



# Integrated radiomic model for predicting the prognosis of esophageal squamous cell carcinoma patients undergoing neoadjuvant chemoradiation

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**Background:** To establish a feasible prediction model for prognoses of esophageal squamous cell carcinoma (ESCC) patients undergoing neoadjuvant concomitant chemoradiation (NACCRT).

**Methods:** Post-chemoradiation computed tomography (CT) radiomics features and clinical parameters were investigated. CT images from advanced thoracic ESCC patients treated with NACCRT and esophagectomy were extracted for radiomics features. Least absolute shrinkage and selection operator regression were used to select features and build signatures. Radiomics signatures and clinical factors were integrated into Cox regression analysis for prognosis; the prediction model's performance was examined via receiver-operating characteristic (ROC) curve analysis.

**Results:** A total of 46 radiomics features and 25 clinical parameters were extracted from 62 cases, of which 59 passed image processing and became eligible for model testing. Eight selected radiomics features showed good prediction power [area under the curve (AUC) =0.851] and reliability in predicting pathological complete response (pCR). The radiomics signature and clinical parameter combination model showed increased prediction power of radiomics signature alone for local regional failure (LRF) (AUC=0.804) and distant failure (DF) (AUC=0.754). Following were the strongest contributors of prediction power for prognostic endpoints: (I) resection status multiplied by long-run emphasis in grey-level run length matrix (GLRLM\_LRE) for progression (hazard ratio=8.776); (II) non-uniformity of the grey-levels (GLRLM\_GLNU) (hazard ratio=6.888); and (III) sphericity (hazard ratio=0.152) for overall survival (OS).

**Conclusions:** The integrated prediction model for prognosis may aid clinicians in decision making regarding post-operative adjuvant therapy for ESCC patients undergoing NACCRT.

**Keywords:** Esophageal cancer; neoadjuvant concurrent chemoradiation; radiomics; prediction model

Received: 12 June 2019; Accepted: 22 July 2019; Published: 12 August 2019.

doi: 10.21037/tro.2019.07.03

View this article at: <http://dx.doi.org/10.21037/tro.2019.07.03>

## Introduction

Esophageal cancer is the fifth leading cause of male cancer mortality in Taiwan, accounting for approximately 1,600 deaths annually (1). In total, 72% of patients are

diagnosed at the advance stage. The most common type of esophageal cancer in Asia is differentiated esophageal squamous cell carcinoma (ESCC), presenting different etiology pattern in the occidental world (2). Neoadjuvant

concomitant chemoradiation (NACCRT) effectively increases the feasibility of complete surgical resection and thus, has become the standard treatment modality for advanced esophageal cancer in the past 20 years (3,4). Clinical studies on NACCRT for esophageal cancer have reported that pathological complete response (pCR) predicts better disease control and survival outcomes (5). Meanwhile, poor responders show poor prognosis despite adjuvant treatment for disease control (6). However, the decision-making factors for the post-operative adjuvant therapy after NACCRT for ESCC remain not well defined. Undoubtedly, predictive models of response to NACCRT can provide crucial information before surgery.

Computed tomography (CT) is widely used in post-NACCRT surveillance of esophageal cancer. However, it has limitations in terms of distinguishing the residual tumor from the post-treatment effect via visual interpretation (7). Interpreting clinical images using a combination of radiomics and machine learning is a new field in artificial intelligence. Radiomics has been proven to be useful in cancer staging and clinical evaluation (8,9). Texture-based radiomics features are potential robust predictors of clinical outcomes of various cancers (10,11).

For establishing a prediction model of response and prognosis in patients undergoing chemoradiation, there are 2 ways of choosing the objects for extracting radiomics features. One way involves the use of pre-treatment images, which have already been used in numerous studies in recent years. Pre-treatment images are closely correlated to individual differences, helpful in identifying high-risk patient groups, and often used in establishing constrain models following chemoradiation (12-15). The other way involves using the image series taken after chemoradiation. Post-treatment images more closely reflect the treatment effect relationship, such as immune reaction and alteration of the tumor microenvironment. However, high rates of false-positive signals are the major technical problem limiting the application of these images. Radiomics becomes valuable in such situations.

The purpose of this study was to establish a prediction model of pathological response, failure pattern, and survival in advance ESCC patients by calculating texture-based radiomics features of CT images after NACCRT. We hypothesized that radiomics would extract more information from post-chemoradiation CT images and thus, could reduce the treatment effect and focus on the tumor-related factors influencing prognosis.

## Methods

### *Patient demographic data and workup*

This retrospective, single-center study evaluated patients with locally advanced esophageal cancer who underwent NACCRT followed by total esophagectomy and gastric tube reconstruction between 2013 and 2018. Those with Eastern Cooperative Oncology Group (ECOG) Performance Status score >1 or biopsy-proven differentiated adenocarcinoma were excluded. Response and outcome were evaluated in 62 patients. Cancer staging was performed using CT, endoscopic ultrasound, esophagogastroduodenoscopy, and whole body 2-deoxy-2-[fluorine-18]-fluoro-D-glucose positron emission tomography integrated with CT (FDG PET/CT). Post-treatment surveillance was assessed 5–8 weeks after completion of NACCRT by performing CT and whole body FDG-PET/CT (optional). Staging was performed according to the TNM classification system (seventh edition) of the American Joint Committee on Cancer. Grade of toxicity was evaluated according to the National Cancer Institute's Common Terminology Criteria for Adverse Events version four.

### *Radiotherapy delivery and chemotherapy administration*

All eligible patients underwent cancer treatment according to the standard operating procedures at our institution. Supine, contrast-enhanced, alpha cradle® kits (Smithers Medical Products) immobilized CT images were performed for simulation. Simulation images included the entire neck, thorax, and upper abdomen; CT was performed with 3-mm slice thickness and OMAR reconstruction (optional) to avoid metallic shadows. Target volume delineation was performed on the Eclipse platform according to the criteria described below:

- ❖ Gross tumor volume (GTV): primary tumor and lymphadenopathy observed on CT and/or <sup>18</sup>F-FDG-PET/CT.
- ❖ High-dose clinical target volume (CTV<sub>High</sub>): entire GTV with the margins extended 1 cm circumferentially and 3–4 cm longitudinally.
- ❖ Low-dose clinical target volume (CTV<sub>Low</sub>): area of the regional lymphatic nodes if suspected of involvement; for instance, the supraclavicular fossa if the primary tumor or any lymphadenopathy were detected in the upper mediastinum or the celiac lymphatics if the primary tumor or

any lymphadenopathy was present close to the esophagogastric junction.

- ❖ High-dose and low-dose planning target volume (PTV<sub>High</sub> and PTV<sub>Low</sub>): addition of CTV<sub>High</sub> and CTV<sub>Low</sub> with a 0.5-cm expanding margin in all directions.

We delivered 6-MV or 10-MV photon beams via intensity-modulated radiation therapy or volumetric arc therapy (VMAT) planning using the Eclipse treatment planning system version 13 (Varian Medical Systems Inc., Palo Alto, CA, USA) or Pinnacle treatment planning system versions 9.2 and 10.1 (Philips Healthcare Inc., Andover, MA, USA). The PTV<sub>High</sub> and PTV<sub>Low</sub> were prescribed radiation doses of 48 and 43.2 Gy, respectively, divided into 24 fractions using a simultaneous integrated boost method. The 100% coverage of the prescribed dose volume should exceed 95% of the targeted PTV whilst meeting the normal organ constraints, including the following: maximum dose to the spinal cord should be <45 Gy, volume of the lung receiving >20 Gy should be <30% of the whole lung; and mean dose to the heart should be <30 Gy.

All patients underwent concomitant chemoradiotherapy with the following weekly platinum-based regimens: (I) cisplatin 30 mg/m<sup>2</sup>; (II) carboplatin (AUC =2); or (III) paclitaxel 50 mg/m<sup>2</sup> plus either cisplatin 30 mg/m<sup>2</sup> or carboplatin (AUC =2).

### *Image processing and extraction of radiomics features*

Three patients were excluded from image processing: two were excluded because of poor quality of post-treatment CT images due to metal implant and one was excluded because of unfinished radiotherapy. The CT or PET/CT images after NACCRT were imported to the Eclipse treatment planning system (version 13). Planning CT images from 59 patients were rigidly co-registered with post-treatment images according to the gray level inside clipbox, sized in whole chest and centrally aligned in planning center, to avoid unwanted deformation of target organs. The planning GTV were duplicated to its relative post-treatment CT images and defined as regions of interest (ROIs). The ROIs of GTV and relative CT series were inserted into the LIFEx software (16) (version 4.7) to extract radiomics features as follow: 3-dimensional Lagrangian polygon interpolation was applied for resampling images into 1×1×3 mm<sup>3</sup> voxels. Hounsfield units (HUs) in all the images were then resampled into 400 discrete values (bins) with absolute

discretization from −1,000 to 3,000 HUs. Grey-level matrices were calculated in 3 dimensions to gather 46 radiomic features.

### *Statistical analysis and computing prediction model*

Statistical analysis was performed using the Statistical Package for the Social Sciences for Windows, SPSS® software v. 18.0 (IBM Corp., New York, NY, USA; formerly SPSS Inc., Chicago, IL, USA) and R computing software (version 3.6.0) under the environment of RStudio (version 1.2.1335.) The correlation among pathological complete remission, recurrence pattern, clinical features, and radiomics features were analyzed using the Pearson correlation coefficient (PCC). Clinical features showing significant correlation with complete remission and recurrence patterns were analyzed as predictors using logistic regression. Radiomics feature reduction and regularization of the regression model was performed using the least absolute shrinkage and selection operator (Lasso) formula. Correlation coefficients of Lasso-selected features were calculated and presented as a correlation matrix for reducing collinearity. The predictability of the radiomic features model was assessed by area under the curve (AUC) analysis of the receiver-operating characteristic (ROC) curve. Cut-off points of the expected value in prediction models were assessed by the Youden index (sensitivity + specificity − 1). The Kaplan-Meier method was used to calculate overall survival (OS) and progression-free survival (PFS), which were determined from the date of biopsy to the date of last follow-up. The association of OS and PFS with clinical or radiomics features was analyzed using Cox regression.

## **Results**

### *Patient characteristics*

Table 1 shows the characteristics of the 62 advanced thoracic ESCC patients treated with NACCRT. Median age at diagnosis was 60 years (range, 40–79 years). The majority of patients had clinical stage III disease (n=50, 80.6%), cT3 disease (n=48, 77.4%), lymph node positivity (n=60, 96.8%), and histological grade 2 differentiation (n=43, 69.4%). Primary tumor at the middle esophagus was observed in 51.6% of patients (n=32). Median follow-up duration was 20.6 months (range, 5.2–60.8 months).

**Table 1** Baseline characteristics of patients with advanced thoracic esophageal cancer who underwent NACCRT and esophagectomy

Characteristic	Case, N (%)
Total	62
Gender (M:F)	60:2
Age	60.8±8.3
Performance status (ECOG score)	
0	39 (62.9)
1	21 (33.9)
2	2 (3.2)
Clinical T stage	
T1	2 (3.2)
T2	10 (16.1)
T3	48 (77.4)
T4a	2 (3.2)
Clinical N stage	
N0	2 (3.2)
N1	27 (43.5)
N2	24 (38.7)
N3	9 (14.5)
Clinical M stage	
M0	59 (95.2)
M1	3 (4.8)
Clinical stage	
IIIB	9 (14.5)
IIIA	22 (35.5)
IIIB	17 (27.4)
IIIC	11 (17.7)
IV	3 (4.8)
Differential grade	
NA	10 (16.1)
1	0 (0.0)
2	43 (69.4)
3	9 (14.5)
Esophagus position	
Upper	14 (22.6)
Middle	32 (51.6)
Lower	16 (25.8)

Data are presented as mean ± standard deviation for age and as case number (percentage) for other variables; NACCRT, neoadjuvant concomitant chemoradiation; ECOG score, Eastern Cooperative Oncology Group performance status score; NA, no assessment of the differential grade owing to the quantity of the endoscopy-aided biopsy sample obtained.

### Treatment compliance, outcomes, and toxicities

The study cohorts showed favorable tolerance, such that 60 of 62 (96.8%) patients completed both radiotherapy and the entire chemotherapy course. Interruption of NACCRT occurred in one patient who developed intolerable fatigue during treatment. Concomitant chemotherapy was discontinued early in one elderly patient who developed severe nausea and weakness after chemotherapy. The clinical treatment outcomes are summarized in *Table 2*. The clinical down-staging rate was 77.4%. The pCR rate was 40.3% (25/62). The 1-, 3-, and 5-year OS rates were 74.7%, 48.8%, and 39.8%, respectively. The 1-, 3-, and 5-year PFS rates were 60.1%, 30.6%, and 18.1%, respectively. The local recurrence rate was 19.4% (12/62), and the distant recurrence rate was 37.1% (23/62). Kaplan-Meier survival curves of OS and PFS are shown in *Figure 1A,B*. In total, 6 patients showed hematological toxicities greater than grade 3. Two patients developed acute radiation pneumonitis ≥ grade 3, one of which died from pneumonitis-induced acute respiratory distress syndrome. One patient died owing to post-surgery infection.

### Response and survival predictive capabilities of clinical features

Correlation coefficients of all 25 clinical features for complete response (CR), local regional failure (LRF), and distant failure (DF) were estimated with double-tail PCC analysis. Clinical T stage and anemia were negatively correlated with CR. CR was the only feature that was negatively correlated with LRF. Clinical N stage, overall clinical stage, pathological T and N stages, clinical response, and resection status were correlated with DF. In Kaplan-Meier analysis, patients with and without CR showed significantly different survival curves (*Figure 1C,D*). Features showed statistical significance of PCC ( $P < 0.05$ ) was further established its log risk model to CR, LRF, and DF by logistic regression (*Table 3*). Furthermore, the hazard ratios of clinical features predicting PFS and OS were calculated using Cox regression (*Table 4*). Adjuvant chemotherapy caused a 2.79-fold reduction in mortality risk, whereas an increase in pathology N stage and anemia grade caused a 1.8-fold increase in hazard ratio of mortality. Resection status was the strongest positive predictor of the hazard ratio of progression whereas anemia and pT and pN stage moderately predicted the hazard ratio of progression. Treatment response negatively predicted progression, with

**Table 2** Treatment, compliance, and outcome among patients with advanced thoracic esophageal cancer who underwent NACCRT and esophagectomy.

Characteristic	Case, N (%)
Concomitant chemotherapy	
Single platinum	34 (54.8)
Platinum-taxol	26 (41.9)
Others	2 (3.2)
Chemotherapy dose adjustment	
No	41 (66.1)
Yes	21 (33.9)
Adjuvant chemotherapy	
Yes	31 (50.0)
No	31 (50.0)
Treatment compliance	
Complete	60 (96.8)
Incomplete CT	1 (1.6)
Incomplete CT and RT	1 (1.6)
Response	
CR	24 (38.7)
PR	24 (38.7)
SD	8 (12.9)
PD	6 (9.7)
Excision status	
R0	58 (93.5)
R1	2 (3.2)
R2	2 (3.2)
Pathology T stage	
T0	24 (38.7)
Tis	1 (1.6)
T1	7 (11.3)
T2	14 (22.6)
T3	15 (24.2)
T4a	1 (1.6)
Pathology N stage	
N0	49 (79.0)
N1	5 (8.1)
N2	6 (9.7)
N3	2 (3.2)

Data are presented as mean  $\pm$  standard deviation for treatment duration and case number (percentage) for other variables; weekly platinum, patients underwent weekly cisplatin or weekly carboplatin treatment; NACCRT, neoadjuvant concomitant chemoradiation; CT, chemotherapy; RT, radiotherapy.

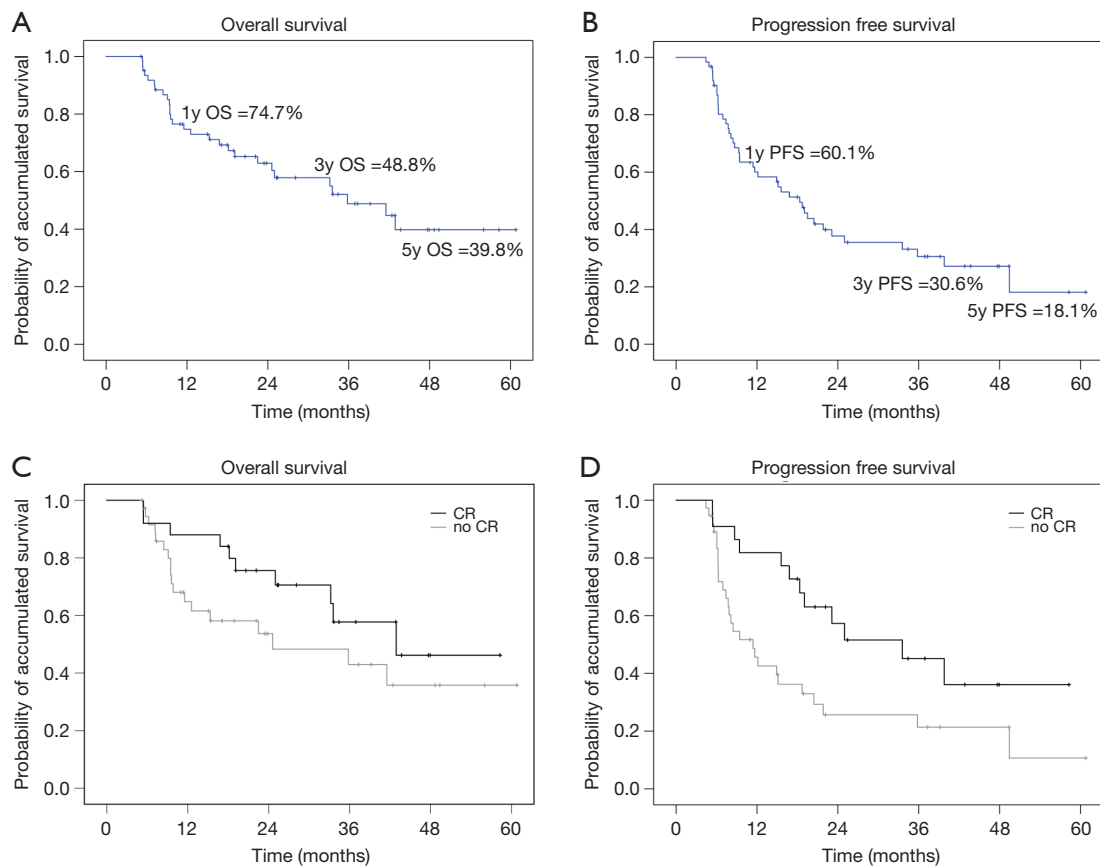
an expected hazard ratio of 0.6.

### *Response- and survival-related predictive capabilities of radiomics features*

Eight features [1 shape feature, 1 neighborhood grey-level different matrix (NGLDM) feature, 2 grey-level co-occurrence matrix (GLCM) features, and 4 grey-level zone length matrix (GLZLM) features], including PCC (*Figure 2A*) were selected using the Lasso regularization formula (*Figure 2B*). The regression model comprising the abovementioned 8 features showed good capability with acceptable representativeness (AUC =0.851, pseudo-R<sup>2</sup> =0.504; ROC curve presented in *Figure 2C*) for predicting CR to NACCRT. The prediction model of LRF and DF with features restricted by the Lasso formula (left panel of *Figure 3*) had medium capability but poor representativeness (AUC =0.773 and 0.721, pseudo-R<sup>2</sup> =0.291 and 0.213, respectively; ROC curve presented in the right panel of *Figure 3*). Using K-fold cross validation by the glmnet package of R software, radiomics features having unreliable coefficients were first excluded (*Figure S1A,B*). The Cox regression model showed that minimal UH, GLRLM\_GLNU, GLZLM\_LZE, and GLZLM\_LZHGE strongly increased the hazard ratio of mortality by 4- to 6-folds, whereas sphericity of shape strongly reduced the hazard ratio of mortality by 6.6-folds. For PFS, 3 GLRLM family and 2 GLZLM family radiomics strongly predicted the hazard ratio of progression.

### *Combining clinical and radiomics features for enhancing prediction power and reliability*

Individually, clinical and radiomics features had problems in terms of prediction power and reliability for predicting LRF and DF. We combined both clinical and radiomic features to develop a new prediction model. In the LRF prediction model that combined the clinical feature “complete response” and 3 radiomics features, the AUC was 0.804 (*Figure 4A*) and pseudo-R square was 0.339. Contrastingly, in the DF prediction model that combined clinical and radiomic features, the AUC and pseudo-R square values were 0.754 (*Figure 4B*) and 0.320, respectively. The formula of LRF and DF prediction models using a combination of clinical and radiomic features were detail in *Table 5*. Both the LRF and DF prediction models with combined clinical and radiomic features had stronger power and better reliability than those using each type of feature



**Figure 1** Overall and progression-free survival. Curves depicting overall survival (A) and progression-free survival (B) in the study cohort. Subgroup survival curve for overall survival (C) and progression-free survival (D) in patients with and without a complete response, obtained using Kaplan-Meier analysis.

individually. Because of the limitation of censored numbers, Cox regression could only proceed with one variable in the overall prediction model in the present study. Therefore, we only tested the combination of one clinical feature and one radiomics feature in the prediction model of PFS. The combination of resection status and selected radiomics, any of the 5 radiomic predictors having significant correlation to hazard ratio of progression, mutually enhanced the expected hazard ratio value of progression. The strongest combination was resection status multiplied by GLRLM\_LRE, which resulted in an expected hazard ratio value of progression of 8.776 (95% CI: 1.959–39.320,  $P=0.005$ ) and good model reliability, tested using the chi-square test of residuals and the omnibus test.

## Discussion

The current esophageal cancer treatment guidelines

recommend CT or PET/CT as the standard of response assessment (4). However, even after 4–6-week recovery intervals, residual tumors at the esophagus on contrast-enhanced CT could be masked by submucosal swelling via visual interpretation (12). It is also difficult to differentiate residual lymph nodes at the pulmonary hilum with the bare human eye (17). Endoscopic ultrasound and biopsy can hardly detect submucosal residual tumors, which are suggested to be predictors of poor pathological outcomes (18). Post-NACCRT surveillance generally involves FDG-PET/CT; although FDG-PET/CT is good in differentiating residual lymph nodes, they tend to get obscured by treatment-induced inflammation (12). A good post-treatment surveillance method is yet to be developed.

Radiomics, known as the next generation technology of computer-aided diagnosis and detection systems, is more capable than answering yes-or-no questions in lesion detection. With the combination of texture

**Table 3** Response and failure patterns and clinical features in the study cohort

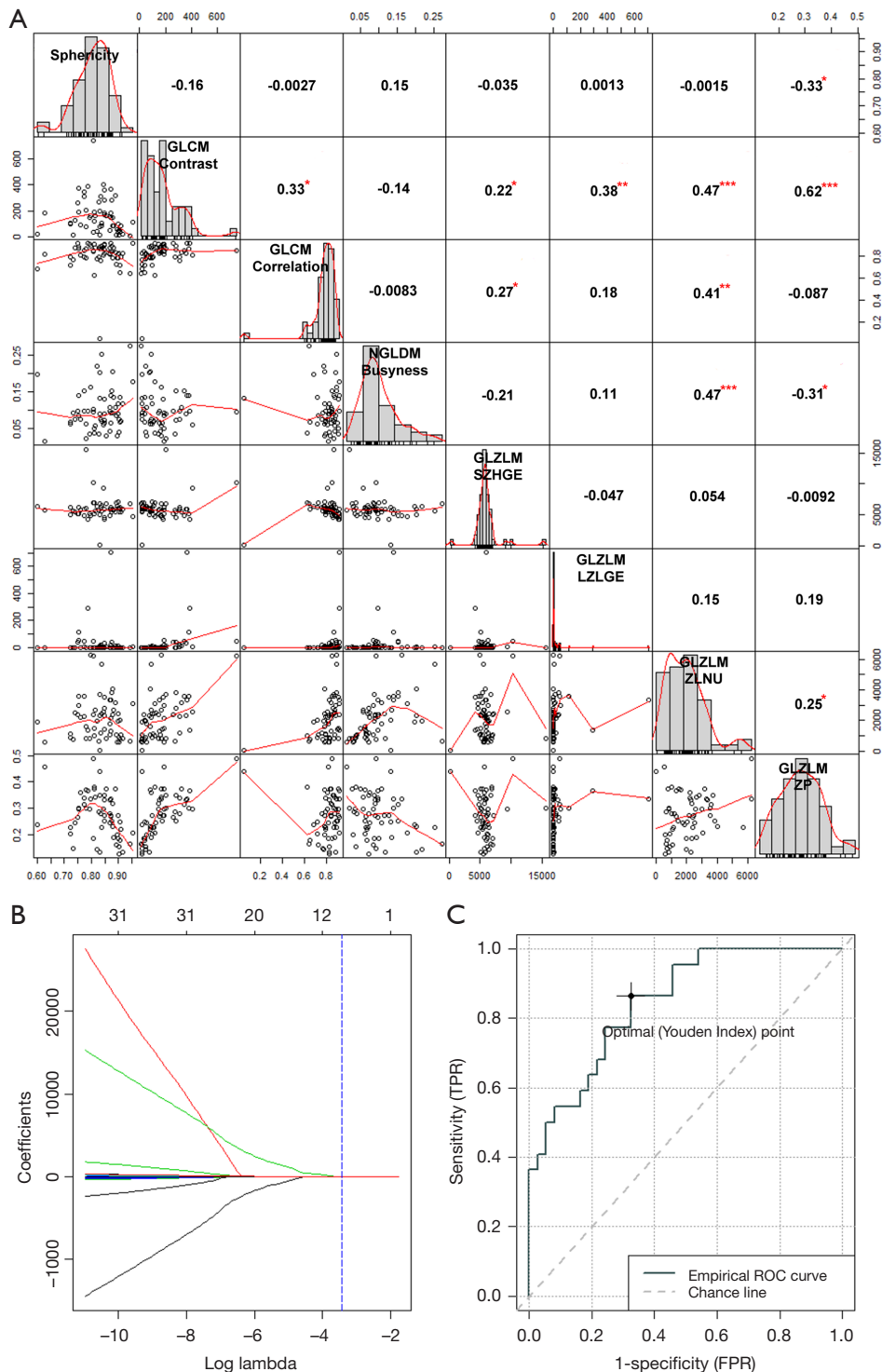
Features	Beta(x)	SE	P value	Exp. (95% CI)
CR prediction model (logistic regression)				
Clinical T stage	-1.586	0.685	0.021*	0.205 (0.054–0.783)
Anemia	-1.477	0.491	0.003*	0.228 (0.087–0.598)
Constant	5.810	2.257	0.010*	–
Pseudo-R2 =0.345				
LRF prediction model (logistic regression)				
CR	-1.349	0.708	0.057	0.259 (0.065–1.038)
Constant	-0.496	0.339	0.143	–
Pseudo-R2 =0.277				
DF prediction model (logistic regression)				
cN stage	0.868	0.427	0.042*	2.381 (1.031–5.501)
pT stage	0.496	0.316	0.117	1.642 (0.884–3.052)
pN stage	0.723	0.520	0.164	2.061 (0.744–5.707)
Response	0.134	0.516	0.795	1.143 (0.416–3.140)
Constant	-3.452	2.220	0.120	–
Pseudo-R2 =0.299				

The expected value of the features was presented through a log risk ratio obtained using logistic regression, followed by the 95% confidence interval. The results of the goodness of fit test were presented as a pseudo-R square value obtained using the Nagelkerke method. \*, star marked as statistical significance. Beta(x), coefficient of variate X; SE, standard error of coefficient, Exp., expected value; CI, confidence interval;

**Table 4** Survival prediction using clinical, radiomic, and combine features

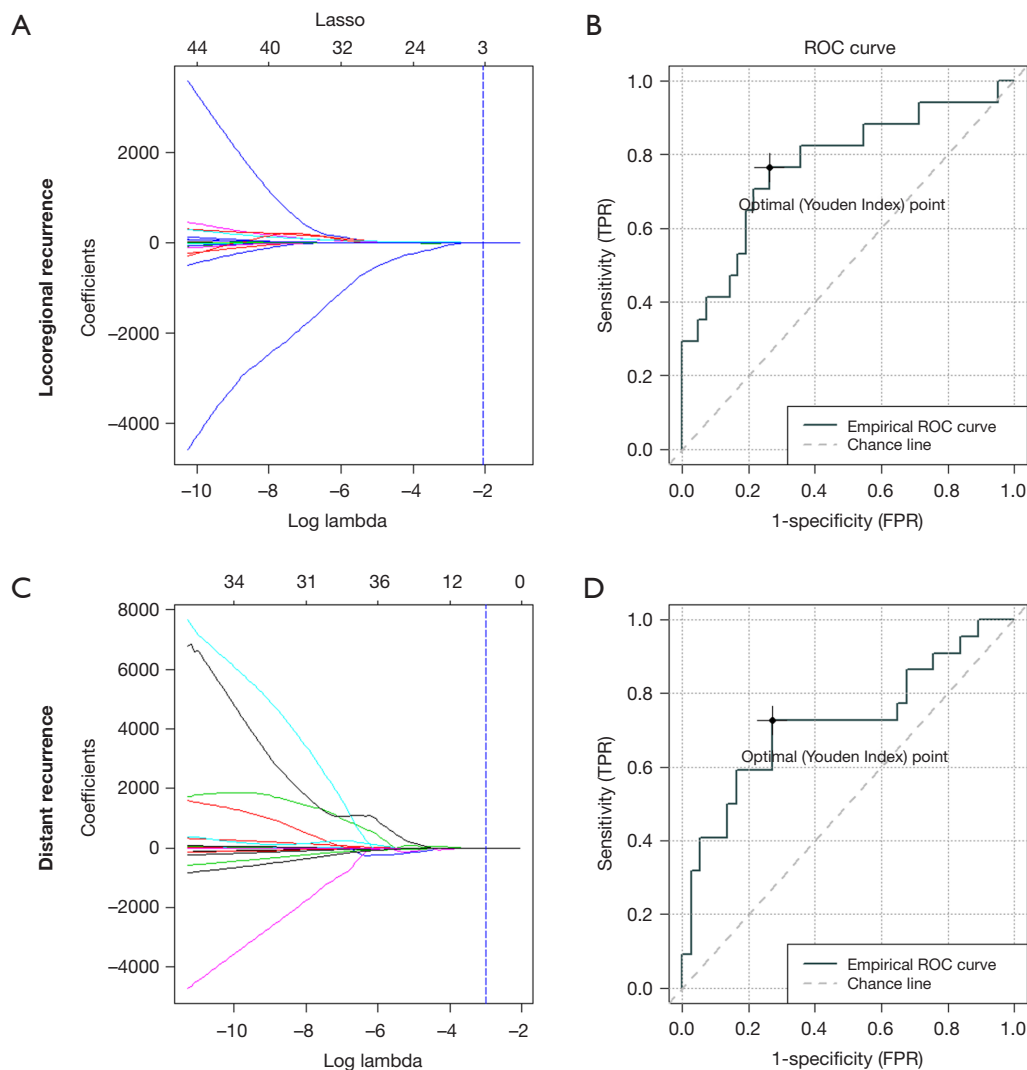
Features	Beta(x)	SE	P value	Exp. (95% CI)
Significant predictors of OS in the logistic regression				
Adj. CTx	-1.024	0.422	0.015	0.359 (0.157–0.821)
Anemia	0.611	0.268	0.023	1.843 (1.089–3.118)
pN stage	0.605	0.191	0.002	1.831 (1.259–2.661)
UHmin	1.747	0.801	0.029	5.737 (1.195–27.549)
SHAPE_Sphericity	-1.881	0.916	0.040	0.152 (0.025–0.917)
GLRLM_GLNU	1.930	0.825	0.019	6.888 (1.368–34.677)
GLZLM_LZE	1.465	0.734	0.046	4.327 (1.026–18.240)
GLZLM_LZHGE	1.454	0.735	0.048	4.281 (1.013–18.095)
PFS prediction model (logistic regression)				
Anemia	0.669	0.219	0.002	1.953 (1.270–3.002)
pT stage	0.294	0.111	0.008	1.341 (1.078–1.669)
pN stage	0.468	0.182	0.010	1.597 (1.118–2.282)
Response	-0.505	0.168	0.003	0.603 (0.434–0.839)
Resection status	0.865	0.348	0.013	2.376 (1.201–4.700)
GLRLM_SRE	-1.585	0.786	0.044	0.205 (0.044–0.955)
GLRLM_LRE	1.688	0.817	0.039	5.410 (1.092–26.810)
GLRLM_RP	-1.667	0.804	0.038	0.189 (0.039–0.914)
GLZLM_LZE	1.429	0.696	0.040	4.175 (1.067–16.340)
GLZLM_LZHGE	1.393	0.702	0.047	4.026 (1.018–15.924)

The expected value of the features was presented as the hazard ratio followed by the 95% confidence interval. The goodness of fit test is presented using the omnibus test. Beta(x), coefficient of variate X; SE, standard error of coefficient, Exp., expected value; CI, confidence interval.



**Figure 2** Prediction of complete response using radiomics features selected by Lasso regression. (A) Person correlation matrix of selected radiomic features presented as a coordinates plot with a trendline and correlation coefficient number; \*, weak; \*\*, moderate; \*\*\*, strong correlation. (B) Radiomic feature selection through Lasso regularization to reduce the interruption of variates with unreasonable coefficients. (C) Receiver-operator characteristic curve of the complete response prediction model and the operating point of the expected value with the maximum Youden index.



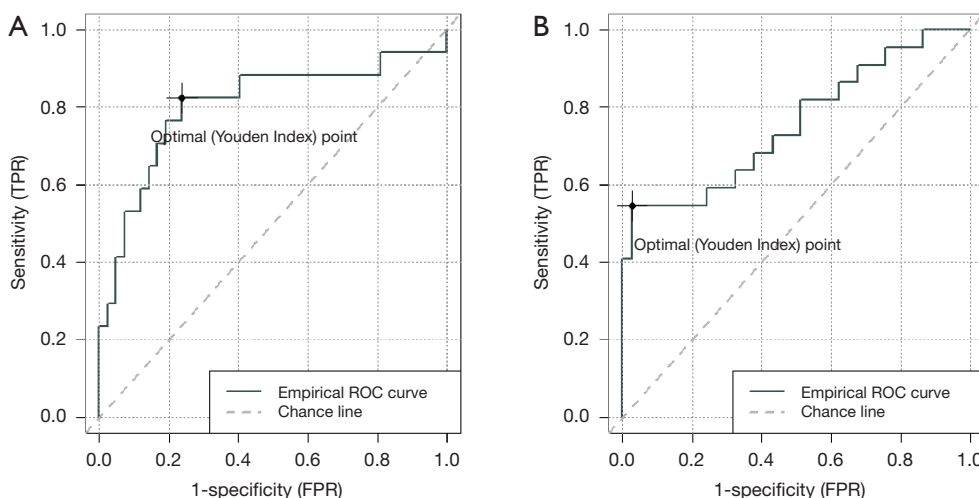


**Figure 3** Prediction of LRF and DF with radiomics. LRF, local regional failure; DF, distant failure. (A,C) Radiomic feature selection through Lasso regularization to reduce the interruption of variates with unreasonable coefficients; (B,D) power of the prediction model, presented with the receiver-operator characteristic curve and the operating point of the expected value with the maximum Youden index. (A,B) Locoregional failure; (C,D) distant failure.

extraction, artificial intelligence, and new algorithms that are able to reduce variates, radiomics can not only be applied in disease diagnosis but also used to extract more information for helping clinical decision making (19). A good example is the application of radiomics in treatment response evaluation after neoadjuvant treatment for nasopharyngeal cancer (NPC) and rectal cancer (20,21); with this technique, the surgeon has more information before performing the operation. Recent studies have also evaluated radiomics characteristics for directly predicting cancer prognosis, for instance in lung cancer, NPC, and

hepatic cancer (10,11,13).

In 2016, van Rossum reviewed the evidence and potential future of applying radiomics in the field of esophageal cancer (22). By using pre-treatment contrast-enhanced CT, Hou *et al.*'s study suggested that radiomics could predict treatment response to definitive chemoradiation in inoperable esophageal cancer patients (23). However, Hou *et al.*'s study had different predictors to PD, SD, PR, and CR, indicating that they did not have a unified prediction model for treatment response. Jin *et al.*' study used a combination of radiotherapy dosimetry and radiomics



**Figure 4** Receiver-operator characteristic curve depicting the power of the clinical-radiomic combination model for prognosis prediction. Receiver-operator characteristic curves for (A) locoregional recurrence and (B) distance recurrence and the operating point of the expected value with the maximum Youden index.

**Table 5** Failure pattern prediction using a combination of clinical and radiomic features

Features	Beta(x)	SE	P value	Exp. (95% CI)
LRF prediction model (logistic regression)				
Complete response	-1.274	0.857	0.137	0.280 (0.052–1.499)
SHAPE_Sphericity	-3.104	1.829	0.090	0.045 (0.001–1.619)
NGLDM_Busyness	4.017	1.702	0.018*	55.533 (1.977–1.560×10 <sup>3</sup> )
GLZLM_LZLGE	2.840	2.264	0.210	17.110 (0.202–1.447×10 <sup>3</sup> )
Constant	-0.119	1.193	0.921	–
Pseudo-R2 =0.339				
DF prediction model (logistic regression)				
pN stage	0.954	0.475	0.045*	2.595 (1.022–6.590)
SHAPE_Sphericity	-4.387	1.963	0.025*	0.012 (0.000–0.583)
GLRLM_LRE	-1.378	3.341	0.680	0.252 (0.000–1.760×10 <sup>2</sup> )
GLRLM_GLNU	-1.063	2.124	0.617	0.345 (0.005–2.222×10 <sup>1</sup> )
GLZLM_LZLGE	3.490	2.485	0.160	32.780 (0.252–4.271×10 <sup>3</sup> )
GLZLM_ZP	-4.957	3.122	0.112	0.007 (0.000–3.200)
Constant	4.405	2.874	0.125	–
Pseudo-R2 =0.320				

The expect value of the features was presented through a log risk ratio obtained using logistic regression, followed by the 95% confidence interval. The results of the goodness of fit test were presented as a pseudo-R square value obtained using the Nagelkerke method. \*, star marked as statistical significance. Beta(x), coefficient of variate X; SE, standard error of coefficient, Exp., expected value; CI, confidence interval.

to predict chemoradiation treatment response (14) and integrated 63 features in one prediction model. However, many of these features showed strong linear correlation in the Person correlation matrix. More than 2 studies did not have good prediction power (AUC =0.6–0.7) and did not mention evidence of robustness and reliability. Larue *et al.*'s study focused on esophageal cancer patients who underwent NACCRT and directly predicted 3-year OS using a model based on radiomics features extracted from pre-treatment CT images (15). However, Larue *et al.*'s model reported an AUC of 0.61 in the validation dataset, and the risk group predicted by the model showed no significant correlation with pathological response, which is not consistent with the current clinical consensus.

According to our review of the literature, this is the first study in esophageal cancer research that attempted to establish a prediction model based on post-NACCRT CT images. With regards to prediction models of pCR and failure patterns, radiomics showed better capability and representativeness than that seen in previous studies based on pre-treatment CT images. LIFEx, a widely accepted freeware with literature support was used to extract 42 histogram, textural and shape radiomics features. As the result (*Figure 1C,D*), significantly longer PFS and OS were observed in the pCR group than in the non-pCR group. Therefore, our high prediction power model showed in *Figure 2* is useful for deciding between esophagectomy and definitive chemoradiation. Furthermore, in the high-risk patient group, the failure pattern prediction model indicated that adjuvant treatment after surgery might have prevented local or distant recurrence.

A few clinical and radiomic predictors were common to the prediction models of DF pattern, PFS, and OS. Residual pathological lymph node significantly affected prognosis, as observed in this study and widely discussed in other studies (6). In the Cox regression model for predicting PFS and OS, radiomics features showed more power for predicting hazard ratios than clinical features. Moreover, both failure pattern and survival prediction models with combined clinical and radiomic predictors showed synergistically enhanced prediction powers. Interestingly, pathological response was a strong predictor of PFS but not OS. This may indicate that, in addition to residual tumors, other factors in the surrounding environment have an influence on patient survival; the adverse effects of chemoradiation as well as immune response may influence patient survival.

Interpreting the meaning of radiomics may reveal its role

in the prediction model. The most important radiomics feature in all prediction models was the long-zone high gray-level emphasis (GLZLM\_LZHGE), defined as the distribution of the long homogeneous zones with high grey-levels, representing poor prognosis with a 4-fold increase in hazard ratios of progression and mortality. High sphericity was a negative predictor of LRF, DF, and mortality. Long homogeneous runs in an image (GLRLM\_LRE) seem like spiculated extended structures that predict DF and progression. Non-uniformity of the grey-levels (GLRLM\_GLNU) is a strong positive predictor of DF and mortality.

The limitation of this study is that the number of patients was too small for separation into the training and validation datasets. Therefore, we were unable to prove the robustness of the prediction model. Furthermore, the retrospective and single-center nature of the study may have introduced selection bias. The established model is still in the preliminary stage. In the future, we aim to test our prediction model by validating the images from the public patient database. We would also improve the algorithm to enhance the accuracy of feature elimination and power of prediction.

In conclusion, we established an integrated model that combined clinical features and post-treatment CT radiomics features for predicting response and survival outcomes of esophageal cancer patients undergoing neoadjuvant chemoradiation. The integrated prediction model may aid clinicians in decision making regarding post-operative adjuvant therapy for ESCC patients who have received neoadjuvant chemoradiation.

## Acknowledgments

*Funding:* This study was funded by MacKay Memorial Hospital (grant number: MMH-E-107-13) and the Ministry of Science and Technology, Taiwan (grant number: MOST 106-2314-B-195-002-MY3 and MOST-106-2623-E-195-001-NU).

## Footnote

*Conflicts of Interest:* The authors have no conflicts of interest to declare.

*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 protocol

was granted institutional review board approval (serial number: 18MMHIS194e). Written informed consent was obtained from all study participants and research was conducted in accordance with the 1975 Declaration of Helsinki, as revised in 2013.

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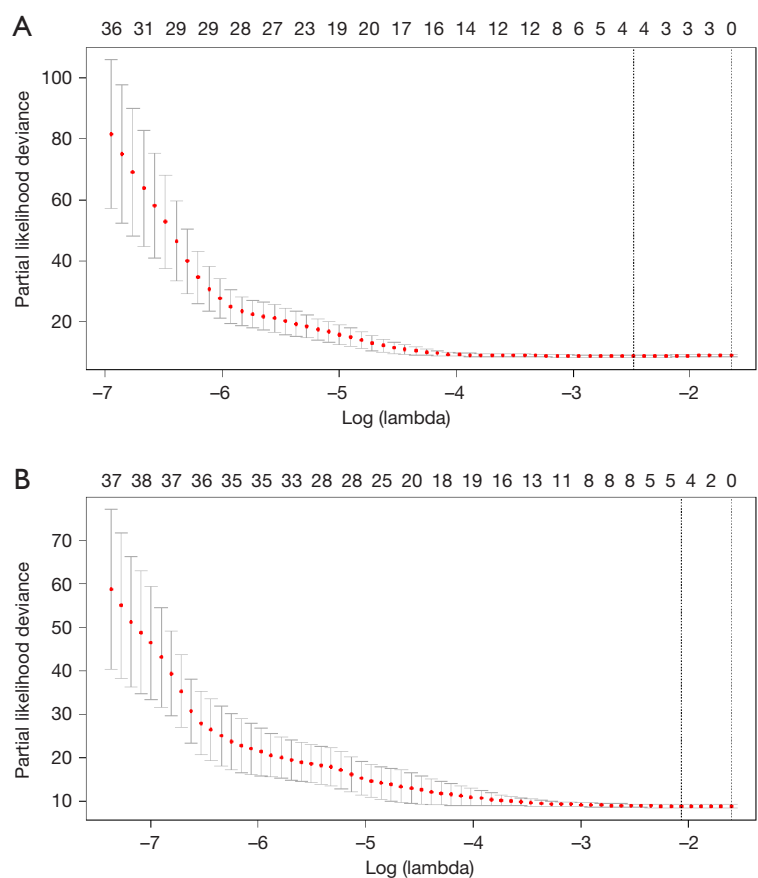
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doi: 10.21037/tro.2019.07.03

**Cite this article as:** Hou TC, Huang WC, Tai HC, Chen YJ. Integrated radiomic model for predicting the prognosis of esophageal squamous cell carcinoma patients undergoing neoadjuvant chemoradiation. *Ther Radiol Oncol* 2019;3:28.



**Figure S1** Survival prediction using radiomic features selected by K-fold cross validation. (A) Predicting overall survival using radiomic features selected by K-fold cross validation; (B) predicting progression-free survival using radiomic features selected by K-fold cross validation.