

## Research Article

# Radiomics Approach for Predicting Esophageal Cancer Pathological Stages Based on MRI

Rihui Yang<sup>1#</sup>; Ting Dong<sup>2#</sup>; Tianhui Zhang<sup>1</sup>; Weixiong Fan<sup>1</sup>; Haodong Qin<sup>3</sup>; Guihua Jiang<sup>2\*</sup>; Haiyang Dai<sup>4\*</sup>

<sup>1</sup>Department of Radiology, Meizhou People's Hospital, PR China

<sup>2</sup>Department of Medical Imaging, Guangdong Second Province General Hospital, PR China

<sup>3</sup>Siemens Healthineers, Guangzhou, PR China

<sup>4</sup>Department of Medical Imaging, Huizhou Municipal Central Hospital, PR China

**\*Corresponding author: Haiyang Dai**

Department of Medical Imaging, Huizhou Municipal Central Hospital, No. 41, North Eling Road, Huizhou 516001, PR China;

Guihua Jiang, Department of Medical Imaging, Guangdong Second Province General Hospital, Guangzhou, 510317, China.

Email: d.ocean@163.com; jianggh@gd2h.org.cn

<sup>#</sup>These authors have been equally contributed to this article.

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## Introduction

Esophageal Cancer (EC) is one of the most common malignant gastrointestinal tumors in China, ranking sixth in malignant tumor incidence and fourth in mortality [1], with a five-year survival rate of 19% [2]. Early (stage I - II) esophageal cancer is primarily treated with minimally invasive endoscopic resection or radical surgical resection, while advanced (stage III - IV) esophageal cancer, which has a poor prognosis, is managed with surgical resection after neoadjuvant chemoradiotherapy,

## Abstract

**Objective:** To explore a rational radiomic approach for predicting preoperative staging and lymph node metastasis based on Magnetic Resonance (MR) images in esophageal cancer patients.

**Materials and Methods:** This retrospective study included 120 patients (primary cohort: n= 84; validation cohort: n= 36) with esophageal carcinoma confirmed by surgery and pathology. All patients underwent a preoperative MR scan from the neck to the abdomen. For each patient, high throughput and quantitative radiomics features were extracted from T2WI and contrast-enhanced T1WI (T1WI\_Gd) images. The radiomics signature of the T2WI and T1WI\_Gd images was constructed using minimal redundancy maximal relevance (mRMR) and the least absolute shrinkage and selection operator (Lasso). In addition, associations between the radiomics signature and esophageal cancer staging and lymph node metastasis were explored. Finally, the diagnostic performance of the radiomics approach and tumor volume for discriminating between stages I - II and III - IV and predicting lymph node metastasis was evaluated and compared using the area under the receiver operating characteristic curve (AUC), Sensitivity (SEN), and Specificity (SPE).

**Results:** A total of 1316 radiomics features were extracted. After feature dimension reduction, eight and nine features were selected for the respective cohorts to build radiomics signatures. Then, based on these signatures, a logistic regression model was built to predict the stages and lymph node metastasis of esophageal cancer. The radiomics signature of T2WI with T1WI\_Gd discriminated better the stages and lymph node metastasis in the primary (AUC: 0.884, 0.858; SEN: 0.953, 0.744; SPE: 0.732, 0.829) and validation cohorts (AUC: 0.765, 0.711; SEN: 0.842, 0.952; SPE: 0.647, 0.533).

**Conclusions:** The MRI-based radiomics signature could identify esophageal cancer stages and lymph node metastasis before treatment.

**Keywords:** Esophageal cancer; Tumor staging; Lymph node metastasis; Multimodal MRI

or chemoradiotherapy alone [3]. Notably, Lymph Node (LN) involvement is generally associated with worse overall survival [4]. Therefore, predicting EC stages and LN metastasis in patients before treatment is clinically essential. Radiomics is gaining importance in cancer research [5]. High-throughput mining extracts quantitative image features from digitally encrypted medical images, and this is coupled with robust image-based signatures that could potentially enhance precision diagnosis

and treatment. Radiomics research recently revealed MRI’s potential to substantially improve the ability to detect or predict LN metastases [4,5]. However, there are no reports on whether a radiomics approach could predict the EC stage based on multimodal MRI.

Computed Tomography (CT), Magnetic Resonance Imaging (MRI), Positron Emission Tomography-CT (PET-CT), and Endoscopic Ultrasonography (EUS) have their respective advantages and disadvantages for evaluating EC preoperative stages and LN status; notably, their sensitivities and specificities are different [6-8]. Currently, the CT-enhanced scan is the most commonly used imaging method for EC diagnosis and efficacy evaluation. However, the detection accuracy of positive LN on preoperative CT remains controversial, and the reported sensitivity, specificity, and accuracy was 37.3-67.2%, 63.9-96.4%, and 85.8-87.2%, respectively [9]. Moreover, the latest version of the Chinese guidelines for EC radiation therapy proposed that the diagnostic value of MRI for LN metastasis was similar to or better than enhanced CT [10].

Relevant studies have shown that radiomics has potential value in predicting tumor stages or LN metastasis [11-13]. However to date, few studies have reported on building models based on MRI radiomics features to predict EC stages and LN metastasis. In this study, the preoperative T2WI and contrast-enhanced T1WI (T1WI\_Gd) sequence radiomics features were extracted. We aimed to construct a model to evaluate their value in predicting EC stages and LN metastasis to provide a reference for the clinical treatment of patients.

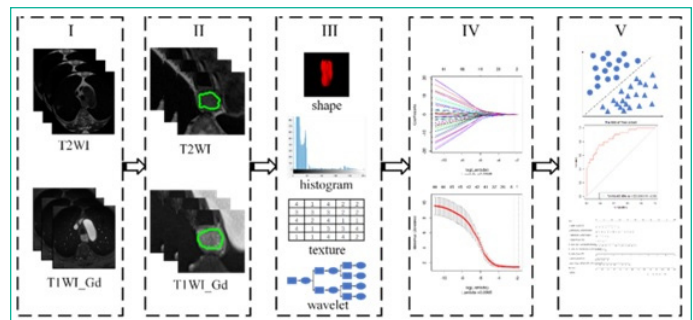
**Result**

**Clinical Data**

The clinicopathologic characteristics of the patients are presented in Table 1. There were no significant differences in age, sex, tumor location, or pathological type between the two groups of patients at EC stages I - II and III – IV, or between the two groups of EC patients with positive and negative LN metastasis ( $P > 0.05$ ).

**Table 1:** Characteristics of patients for TNM stages and LN status.

Characteristics	TNM stages [n (%)]		P Value	LN status [n (%)]		P Value
	I-II stage	III-IV stage		LN (+)	LN (-)	
Age ( $\bar{x} \pm s$ ) (year)	63.4±8.4	63.5±7.8	0.949	63.8±7.8	63.1±8.6	0.64
Gender			0.094			0.14
Male	39 (67.2)	50 (80.6)		51 (79.7)	38 (67.9)	
Female	19 (32.8)	12 (19.4)		13 (20.3)	18 (32.1)	
Location			0.597			0.64
Upper	4 (6.9)	3 (4.8)		3 (4.7)	4 (7.2)	
Middle	33 (56.9)	31 (50.0)		32 (50.0)	32 (57.1)	
Lower	20 (34.5)	22 (35.5)		23 (35.9)	19 (33.9)	
Middle and lower	1 (1.7)	6 (9.7)		6 (9.4)	1 (1.8)	
Pathological type			0.573			0.39
Squamous cell carcinoma	55 (94.8)	61 (98.4)		63 (98.4)	53(94.6)	
Adenocarcinoma	1 (1.7)	0 (0)		0 (0)	1 (1.8)	
Mixed carcinoma	2 (3.5)	1 (1.6)		1 (1.6)	2 (3.6)	



**Figure 1:** Flowchart of this study: (I) Original MR images: T2WI and T1WI\_Gd. (II) Segmentation was performed on both T2WI and T1WI\_Gd images to define the tumor region. (III) Radiomic features were extracted from the tumor region, including shape, first-order histogram, texture and wavelet group analysis. (IV) Several features were selected to build the radiomic signature using the minimal redundancy maximal relevance (mRMR) and the Least Absolute Shrinkage and Selection Operator (LASSO) method. (V) Finally, the classification ability of the radiomic signature was tested by the Receiver-Operating Characteristics (ROC) curves in both the primary and validation cohort.

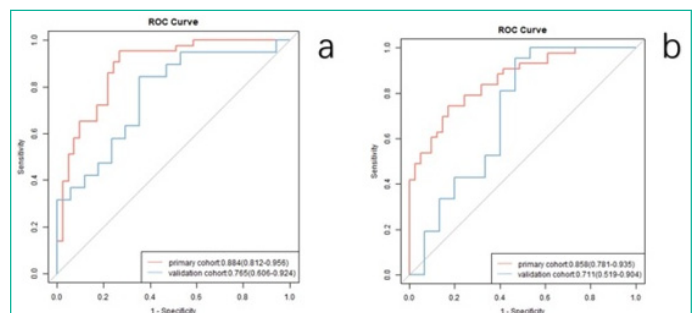
**Radiomic Features**

We used the variance method (excluding variance as 0), mRMR method (select the top 50), and LASSO method (lambda selection standard  $\lambda_{min}$ ) for dimensionality reduction. As a result, eight and nine of 1316 radiomic features were included for EC staging and LN metastasis prediction, respectively. For EC staging, these included logarithm\_glcm\_Maximum Probability, logarithm\_firstorder\_Interquartile Range, exponential\_glszm\_Gray Level Non-Uniformity Normalized, wavelet. HHL\_glszm\_Size Zone Non-Uniformity Normalized, gradient\_glszm\_Low Gray Level Zone Emphasis, logarithm\_firstorder\_Skewness, logarithm\_ngtdm\_Contrast, logarithm\_glcm\_Idn. For LN metastasis prediction, these were the included wavelet. LLH\_glcm\_MCC, wavelet. HHL\_glcm\_MCC, logarithm\_glcm\_Maximum Probability, wavelet. HHL\_glcm\_Imc2, exponential\_glszm\_Gray Level Non-Uniformity Normalize, gradient\_glszm\_Low Gray Level Zone Emphasis, logarithm\_ngtdm\_Contrast, gradient\_gldm\_Dependence Non-Uniformity Normalized, and logarithm\_gldm\_Dependence Variance.

The names and descriptions of the selected features are listed in Table 2. Distribution of eight and nine features that could distinguish between stages I - II and III - IV, and positive and negative LN metastasis, were analyzed using the  $\chi^2$ -test, respectively. Of these, the feature “wavelet.HHL\_glcm\_Imc2” had a  $P$  value  $< 0.05$ .

**Radiomic Signature Discrimination**

The radiomic signature was built by employing the selected features in the previous section, forming a linear combination



**Figure 2:** ROCs were employed to assess the MRI radiomic signature discriminative performance of the TNM stages and the LN metastasis in preoperative EC patients.

**Table 2:** Selected features with descriptions.

Task	MRI	Feature Name	OR (95%CI)	P Value
TNM stages prediction	T2WI	logarithm_glcm_MaximumProbability	1.32(0.64~1.86)	0.251
		logarithm_firstorder_InterquartileRange	0.86(0.39~1.36)	0.054
	T1WI-Gd	exponential_glszm_GrayLevelNonUniformityNormalized	1.16(0.84~2.54)	0.654
		wavelet.HHL_glszm_SizeZoneNonUniformityNormalized	0.63(0.74~0.86)	0.075
		gradient_glszm_LowGrayLevelZoneEmphasis	0.95(0.55~1.34)	0.274
		logarithm_firstorder_Skewness	0.76(0.63~2.16)	0.064
		logarithm_ngtdm_Contrast	1.75(0.81~3.68)	0.312
logarithm_glcm_Idn	0.72(0.33~0.93)	0.476		
LN status prediction	T2WI	wavelet.LLH_glcm_MCC	0.92(0.55~1.53)	0.765
		wavelet.HHL_glcm_MCC	0.55(0.19~1.44)	0.231
		logarithm_glcm_MaximumProbability	1.52(0.84~2.86)	0.175
		wavelet.HHL_glcm_Imc2	0.5(0.24~0.93)	0.045*
	T1WI-Gd	exponential_glszm_GrayLevelNonUniformityNormalize	0.96(0.54~1.66)	0.879
		gradient_glszm_LowGrayLevelZoneEmphasis	0.71(0.26~1.76)	0.474
		logarithm_ngtdm_Contrast	1.35(0.61~3.07)	0.467
		gradient_gldm_DependenceNonUniformityNormalized	0.56(0.29~1.06)	0.084
		logarithm_gldm_DependenceVariance	0.62(0.25~1.43)	0.276

of the logistic regression model of these features. The radiomic signature's discriminative power for TNM staging and LN metastasis was assessed by two ROCs in the primary and validation cohorts (Figure 2), respectively. Then, the model's ROCs were tested by DeLong, and the AUC, SEN, and SPE were calculated (Table 3). The radiomic scores (Rad scores) of the EC patients in the primary and validation cohorts were calculated through the elastic net model. Each patient's Rad scores in both the primary and validation cohorts for two tasks.

## Discussion

This study showed that a quantitative approach based on multimodal MRI could identify esophageal cancer stages and lymph node metastasis before treatment initiation. Based on significant differences in the Rad-score, the radiomics signature successfully differentiated between ESCC stages I - II and III - IV before the preoperative period. In addition, by comparing the radiomics characteristics and those of the combined models, our results showed that MRI-based radiomics could predict lymph node metastasis of esophageal cancer before the preoperative period. These observations were based on MRI's excellent soft tissue resolution and multimodal imaging, which can

provide a high-quality image basis for accurate tumor delineation and radiomic analysis.

Surgery is the most essential treatment for esophageal cancer. However, operating on advanced esophageal cancer (III ~ IV stages) is challenging, and the recurrence rate is high after surgical treatment. The lymph node status in esophageal cancer is a significant independent predictor of prognosis, and the method used for lymph node dissection is crucial for esophageal cancer treatment [4]. Some studies have suggested that systematic lymph node dissection in esophageal cancer patients should be conducted only when the tumor invades the submucosa and that extended lymph node resection may increase postoperative complications [14-15]. In addition, lymph node staging is also crucial for implementing neoadjuvant therapy [16]. Therefore, it is essential to accurately predict the stage and lymph node metastasis in esophageal cancer patients before surgery to allow the formulation of a treatment plan and prognostic evaluation. CT is a common imaging method for preoperative esophageal cancer staging, but its effect on staging is poor. By extracting high-throughput features from medical images, radiomics can reveal tumor heterogeneity and biological behavior and improve the accuracy of tumor diagnosis and patient prognosis.

This study demonstrated the feasibility of predicting esophageal cancer's stage and lymph node metastasis with MR radiomic characteristics. We used T2WI and T1WI-Gd to extract 1316 quantitative image features from tumors. Eight features that contributed most to the differentiation of EC stages and nine that contributed most to the differentiation of lymph node metastasis were selected to construct the models. T2WI radiomics features are based on the extraction of water molecule signal strength, which primarily reflects the characteristics of the water content of the tumor.

The obtained high-order radiomics feature is mainly the gray level co-occurrence matrix, which reflects the gray level difference, including the gray level distribution uniformity and texture thickness. The more uneven the gray level and the more complex the texture, the greater the heterogeneity of the tumor. The histological features of the T1WI-Gd images were obtained based on the solid part of tumor enhancement and most reflected the tumor's microvessel density, histological grading, and invasiveness [17]. The extracted higher-order features are more diverse, including the degree of difference between adjacent pixels, a gray area matrix, and dependence on the advantages of high gray, which can better reflect the heterogeneity of tumors in multiple dimensions [18].

As in other studies [19-21], we combined the radiomics characteristics of two sequence images to distinguish the esophageal cancer stages I - II and III - IV and lymph node metastasis and achieved satisfactory results (AUC = 0.884, 0.858 in the main cohort, AUC = 0.765, 0.711 in the verification cohort). Compared with previous studies on esophageal cancer tumor staging and lymph node metastasis, the prediction efficiency of the radiologic features as we built was better than that of the CT-based radiomics methods used for predicting esophageal cancer stages I - II and III - IV (AUC = 0.795 in the main cohort, AUC = 0.694 in the validation cohort) [22]. These could be because the model built by combining the radiomics features of the MR sequence images can mutually complement the image information and be more comprehensive, objectively reflecting tumor heterogeneity. In addition, our results were similar to those of other studies on the prediction of lymph node me-



**Table 3:** The performance of model for predicting TNM stages and LN metastasis in EC.

Task	Primary cohort			Validation cohorts		
	AUC (95% CI)	SEN (95% CI)	SPE (95% CI)	AUC (95% CI)	SEN (95% CI)	SPE (95% CI)
TNM stages prediction	0.884 (0.812-0.956)	0.953 (0.891-1.000)	0.732 (0.596-0.867)	0.765 (0.606-0.924)	0.842 (0.678-1.000)	0.647 (0.420-0.874)
LN status prediction	0.858 (0.781-0.935)	0.744 (0.614-0.875)	0.829 (0.714-0.944)	0.711 (0.519-0.904)	0.952 (0.861-1.000)	0.533 (0.281-0.786)

tastasis of esophageal cancer based on MR radiomics characteristics (AUC = 0.821 in the main cohort, AUC = 0.762 in the validation cohort) [19], further confirming the feasibility of MR imaging characteristics to predict lymph node metastasis of esophageal cancer.

This study had several limitations. First, the sample size was relatively small, and our results should be further verified with a multicenter, large-sample study. Second, we only discussed whether the MRI radiomics could predict lymph node metastasis of esophageal cancer. However, the prediction efficiency of lymph node metastasis and its extent in different regions has not been studied. Again, a prospective, large-sample, multicenter prospective study is needed to verify our results. Last, the images in this study were all produced using the same MRI scanner. In a future study, we plan to discuss the influence of different types of equipment, field strength, and scanning parameters on the radiomics model.

In conclusion, this study explored the feasibility of using MR image-based radiomics methods to identify esophageal cancer stages I - II and III - IV and lymph node metastasis before treatment. We constructed a multi-feature radiomics model from the extracted radiomics features. This model has shown satisfactory performance in identifying esophageal cancer stages I - II and III - IV and lymph node metastasis early in surgery. With the popularization of magnetic resonance examination, we believe that MR-based radiomics might be a noninvasive and quantitative method for predicting the preoperative staging of esophageal cancer and guiding individual treatment decisions.

## Materials and Methods

### Patients

The clinical and MRI data of EC patients underwent lymph node dissection confirmed by surgery and pathology from January to December 2020 were retrospectively collected. The inclusion criteria were: ① The postoperative pathology was confirmed to be EC; ② Radical resection of esophageal cancer and lymph node dissection, with clear postoperative pathological stage and clear pathological lymph node results; ③ MRI scan (T2WI and T1WI-Gd) was performed within two weeks before surgery; ④ No EC treatment before MRI scan; and ⑤ No previous history of other malignant tumors. The exclusion criteria were: ① Preoperative EC targeted for treatment; ② EC associated with other malignant tumors, and ③ Poor MRI image quality, affecting observation and subsequent feature extraction. This study was approved by the Ethics Committee on Clinical Research and Novel Technologies of Meizhou People's Hospital, and patient informed consent was waived for this retrospective study. All experiments were performed in accordance with relevant guidelines and regulations.

Ultimately, 120 patients with EC (89 males and 31 females, ages 44 to 82 years old, mean age 63.4±8.17 years) were included. Based on TNM staging, 58 patients were EC stages I - II (39 males, 19 females, mean age 63.4±8.4 years old), and 62 were EC stages III - IV (50 males and 12 females, mean age 63.5±7.8 years old). According to the postoperative lymph node pathology results, 56 patients were negative (38 males, 18 females,

mean age 63.1±8.6 years old), and 64 patients were positive (51 males and 13 females, mean age 63.8±7.8 years old). The clinicopathologic characteristics of the patients are presented in Table 1.

### Image Acquisition

Imaging was performed with a 3.0T Siemens (Erlangen, Germany) Skyra MR scanner equipped with 18-channel phased array coil. The patient was placed supine and told to breathe as calmly as possible to reduce motion artifacts. The scanning range was from the level of the supraclavicular fossa to the gastric cardia. The scanning sequence and parameters were: (1) T2WI: slice thickness 3 mm, TR 3000 ms, TE 91 ms, NEX=1, matrix 256×256, field of view (FOV) 210 mm×210 mm and scanning time 2 min; and (2) T1WI-Gd: slice thickness 1.5 mm, TR 4.4 ms, TE 2.0 ms, NEX=1, matrix 160×160, FOV 150 mm×150 mm, and flip angle 12. During enhanced scanning, Gd-DTPA contrast agent was injected intravenously through the elbow via a high-pressure syringe at a 0.1 mmol/kg dose. The scanning time was 3 minutes 26 seconds.

### Tumor Segmentation

Manual segmentation of the EC was performed on each patient's MR images utilizing an A.K. (Artificial Intelligent Kit), a new application platform for the overall solution of artificial intelligence (AI) in medical imaging radiomics launched by GE Healthcare. Two experienced associate senior radiologists (readers 1 and 2) carefully contoured the tumor on all T2WI and T1WI-Gd images to generate two 3D segmentations of the entire tumor. These regions of interest (ROIs) were performed for subsequent feature extraction and further analysis.

### Radiomic Feature Extraction

Patients were randomly divided into primary (84 patients) and validation (36 patients) cohorts. Radiomic features were calculated based on the segmentation results from the primary cohort using a homemade program in the A.K. For each MRI sequence, the 1316 extracted features covered the major feature pool in recent radiomic studies. The features were categorized into six primary types: (1) histogram features; (2) form factor features; (3) Gray Level Co-occurrence Matrix (GLCM) and Harilick features; (4) Gray Level Run Length Matrix (GLRLM) features; (5) Grey Level Size Zone Matrix (GLSZM) features; and (6) gray level dependence matrix (GLDM) features (Figure 1).

The appropriate feature selection procedure was used to simplify the building model and avoid over-fitting issues. In addition, the minimal redundancy maximal relevance (mRMR) and Least Absolute Shrinkage and Selection Operator (LASSO) method were used for feature dimensionality reduction. For tuning coefficients  $\lambda$  and  $\alpha$ , the respective minimum standard deviation and maximum AUC criteria were followed. For accurate prediction of the EC stages, eight features selected from the T2WI with T1WI-Gd images made the greatest contribution. On the other hand, nine features selected from T2WI with T1WI-Gd images were most useful for precise detection of LN metastasis. Then, each patient's two radiomic signature sets were inputted into the logistic regression model. After the training in the pri-

mary cohort, a linear combination of the selected features was extracted. The sums of those linear combinations formed the radiomic signatures of each patient.

### Statistics Analysis

The statistical analysis was performed in R (version 3.5.1; <http://www.Rproject.org>). The Shapiro-Wilk test was used to test whether the quantitative data were normally distributed. The  $\chi^2$ -test was used to compare the differences in age, sex, tumor location, and pathological type between patients with EC stages I - II and stages III - IV and positive and negative LN metastasis. In addition, the primary cohort was used to train the logistic regression model, and the validation cohort was used to test the model. Finally, the Receiver operating Characteristic Curve (ROC), Sensitivity (SEN), and Specificity (SPE) were used to evaluate the performance of the prediction model, and the DeLong test was used to measure the significance of the Area Under the Curve (AUC) between the single-sequence and mixed-sequence models.  $P < 0.05$  indicated statistical significance.

### Author Statements

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#### Author Contributions

All authors participated in the study design, interpretation of the data, and critically reviewing the paper. Guihua Jiang put forward the study concepts and Rihui Yang wrote the first draft, and contributed to the writing of subsequent versions. Data was acquired and prepared for analysis by Ting Dong and Tianhui Zhang. Statistical analyses and preparation of tables and figures was performed by Weixiong Fan and Haodong Qin. Haiyang Dai was responsible for project administration and supervision.

#### Data Availability Statement

All data generated or analyzed during this study are included in this article.

#### Conflict of Interest statement

The authors have no relevant conflicts of interest to disclose.

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