imaging modality to identify additional tumor features that are difficult to identify or quantify with the naked eye, such as shape and heterogeneity (11). The tumor region of interest (ROI) is delineated by automated segmentation methods or an experienced radiologist, and the radiomics features are subsequently extracted from the ROI. These features include the shape feature, first-order feature, grayscale co-occurrence matrix feature (GLCM), grayscale region size matrix feature (GLSZM), grayscale distance region matrix feature (GLDZM), gravscale runlength matrix feature (GLRLM), neighborhood grayscale difference matrix feature (NGTDM), grayscale correlation matrix, and wavelet feature (12). The extracted radiomics features can be further screened, and the relationship between the selected features can then be analyzed (11).

In this study, we segmented the ROI of the left RLNN from the contrast-enhanced CT scans of patients with ESCC, extracted and then filtered the radiomics features of the left RLNNs, and finally included the patient's clinical data in the prediction model to predict the metastasis of the left RLNNs and analyzed its accuracy through internal verification. We present the following article in accordance with the TRIPOD reporting checklist (available at https://atm.amegroups.com/ article/view/10.21037/atm-22-5628/rc).

## **Methods**

#### Study design

The sample size of this study was initially set based on 10 events per variable (EPV). Due to the large number of radiomics features, the sample size required according to this principle is very large. However, the amount of data that patients diagnosed with ESCC who underwent radical resection of esophageal cancer in our hospital is limited. There were 511 patients with ESCC who underwent radical resection of esophageal cancer in our hospital from 2015 to 2020. Excluding patients who had received preoperative anti-tumor treatment (104 cases), combined with other malignant tumors (18 cases), and missing clinical data (39 cases), the clinical data of 350 patients were finally collected and the radiomics features were segmented and extracted. Among them, 55 patients had left RLNN metastasis. The postoperative pathology confirmed that the left RLNN metastasis rate is about 11.7%, which is consistent with the study of Kanemura et al. (13). In order to solve the problem of too many independent variables and small sample size, we used least absolute shrinkage and selection operator (LASSO) analysis and stepwise regression to preprocess the radiomics features, and finally screened out 6 radiomics features to be included in the construction of the model. Due to the small sample size, this study did not divide the training set and validation set at the beginning but used 10-fold cross-validation and bootstrap sampling verification for internal validation after the model was established.

## Patient information

The study was conducted in accordance with the Declaration of Helsinki (as revised in 2013). The study was approved by Ethics Committee-Scientific Research Subcommittee of Zhongshan Hospital Affiliated to Xiamen University (No. 2022-256). Individual consent for this retrospective analysis was waived. This study included 350 patients who were diagnosed with ESCC and underwent surgical treatment at the Zhongshan Hospital Affiliated to Xiamen University from 2015 to 2020. Of the patients, 282 were male and 68 were female. The patients were aged 36–82 years, and had an average age of 61.55±8.41 years. Further, 55 patients tested positive for metastasis to the left RLNN, and 295 patients tested



Figure 1 3D reconstruction of the left RLNN: the ROI of the left RLNN was segmented manually by an experienced clinical, and 3D reconstruction was performed. RLNN, recurrent laryngeal nerve lymph node; ROI, region of interest.

negative for metastasis to the left RLNN.

To be eligible for inclusion in this study, patients had to meet the following inclusion criteria: (I) have been diagnosed with ESCC and treated with surgery; (II) have a gold standard postoperative pathology report; (III) have undergone a contrast-enhanced CT examination within 1 month of surgery, and have image data that could be exported; and (IV) have complete clinical data. Patients were excluded from the study if they met any of the following exclusion criteria: (I) had other concurrent malignant tumors; (II) had undergone preoperative radiotherapy, chemotherapy, and other anti-tumor treatments.

In this study, age, gender, smoking history, drinking history, the contrast-enhanced CT value of the primary lesion, the contrast-enhanced CT value of the left RLNN, the difference between the contrast-enhanced CT value and the non-enhanced CT value of the left RLNN, the difference of contrast-enhanced CT value between the left RLNN and the primary lesion, tumor location, lesion length, and degree of tumor differentiation were included.

## CT image

All the patients received routine and contrast-enhanced CT of the thorax using a multi-detector CT system (Brilliance ICT; Philips Medical Systems). The scanning range was from the mandible to the costophrenic angle level. The contrast-enhanced CT scan parameters were 120 kVp,

with automatic tube current modulation, and a matrix size of 512×512. An injection of Iopamidol (300 mg/mL) was administered as a contrast medium at a dose of 70 mL and a rate of 4.0 mL/s.

# Image processing and radiomics feature extraction and screening

The chest contrast-enhanced CT scans (venous-phase scans with a slice thickness of 2.5 mm) of the patients were exported in the Digital Imaging and Communications in Medicine (DICOM) format and then imported into the 3D Slicer 4.11, an open-source radiomics extraction toolkit, and the ROI of the left RLNNs was segmented manually by 2 experienced thoracic surgeons (*Figure 1*). Next, 3D reconstruction was performed, and the radiomics plugin was used to automatically calculate and extract the radiomics of the left RLNNs, including the shape features, first-order feature, GLCM, GLSZM, GLDZM, GLRLM, and NGTDM. The extracted radiomics features were included in the LASSO regression for variable screening, and the variables corresponding to the optimal  $\lambda$  were selected.

#### Construction of the nomogram and evaluation

The radiomics features screened by LASSO-logistic regression and the variables with P values <0.05 among clinical indicators were included in the multivariate logistic

regression model, and the variables were screened using the stepwise method, after which, the nomogram for predicting the metastasis of the left RLNNs was finally established. The receiver operator characteristic (ROC) curve was used to evaluate the predictive ability of the model. The 10-fold crossvalidation was used for the internal validation of the model. The Bootstrap method was repeatedly used 2,000 times to draw the calibration curve. Harrell's C statistic was used to calculate the concordance index (C-index) to evaluate the calibration of the nomogram model.

#### Statistical analysis

SAS 9.4 was used for the statistical analysis. The counting data are expressed as the rates, and the comparisons between groups were performed using the chi-square test. The normally distributed measurement data are described as the mean plus or minus the standard deviation, and the comparisons between groups were performed by the 2 independent samples t-test. P value <0.05 (two sides) was considered statistically significant. The radiomics features were screened by LASSO-logistic regression. The variables were screened by the stepwise method, and the nomogram was constructed by the multivariate logistic regression. The R 3.6.1 software "rms" package was used to establish the nomogram. ROC curves were used to evaluate the predictive ability of the model. The 10-fold cross-validation was used for the internal validation of the model. The Bootstrap method was used to draw the calibration curve. Harrell's C statistic was used to evaluate the calibration of the model.

## **Results**

## Analysis of clinical indicators

The clinical indicators of the 350 patients were included in this study, and the results are shown in *Table 1*. According to the univariate analysis results, the enhanced CT value of the left RLNNs, the difference between the contrast-enhanced CT values between the left recurrent nerve and the primary lesion, and the tumor location were statistically significant (all P<0.05).

## Screening of radiomics features

The 3D slicer 4.11 extracted 107 radiomics features. The extracted radiomics features were included in the LASSO-

logistic regression for variable screening, and the curve of the regression coefficient changes with log ( $\lambda$ ) is shown in *Figure 2A*. The results showed that the optimal  $\lambda$  value with log ( $\lambda$ ) =-3.22 was chosen when the deviation was the smallest. The 6 variables with non-zero coefficients were selected, and the area under the curve (AUC) value was 0.884, which was a small distance from the maximum AUC, as shown in *Figure 2B*. The 6 variables identified were the least axis length, maximum 2D diameter column, maximum 2D diameter row, surface volume ratio, difference entropy, and run-length non-uniformity.

#### Analysis results of the logistic regression model

The statistically significant variables (P<0.05) among the clinical indicators (i.e., the contrast-enhanced CT value of the left RLNNs, the difference of contrast-enhanced CT value between the left RLNNs and the primary lesion, and tumor location) and the variables identified by the LASSO-logistic regression (i.e., the least axis length, maximum 2D diameter column, maximum 2D diameter row, surface volume ratio, difference entropy, and runlength non-uniformity) were included in the multivariate logistic regression model, and the stepwise method was used to screen the variables. Finally, we identified the most meaningful predictors, including the tumor location, surface volume ratio, and run-length non-uniformity. The results are shown in *Table 2*.

## Model construction and evaluation

#### Nomogram construction and use

According to the results of the multivariate logistic regression analysis, an individualized and visualized nomogram model for predicting left RLNNs metastasis in patients with ESCC was constructed. As *Figure 3* shows, the scores (points) corresponding to the individual indicators of the patient in the model were added to obtain the total score (total points), and a vertical curve was drawn to the predicted probability axis to obtain the probability of metastasis of left RLNN. The result suggests that the patient has left RLNNs metastasis when the total points are  $\geq$ 32 points [P (metastasis)  $\geq$ 0.10; the cut-off value].

## Evaluation of the model

The ROC curve was plotted, and the AUC value was used to assess model prediction performance. The AUC value of the nomogram model in this study was 0.903 [95%

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Variables	Total	Negative (n=295)	Positive (n=55)	$t/Z/\chi^2$	Р
Age, mean ± SD (years)	61.55±8.41	61.43±8.47	62.16±8.11	-0.591	0.555
Gender, n (%)					
Male	282 (80.57)	236 (80.00)	46 (83.64)	0.392	0.531
Female	68 (19.43)	59 (20.00)	9 (16.36)		
Smoking history, n (%)					
No	124 (35.43)	105 (35.59)	19 (34.55)	0.022	0.881
Yes	226 (64.57)	190 (64.41)	36 (65.45)		
Drinking history, n (%)					
No	242 (69.14)	199 (67.46)	43 (78.18)	2.499	0.114
Yes	108 (30.86)	96 (32.54)	12 (21.82)		
Contrast-enhanced CT value of primary lesion, mean $\pm$ SD	86.09±14.06	85.7±14.06	88.16±14.05	-1.191	0.234
Contrast-enhanced CT value of left RLNN, mean $\pm$ SD	37.86±30.26	35.87±32.38	48.52±8.61	-5.712	<0.001*
Difference between the contrast-enhanced CT value and the non-enhanced CT value of left RLNN, mean $\pm$ SD	25.53±16.23	26.26±16.48	21.62±14.29	1.955	0.051
Difference of contrast-enhanced CT value between left RLNN and the primary lesion, mean $\pm$ SD	48.22±30.29	49.83±32.35	39.64±11.94	4.112	<0.001*
Tumor location, n (%)				13.333	0.001*
Upper	34 (9.71)	32 (10.85)	2 (3.64)		
Middle	246 (70.29)	196 (66.44)	50 (90.91)		
Low	70 (20.00)	67 (22.71)	3 (5.45)		
Lesion length, n (%)				2.038	0.361
<3 cm	68 (19.43)	57 (19.32)	11 (20.00)		
3–5 cm	206 (58.86)	170 (57.63)	36 (65.45)		
>5 cm	76 (21.71)	68 (23.05)	8 (14.55)		
Degree of tumor differentiation, n (%)				0.581	0.748
G1	14 (4.00)	11 (3.73)	3 (5.45)		
G2	279 (79.71)	237 (80.34)	42 (76.36)		
G3	57 (16.29)	47 (15.93)	10 (18.18)		

\*, P<0.05 was considered statistically significant. RLNN, recurrent laryngeal nerve lymph node; CT, computed tomography.

confidence interval (CI): 0.861–0.945; P<0.001; Figure 4], indicating that the model had an excellent ability to predict left RLNN metastasis. The Brier score of this model was 0.074, which was <0.25, indicating that the model's predicted values were close to the actual values. The accuracy of the model was evaluated by 10-fold cross-validation, and the of are shown in *Table 3*. The validation set had an accuracy of 76.60%, a sensitivity of 85.08%, and

a specificity of 75.49%. The results of the training set and the validation set were similar, indicating that the model was stable. After the Bootstrap sampling verification of the nomogram model (n=2,000), the C-index was 0.904, indicating good calibration. The correction curve (*Figure 5*) shows that the prediction results obtained by the model were in good agreement with the actual observations, with a mean absolute error of 0.028.



**Figure 2** Selection of radiomics features by the LASSO-logistic regression. (A) The curve of the regression coefficient changes with log  $(\lambda)$ . The abscissa is log Lambda, and the ordinate is the respective radiomics coefficient. Each curve represents the trajectory of the coefficients of each radiomics feature. (B) The AUC curve versus log  $(\lambda)$ ; The red dots indicate the AUC values for each model with a given  $\lambda$ . The dotted vertical line on the left was drawn at the highest point of the AUC value, and the dotted vertical line on the right was drawn at the optimal  $\lambda$  value corresponding to the most significant value of AUC. When the AUC value was the largest, 16 features were selected, and there were too many radiomics features to apply. When the optimal  $\lambda$  value corresponding to the most significant suitable for practical applications. Finally, the optimal  $\lambda$  value with a log  $(\lambda) = -3.22$  was chosen when the deviation was the smallest. The 6 features with non-zero coefficients were selected in the plot (A) when  $\lambda$  was optimal and the value of AUC was 0.884, which was a small distance from the largest value of AUC (B). LASSO, least absolute shrinkage and selection operator; AUC, the area under the curve.

Table 2 Multivariate	logistic	regression	model	analysis	of the	left RL	NN	I metastasis
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Variables	β	Standard error	Wald $\chi^2$	Р	OR (95% CI)
Constant	-3.651	1.32	7.65	0.006	
Tumor location					
Upper	-0.341	1.367	0.062	0.803	0.711 (0.049, 10.352)
Middle	1.923	0.718	7.182	0.007	6.841 (1.676, 27.921)
Low					1.0 (reference)
Surface volume ratio	-1.084	0.498	4.732	0.03	0.338 (0.127, 0.898)
Run-length non-uniformity	0.036	0.007	27.828	<0.001	1.037 (1.023, 1.051)

RLNN, recurrent laryngeal nerve lymph node.

#### Discussion

Esophageal cancer is one of the most aggressive types of cancer, and due to the high incidence of LNM, patients have a poor prognosis. LNM is the most critical factor affecting the prognosis of esophageal cancer patients, and lymph node dissection is an integral part of radical surgery for esophageal cancer. However, the use of lymph node dissection in esophageal cancer is still controversial. According to a meta-analysis, a 3-field lymphadenectomy is associated with a significantly improved 5-year OS, but no significant differences between 3-field lymphadenectomy and 2-field lymphadenectomy were found in terms of postoperative mortality, anastomotic leak, pulmonary complications, chylothorax, and recurrent nerve palsy (14). Conversely, Koterazawa's meta-analysis showed that OS did not differ significantly between the 3- and 2-field lymphadenectomy groups, but the incidence of RLN palsy was significantly higher in the 3-field lymphadenectomy than the 2-field lymphadenectomy group (15).

Regional lymph node stations of metastasis are most frequently observed in the RLNN; however, the rates of metastasis to the to the left and right RLNN are similar. The left RLNNs, whose anatomical pathway is longer that its



**Figure 3** Nomogram for predicting the left RLNN metastasis. The point axis represents the score of each predictor in the plot at different values, and the total points axis represents the sum of the scores of all the predictors. We found the value of the patient's indicator on each variable axis, drew a vertical line at this position to the point axis, and obtained the score of the indicator, then added the scores (point) corresponding to the individual indicators of the patient in the model to obtain the total score (total points), and drew the vertical curve from the Total point axis to the probability of metastatic axis to get the probability of left RLNN metastasis. It suggests that the patient has left RLNN metastasis when the total points are  $\geq$ 32 points [P (metastasis)  $\geq$ 0.10, the cut-off value]. RLNN, recurrent laryngeal nerve lymph node.



**Figure 4** The ROC curve of the logistic regression model predicting left RLNN metastasis. The abscissa is 1 – specificity, and the ordinate is sensitivity, AUC =0.903, 95% CI: 0.861–0.945. ROC, receiver operator characteristic curve; RLNN, recurrent laryngeal nerve lymph node; AUC, the area under the curve.

Table 3 Result	s of 10-fold	cross-validation
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	Correct rate, %	Sensitivity, %	Specificity, %
Training set	77.74	88.08	75.81
Validation set	76.60	85.08	75.49



**Figure 5** Calibration curve of the nomogram for predicting left RLNN metastasis. The abscissa coordinate represents the predicted probability of left recurrent laryngeal LNM in the nomogram, the ordinate indicates the probability of actual metastasis of left RLNN, the diagonal line represents the perfect prediction curve of the ideal model, the dotted line represents the prediction curve of this nomogram, and the solid line represents the bias-corrected prediction curve. RLNN, recurrent laryngeal nerve lymph node; LNM, lymph node metastasis.

right counterpart, runs over a broad region, along the RLN, from the upper mediastinum to the neck, and thus can be more difficult to remove surgically. Moreover, it is located in front of the trachea, which increases the difficulty of the right thoracotomy operation (13). Additionally, left RLNN

adjacent to the aorta and a left RLN lymphadenectomy also increase postoperative respiratory complications (including dyspnea, pneumonia, aspiration, and acute respiratory distress syndrome), increase the incidence of esophagogastric anastomotic fistulas, and prolong postoperative hospital stays. Left RLN palsy is more common than right RLN because of its longer path and deeper location (16). According to Pai's study of patients with clinical positive RLN lymph node involvement before treatment, there were significant differences in the OS and disease-free survival (DFS) between patients with and without RLN lymphadenectomy; however, no significant differences were found in the 3-year OS and DFS between patients with and without RLN lymphadenectomy in the entire cohort or the neoadjuvant chemoradiotherapy group (17).

Preoperative examinations of the left RLNN metastasis determine the choice of lymph node dissection in patients with esophageal cancer. Thus, predicting the left RLNNs metastasis using a non-invasive method would enable clinicians to formulate individualized lymph node dissection plans for patients with ESCC before surgery. This study sought to establish a prediction model for left RLNN metastasis in ESCC to guide clinicians in formulating personalized and precise lymph node dissection plans for patients with esophageal cancer before surgery.

Radiomics focuses on improving image analysis, using an automated high-throughput extraction of large amounts of quantitative features of medical images, and is a recent innovation in medical imaging analysis. Radiomics, as the advanced image analysis of novel medical imaging, can capture additional information not currently used, and more specifically, genomics and proteomic patterns that can be expressed in terms of macroscopic image-based features (9). Thus, this study constructed a predictive model for left RLNN metastasis in ESCC patients based on radiomics and clinical data.

Relevant study has shown that LNM of esophageal cancer is associated with the tumor location, lesion length, depth of invasion, degree of differentiation, and the presence of vascular tumor thrombus (18). A previous study showed that PIM-1, a serine/threonine protein kinases of the PIM (Proviral Integration of Moloney virus) family, is a promising molecular marker for predicting LNM in early esophagus cancer (19). However, as only 67 samples were included in that research, and a single data source was used, the study lacked generalisability. As clinical data for molecular marker was lacking at our research center, it was not included in the univariate analysis of the clinical data

in this study. In clinical practice, the short diameter of the lymph node  $\geq$ 1.0 cm is often used as the diagnostic criterion for positive LNM, but the clinical diagnostic criterion for the short diameter of metastatic lymph nodes is still controversial (20). Additionally, as lymph node enlargement cannot exclude other possibilities, such as inflammatory cell infiltration and tissue hyperplasia, this indicator was not included in the univariate analysis of the clinical data in this study.

The following preoperative clinical indicators were included in this study: age, gender, smoking history, drinking history, the contrast-enhanced CT value of the primary lesion, the contrast-enhanced CT value of the left RLNNs, the difference between the contrast-enhanced CT value and the non-enhanced CT value of the left RLNNs, the difference between the contrast-enhanced CT value between the left RLNNs and the primary lesion, tumor location, lesion length, and degree of tumor differentiation. According to the univariate analysis results, the contrastenhanced CT value of the left RLNNs, the difference of contrast-enhanced CT value between the left recurrent nerve and the primary lesion, and the tumor location were statistically significant (all P<0.05).

LASSO is a widely used data screening method that shrinks the coefficients and changes some coefficients to 0, and LASSO can be used for feature reduction and selection (21). Studies have shown that the method is suitable for processing large numbers of radiomics with relatively small sample sizes and avoiding overfitting (22,23).

The AUC of the prediction model reached 0.903 (95% CI: 0.861–0.945), indicating that the prediction model had excellent prediction performance. Further, 10-fold cross-validation was used to verify the internal validation of the model. The results of the training set and the validation set were similar, indicating that the model was stable. The Bootstrap method was used to draw the calibration curve, and the nomogram model was internally verified. Harrell's C statistic was used to calculate the C-index to evaluate the calibration of the nomogram model, and the C-index for predicting malignancy was 0.904, indicating that the model had a good degree of calibration, and the prediction results obtained by the model were close to the actual observations.

This study had a number of limitations. First, the data in this study were all obtained from the Zhongshan Hospital Affiliated to Xiamen University, and the sample size was not sufficiently large. In future research, it is hoped that data from multicenter sources can be collected, and the external verification of the model can be further improved.

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Second, the elevation of some tumor markers, such as abnormal tumor protein PMI-1, is associated with LNM in esophageal cancer. However, as the study lacked complete clinical data for tumor markers, some tumor markers were not included in the univariate analysis. The inclusion of specific tumor markers in subsequent studies may increase the accurate predictive ability and clinical application value of the constructed predictive models. Third, in the process of radiomics extraction in this study, the ROIs of the left RLNNs on chest contrast-enhanced CT scans in patients with ESCC were manually outlined, which took a great deal of time and effort and were affected by some fatty tissues next to the lymph nodes. Further developments need to be made in medical imaging techniques and the accuracy and precision of lymph node segmentation needs to be improved. Thus, in future research, we hope to obtain data from multicenter sources, increase the sample size, include more meaningful clinical indicators, and extract radiomics features in a more scientific and effective way to improve the predictive ability and clinical application value of the predictive model.

## Conclusions

A visual nomogram model combining clinical data and radiomics features was established to predict left RLNN metastasis in patients with ESCC in a non-invasive way. This model can provide a reference for clinicians to formulate individualized lymph node dissection plans for patients with ESCC. According to the ROC curve, the AUC of the model was 0.903, which means that the model had an excellent predictive ability. Combined with the internal verification results, our results indicated that the predictive model was stable and had a high degree of calibration.

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## Footnote

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*Ethical Statement:* The authors are accountable for all aspects of the work in ensuring that questions related to the accuracy or integrity of any part of the work are appropriately investigated and resolved. The study was conducted in accordance with the Declaration of Helsinki (as revised in 2013). The study was approved by Ethics Committee-Scientific Research Subcommittee of Zhongshan Hospital Affiliated to Xiamen University (No. 2022-256). Individual consent for this retrospective analysis was waived.

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