



Development and validation of a model to predict the risk of recurrence in patients with laryngeal squamous cell carcinoma after total laryngectomy

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Background: Recurrence is still the main obstacle to the survival of laryngeal squamous cell carcinoma (LSCC) patients who have undergone a total laryngectomy. Previous models for recurrence prediction in patients with LSCC were based on pathological information, while the role of easily accessible inflammatory markers in the prognosis of LSCC patients has rarely been reported. This study sought to develop and validate a model to predict the risk of recurrence in LSCC patients who underwent total laryngectomy.

Methods: A total of 204 LSCC patients who underwent a total laryngectomy were included in this retrospective cohort study. Demographics, pathology, and inflammatory markers of patients were collected. All the patients were randomly divided into a training set and a test set at a ratio of 4:1. Patients were followed up for 3 years after surgery or until death occurred during this period. The random-forest method was used to develop a predictive model. The performance of the model was evaluated by calculating the area under the receiver operating characteristic (ROC) curve (AUC) with the 95% confidence interval (CI), sensitivity, specificity, positive predictive value (PPV), and negative predictive value (NPV).

Results: Of the 204 LSCC patients, 56 (27.45%) patients had a recurrence. The random-forest prediction model was an all-factor model, and the most important predictors of the model were the albumin/globulin ratio (AGR), neutrophil/lymphocyte ratio (NLR), and platelet/lymphocyte ratio (PLR), with proportions of 0.121, 0.100, and 0.092, respectively. The AUCs of the model in predicting the recurrence of LSCC in the training set and the test set were 0.960 (95% CI, 0.931–0.989) and 0.721 (95% CI, 0.716–0.726), respectively. The sensitivity, specificity, accuracy, PPV, and NPV of the model in the test set were 0.750 (95% CI, 0.505–0.995), 0.690 (95% CI, 0.521–0.858), 0.707 (95% CI, 0.568–0.847), 0.500 (95% CI, 0.269–0.921), and 0.870 (95% CI, 0.732–1.000), respectively.

Conclusions: A model to predict the risk of recurrence in LSCC patients who have undergone a total laryngectomy was established, and inflammatory markers AGR, NLR, and PLR play an important role in the predictive model.

Keywords: Laryngeal squamous cell carcinoma (LSCC); prediction model; recurrence; total laryngectomy

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Introduction

Laryngeal squamous cell carcinoma (LSCC) is one of the most common cancers in the head and neck area, and accounts for 85–95% of all laryngeal cancers (1). It is estimated that 12,470 new cases of laryngeal cancer will be diagnosed and approximately 3,820 patients will die from the disease in the United States in 2022 (2). Approximately 60% of patients have advanced (stage III or IV) disease at the time of diagnosis (3,4). Current treatment methods for LSCC include surgery, radiotherapy, and chemotherapy (5). Total laryngectomy is used as the primary therapy for advanced laryngeal cancer (6). Several studies have suggested that undergoing a total laryngectomy for advanced-stage laryngeal cancer can improve patient survival (7,8). However, even after a total laryngectomy of the LSCC, patients are still at risk of recurrence.

A review reported that the local recurrence rate after a total laryngectomy is between 30–66% in patients with recurrent or persistent laryngeal cancer (9). Importantly, the survival of patients who have undergone a salvage total laryngectomy has been reported to be significantly lower than that of patients who do not require salvage surgery (9). Thus, it is important to identify the prognostic factors influencing the oncologic outcomes after a total laryngectomy, and to predict the likelihood of recurrence to optimize patient treatment and follow-up. Many factors have been reported to have predictive value in LSCC, including clinical (e.g., gender and age) (10), pathological (e.g., tumor stage, size, and grade) (11,12), and genetic variables (e.g., tumor protein p53 mutations and long non-coding RNA AC008440.10) (13,14). Due to the variability of clinicopathological features and tumor biology, a single feature has limited predictive effect in clinical practice. Prediction models can improve the accuracy of prediction by integrating multiple clinical variables and provide clinicians with prognostic prediction tools for individualized patients. Cui *et al.* developed a nomogram for predicting the risk of recurrence after curative-intend surgery in patients with LSCC (15). However, their model is mainly based on pathological data, and the inclusion of some routine inflammatory indicators related to the prognosis of LSCC patients may further improve the ability of the model. Several studies have shown that inflammation markers such as the neutrophil/lymphocyte ratio (NLR), platelet/lymphocyte ratio (PLR), and lymphocyte/monocyte ratio (LMR) are independently correlated with poor outcomes in LSCC patients (16–19). However, the role of these

inflammation indicators in the prediction of recurrence in LSCC patients with total laryngectomy has not been reported.

In the present study, we sought to develop and internally validate a model using systematic statistical methods to predict the risk of recurrence in LSCC patients who have undergone a total laryngectomy. We present the following article in accordance with the TRIPOD reporting checklist (available at <https://atm.amegroups.com/article/view/10.21037/atm-22-4802/rc>).

Methods

Study design and population

Patients with LCSS who underwent total laryngectomy between 2012 and 2019 at the Shanxi Province Cancer Hospital were included in this retrospective cohort study. To be eligible for inclusion in this study, patients had to meet the following inclusion criteria: (I) have been diagnosed with LCSS by histology or cytology; (II) have undergone a total laryngectomy; and (III) have complete pathological examination data and follow-up data. Patients were excluded from the study if they met any of the following exclusion criteria: (I) had undergone an emergency tracheotomy; (II) had an active systemic inflammation (such as a pneumonia); (III) had a severe uncontrolled cardiovascular disease, or a recent history of myocardial infarction (within the last 3 months); and/or (IV) had suffered from other malignant tumors within the last 5 years. The study was conducted in accordance with the Declaration of Helsinki (as revised in 2013). This study was approved by the Ethics Committee of Shanxi Province Cancer Hospital (No. 202130) and individual consent for this retrospective analysis was waived.

Outcomes

The primary outcome of this study was the recurrence measured from the time of surgery. All the patients received regular follow-up every 3 months in the first 6 months after resection, and every 6 months thereafter; the length of follow-up was 3 years or until death. Each patient's recurrence and survival status were recorded during the follow-ups. Recurrence was defined as a new laryngeal mass found by imaging examination and confirmed by biopsy or surgical pathology to be laryngeal cancer, or distant lymph node metastasis.

Sample size

According to previous studies (9), the recurrence rate was set as 40%, and the area under the receiver operating characteristic (ROC) curve (AUC) of the recurrence prediction model in previous studies was 0.778 as a reference (15). The sample size was calculated by PASS 15.0.5 software (NCSS, LLC, Kaysville, UT, USA), the AUC of the prediction model in this study was set as 0.90, and 114 patients were assigned to the training set with an α -error of 0.05 and a power of 0.8 (two-sided). The training set and testing set were randomly assigned in a ratio of 2:1, with a total sample size of at least 171.

Data collection and definition

Patients' demographic and clinical data were collected, including data on their age, gender, tumor size, tumor (T) stage, node (N) stage, grade, primary site, metastatic, surgical excision margins extracapsular invasion, lymphatic vascular invasion, NLR, PLR, LMR, albumin/globulin ratio (AGR), prognostic nutritional index (PNI), the combination of the platelet count and the NLR (COP-NLR), the combined score of the plasma fibrinogen level and the NLR (F-NLR), and recurrence.

The PNI was calculated based on the albumin (Alb) and the absolute lymphocyte count (ALC) (20) using the following formula: $PNI = 10 \times Alb + 0.005 \times ALC$. The COP-NLRs were defined as follows: COP-NLR 2—patients with an elevated platelet count $>300 \times 10^9/L$ and a NLR >3 ; COP-NLR 1—patients with 1 abnormal value; and COP-NLR 0—patients with no abnormal value (21,22). The F-NLRs were defined as follows: F-NLR 2—patients with fibrinogen ≥ 341 mg/dL and a NLR ≥ 3.59 ; F-NLR 1—patients with fibrinogen ≥ 341 mg/dL or a NLR ≥ 3.59 ; and F-NLR 0—patients with fibrinogen <341 mg/dL and a NLR <3.59 (23).

Statistical analysis

The continuous variables were tested for normality using the Shapiro-Wilk test; the continuous variables with a normal distribution are expressed as the mean \pm standard deviation (SD), and the *t*-test was used for comparisons between groups. The non-normal variables are expressed as median (interquartile range) [M (Q1, Q3)], and the Wilcoxon rank-sum test was used for comparisons between groups. The categorical variables are expressed

as numbers and percentages, and the chi-square test (χ^2) or Fisher's test was used for comparisons between groups. The data were randomly assigned to a training set and a test set by Python at a ratio of 4:1. The random-forest model was used to construct the prediction model. The performance of the model was quantified by calculating the AUC with the 95% confidence interval (CI), and the sensitivity, specificity, positive predictive value (PPV), and negative predictive value (NPV). The AUC value of the prediction model was over 0.80, indicating good predictive performance of the model.

All the statistical analyses were two-sided, and a P value <0.05 was considered statistically significant. The baseline characteristics were analyzed by the SAS 9.4 software (SAS Institute Inc., Cary, NC, USA). Python 3.7 software (Python Software Foundation, Delaware, USA) was used to develop the random-forest model and plot the importance of the predictors and the ROC curves.

Results

Baseline characteristics

A total of 204 LSCC patients were included in this study. The patients had a mean age of 59.89 ± 8.90 years, and a mean tumor size of 2.93 ± 1.01 cm. Of these patients, 193 (94.61%) were male, 11 (5.39%) were female, and 4 (1.96%) were T1 stage, 28 (13.73%) were T2 stage, 107 (52.45%) were T3 stage, and 65 (31.86%) were T4 stage. In terms of tumor grade, 28 (13.73%) patients had high differentiation, 123 (60.29%) had moderate differentiation, and 53 (25.98%) had poor differentiation. The median NLRs, PLRs, and LMRs of the patients were 2.15 (1.60, 3.14), 123.38 (96.92, 152.58), and 4.26 (3.07, 5.63), respectively. The mean AGRs and PNIs of the patients were 1.57 ± 0.26 and 427.31 ± 37.99 , respectively. The numbers of patients with COP-NLR scores of 0, 1, and 2 were 120 (58.82%), 70 (34.31%), and 14 (6.86%), respectively. The numbers of patients with F-NLR scores of 0, 1, and 2 were 111 (54.41%), 25 (12.25%), and 68 (33.33%), respectively. At the end of the follow-up, 56 (27.45%) patients had relapsed. More detailed characteristics are shown in *Table 1*.

Comparison of differences between the training set and the test set

A total of 204 patients were randomly divided into a training

Table 1 Characteristics of all patients

Variables	Total (n=204)
Age (years), mean \pm SD	59.89 \pm 8.90
Gender, n (%)	
Male	193 (94.61)
Female	11 (5.39)
Tumor size (cm), mean \pm SD	2.93 \pm 1.01
T stage, n (%)	
T1	4 (1.96)
T2	28 (13.73)
T3	107 (52.45)
T4	65 (31.86)
N stage, n (%)	
N0	129 (63.24)
N1	28 (13.73)
N2a	5 (2.45)
N2b	21 (10.29)
N2c	21 (10.29)
Grade, n (%)	
High differentiation	28 (13.73)
Moderate differentiation	123 (60.29)
Poor differentiation	53 (25.98)
Primary site, n (%)	
Glottis	57 (27.94)
Supraglottis	118 (57.84)
Subglottis	13 (6.37)
Transglottis	16 (7.84)
Surgical excision margins, n (%)	
Positive	6 (2.94)
Negative	198 (97.06)
Metastatic, n (%)	
Yes	76 (37.25)
No	128 (62.75)
Extracapsular invasion, n (%)	
Yes	23 (11.27)
No	181 (88.73)

Table 1 (continued)**Table 1** (continued)

Variables	Total (n=204)
Lymphatic vascular invasion, n (%)	
Yes	17 (8.33)
No	187 (91.67)
NLR, M (Q ₁ , Q ₃)	2.15 (1.60, 3.14)
PLR, M (Q ₁ , Q ₃)	123.38 (96.92, 152.58)
LMR, M (Q ₁ , Q ₃)	4.26 (3.07, 5.63)
AGR, mean \pm SD	1.57 \pm 0.26
PNI, mean \pm SD	427.31 \pm 37.99
COP-NLR (score), n (%)	
0	120 (58.82)
1	70 (34.31)
2	14 (6.86)
F-NLR (score), n (%)	
0	111 (54.41)
1	25 (12.25)
2	68 (33.33)
Group, n (%)	148 (72.55)
Relapse group	56 (27.45)
Non-relapse group	148 (72.55)

SD, standard deviation; T, tumor; N, node; NLR, neutrophil/lymphocyte ratio; M (Q₁, Q₃), median (interquartile range); PLR, platelet/lymphocyte ratio; LMR, lymphocyte/monocyte ratio; AGR, albumin/globulin ratio; PNI, prognostic nutritional index; COP-NLR, combination of the platelet count and the NLR; F-NLR, combined score of the plasma fibrinogen level and the NLR.

set and a test set at a ratio of 4:1. The results showed that there was no statistical difference (all $P > 0.05$) among all the characteristics between the training set and the test set (Table 2). Thus, the sampling of the training set and the test set was balanced, and the data of the test set could be used to test the training set.

Comparison of the characteristics between the relapsed and non-relapsed patients

The training set data were divided into the relapse group and non-relapse group. A difference analysis was performed

Table 2 Comparison of the differences between the training set and the test set

Variables	Training group (n=163)	Testing group (n=41)	Statistic	P
Age (years), mean ± SD	59.43±8.97	61.71±8.50	t=-1.470	0.143
Gender, n (%)			-	1.000
Male	154 (94.48)	39 (95.12)		
Female	9 (5.52)	2 (4.88)		
Tumor size (cm), mean ± SD	2.912±1.006	3.020±1.050	t=-0.60	0.546
T stage, n (%)			-	0.436
T1	3 (1.84)	1 (2.44)		
T2	20 (12.27)	8 (19.51)		
T3	85 (52.15)	22 (53.66)		
T4	55 (33.74)	10 (24.39)		
N stage, n (%)			-	0.560
N0	98 (60.12)	31 (75.61)		
N1	24 (14.72)	4 (9.76)		
N2a	5 (3.07)	0 (0.00)		
N2b	18 (11.04)	3 (7.32)		
N2c	18 (11.04)	3 (7.32)		
Grade, n (%)			$\chi^2=0.106$	0.949
High differentiation	23 (14.11)	5 (12.20)		
Moderate differentiation	98 (60.12)	25 (60.98)		
Poor differentiation	42 (25.77)	11 (26.83)		
Primary site, n (%)			-	0.550
Glottis	45 (27.61)	12 (29.27)		
Supraglottis	92 (56.44)	26 (63.41)		
Subglottis	11 (6.75)	2 (4.88)		
Transglottis	15 (9.20)	1 (2.44)		
Surgical excision margins, n (%)			-	0.602
Positive	6 (3.68)	0 (0.00)		
Negative	157 (96.32)	41 (100.00)		
Metastatic, n (%)			$\chi^2=2.386$	0.122
Yes	65 (39.88)	11 (26.83)		
No	98 (60.12)	30 (73.17)		
Extracapsular invasion, n (%)			-	0.787
Yes	18 (11.04)	5 (12.20)		
No	145 (88.96)	36 (87.80)		

Table 2 (continued)

Table 2 (continued)

Variables	Training group (n=163)	Testing group (n=41)	Statistic	P
Lymphatic vascular invasion, n (%)			–	0.344
Yes	12 (7.36)	5 (12.20)		
No	151 (92.64)	36 (87.80)		
NLR, M (Q ₁ , Q ₃)	2.21 (1.65, 3.10)	2.01 (1.34, 3.67)	Z=-0.900	0.368
PLR, M (Q ₁ , Q ₃)	124.34 (98.24, 153.03)	122.66 (93.20, 149.68)	Z=-0.755	0.450
LMR, M (Q ₁ , Q ₃)	4.04 (3.03, 5.48)	5.10 (3.70, 6.18)	Z=1.885	0.059
AGR, mean ± SD	1.57±0.27	1.56±0.25	t=0.110	0.910
PNI, mean ± SD	426.35±39.74	431.13±30.14	t=-0.850	0.399
COP-NLR (score), n (%)			$\chi^2=0.133$	0.715
0	97 (59.51)	23 (56.10)		
1	55 (33.74)	15 (36.59)		
2	11 (6.75)	3 (7.32)		
F-NLR (score), n (%)			$\chi^2=4.285$	0.117
0	94 (57.67)	17 (41.46)		
1	17 (10.43)	8 (19.51)		
2	52 (31.90)	16 (39.02)		
Group, n (%)			$\chi^2=0.085$	0.771
Non-relapse group	119 (73.01)	29 (70.73)		
Relapse group	44 (26.99)	12 (29.27)		

SD, standard deviation; T, tumor; N, node; NLR, neutrophil/lymphocyte ratio; M (Q₁, Q₃), median (interquartile range); PLR, platelet/lymphocyte ratio; LMR, lymphocyte/monocyte ratio; AGR, albumin/globulin ratio; PNI, prognostic nutritional index; COP-NLR, combination of the platelet count and the NLR; F-NLR, combined score of the plasma fibrinogen level and the NLR.

between the relapse group and non-relapse group. The results showed that there were significant differences in the PLR (Z=2.275, P=0.023), AGR (t=2.420, P=0.017), and proportion of males (P=0.002), metastatic status ($\chi^2=5.409$, P=0.020), COP-NLR ($\chi^2=4.192$, P=0.041), and F-NLR ($\chi^2=4.242$, P=0.039). These 6 factors may be associated with recurrence in LSCC patients who have undergone a total laryngectomy (see Table 3).

Development and validation of the random-forest model

The random-forest method was used to develop a model to predict the recurrence of LSCC in patients who have undergone a total laryngectomy. Under the random-forest model, the most important predictors were the AGR, NLR, and PLR, which contributed 0.121, 0.100, and 0.092 to the

model, respectively (see Figure 1).

In the training set, the AUC of the random-forest model was 0.960 (95% CI, 0.931–0.989); according to the Youden Index, a cutoff value of 0.253 was selected (see Figure 2A). The sensitivity, specificity, accuracy, PPV, and NPV of the model in the training set were 0.955 (95% CI, 0.893–1.000), 0.832 (95% CI, 0.765–0.899), 0.865 (95% CI, 0.813–0.917), 0.677 (95% CI, 0.561–0.948), and 0.980 (95% CI, 0.953–1.000), respectively. When the data of the test set was substituted into the prediction model of the training set, the AUC, sensitivity, specificity, accuracy, PPV, and NPV of the model in the test set were 0.721 (95% CI, 0.716–0.726), 0.750 (95% CI, 0.505–0.995), 0.690 (95% CI, 0.521–0.858), 0.707 (95% CI, 0.568–0.847), 0.500 (95% CI, 0.269–0.921), and 0.870 (95% CI, 0.732–1.000), respectively (see Figure 2B and Table 4).

Table 3 Comparison of the characteristics between relapsed patients and non-relapsed patients

Variables	Relapse group (n=44)	Non-relapse group (n=119)	Statistic	P
Age (years), mean \pm SD	61.27 \pm 10.26	58.75 \pm 8.38	t=-1.60	0.111
Gender, n (%)			-	0.002
Male	37 (84.09)	117 (98.32)		
Female	7 (15.91)	2 (1.68)		
Tumor size (cm), mean \pm SD	3.13 \pm 1.02	2.83 \pm 0.99	t=-1.65	0.101
T stage, n (%)			-	0.799
T1	1 (2.27)	2 (1.68)		
T2	5 (11.36)	15 (12.61)		
T3	21 (47.73)	64 (53.78)		
T4	17 (38.64)	38 (31.93)		
N stage, n (%)			-	0.103
N0	22 (50.00)	76 (63.87)		
N1	5 (11.36)	19 (15.97)		
N2a	1 (2.27)	4 (3.36)		
N2b	7 (15.91)	11 (9.24)		
N2c	9 (20.45)	9 (7.56)		
Grade, n (%)			$\chi^2=5.544$	0.063
High differentiation	2 (4.55)	21 (17.65)		
Moderate differentiation	27 (61.36)	71 (59.66)		
Poor differentiation	15 (34.09)	27 (22.69)		
Primary site, n (%)			-	0.630
Glottis	9 (20.45)	36 (30.25)		
Supraglottis	27 (61.36)	65 (54.62)		
Subglottis	3 (6.82)	8 (6.72)		
Transglottis	5 (11.36)	10 (8.40)		
Surgical excision margins, n (%)			-	0.661
Positive	2 (4.55)	4 (3.36)		
Negative	42 (95.45)	115 (96.64)		
Metastatic, n (%)			$\chi^2=5.409$	0.020
Yes	24 (54.55)	41 (34.45)		
No	20 (45.45)	78 (65.55)		
Extracapsular invasion, n (%)			-	0.093
Yes	8 (18.18)	10 (8.40)		
No	36 (81.82)	109 (91.60)		

Table 3 (continued)

Table 3 (continued)

Variables	Relapse group (n=44)	Non-relapse group (n=119)	Statistic	P
Lymphatic vascular invasion, n (%)			–	0.736
Yes	4 (9.09)	8 (6.72)		
No	40 (90.91)	111 (93.28)		
NLR, M (Q ₁ , Q ₃)	2.64 (1.80, 3.69)	2.12 (1.59, 2.92)	Z=1.927	0.054
PLR, M (Q ₁ , Q ₃)	137.01 (108.62, 179.59)	120.34 (96.89, 149.65)	Z=2.275	0.023
LMR, M (Q ₁ , Q ₃)	3.88 (2.84, 5.46)	4.11 (3.14, 5.56)	Z=-0.716	0.474
AGR, mean ± SD	1.49±0.23	1.60±0.27	t=2.42	0.017
PNI, mean ± SD	421.96±44.43	427.97±37.93	t=0.86	0.393
COP-NLR (score), n (%)			$\chi^2=4.192$	0.041
0	20 (45.45)	77 (64.71)		
1	20 (45.45)	35 (29.41)		
2	4 (9.09)	7 (5.88)		
F-NLR (score), n (%)			$\chi^2=4.242$	0.039
0	30 (68.18)	64 (53.78)		
1	6 (13.64)	11 (9.24)		
2	8 (18.18)	44 (36.97)		

SD, standard deviation; T, tumor; N, node; NLR, neutrophil/lymphocyte ratio; M (Q₁, Q₃), median (interquartile range); PLR, platelet/lymphocyte ratio; LMR, lymphocyte/monocyte ratio; AGR, albumin/globulin ratio; PNI, prognostic nutritional index; COP-NLR, combination of the platelet count and the NLR; F-NLR, combined score of the plasma fibrinogen level and the NLR.

Discussion

Total laryngectomy is commonly performed in locally advanced LSCC patients or with recurrent or persistent cancer after radiation or chemoradiation treatment. However, these patients are still at risk of recurrence after a total laryngectomy. In this study, we used several easily available clinical variables to establish a random-forest model to predict the risk of recurrence of LSCC in patients who have undergone a total laryngectomy. The results showed that certain clinical indicators, including the AGR, NLR, and PLR, were the most important factors in predicting the risk of recurrence of LSCC in patients who have undergone a total laryngectomy. The AUCs of the random-forest model in the training set and the test set were 0.960 and 0.721, respectively.

Several models have been reported to predict the risk of LSCC recurrence in patients. Yang *et al.* established a scoring model based on the 2 independent predictors of CDGSH iron-sulfur domain 2 and N stage (24). Jover-

Esplá *et al.* developed a risk-prediction model based on all laryngeal cancer patients, and showed that age, alcohol consumption, lymph node stage, and stage were associated with a 5-year risk of recurrence (25). Recently, Cui *et al.* (15) developed a nomogram to predict the recurrence risk of LSCC in patients that included 6 factors (i.e., age, tumor site, smoking, alcohol, N stage, and hemoglobin). However, few studies have reported the risk of recurrence in patients with LSCC after a total laryngectomy. In the present study, we used the random-forest method to develop a model to predict the risk of recurrence in LSCC patients who have undergone a total laryngectomy. The AUCs of our predictive model in the training set and the test set were 0.960 and 0.721, which indicated that our model was able to predict the risk of recurrence in LSCC patients after a total laryngectomy well.

Our random-forest model indicated that clinical indicators, such as the AGR, NLR, and PLR, were important predictors for predicting the risk of recurrence of LSCC in patients. Several studies have indicated that

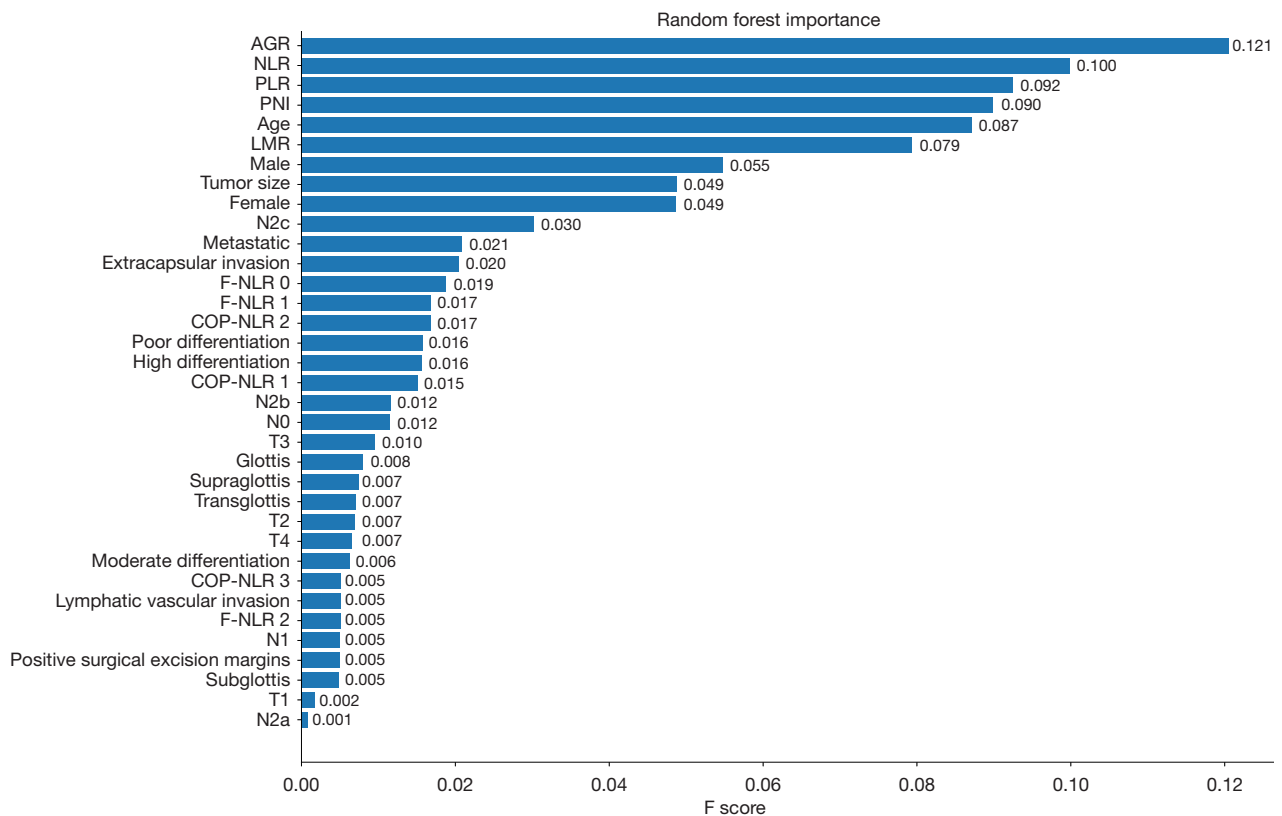


Figure 1 Important predictors of recurrence risk in LSCC patients who have undergone a total laryngectomy. AGR, albumin/globulin ratio; NLR, neutrophil/lymphocyte ratio; PLR, platelet/lymphocyte ratio; PNI, prognostic nutritional index; LMR, lymphocyte/monocyte ratio; F-NLR, combined score of the plasma fibrinogen level and the NLR; COP-NLR, combination of the platelet count and the NLR; N, node; T, tumor; LSCC, laryngeal squamous cell carcinoma.

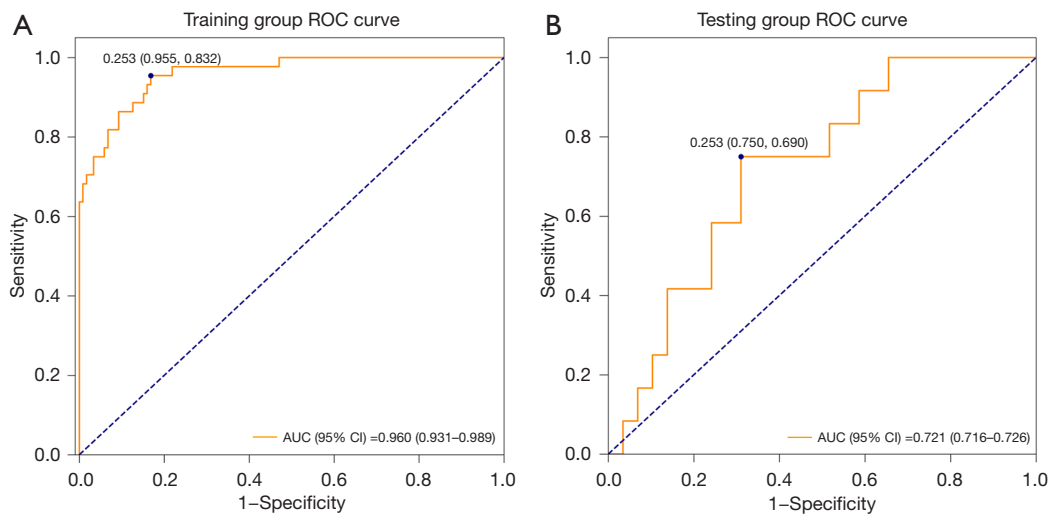


Figure 2 ROC curves with AUC values of the random-forest model. (A) ROC curves in the training set; (B) ROC curves in the test set. ROC, receiver operating characteristic; AUC, area under the ROC curve; CI, confidence interval.

Table 4 The performance of the random-forest model in the training set and the test set

Variables	Training set (95% CI)	Test set (95% CI)
AUC	0.960 (0.931, 0.989)	0.721 (0.716, 0.726)
Accuracy	0.865 (0.813, 0.917)	0.707 (0.568, 0.847)
Specificity	0.832 (0.765, 0.899)	0.690 (0.521, 0.858)
Sensitivity	0.955 (0.893, 1.000)	0.750 (0.505, 0.995)
PPV	0.677 (0.561, 0.948)	0.500 (0.269, 0.921)
NPV	0.980 (0.953, 1.000)	0.870 (0.732, 1.000)

AUC, area under the ROC curve; ROC, receiver operating characteristic; PPV, positive predictive value; NPV, negative predictive value; CI, confidence interval.

some inflammation markers, such as the NLR, PLR, and LMR, are independently correlated with poor outcomes in LSCC patients (16-19). It has been reported that cancer-associated inflammation plays an important role in tumor progression (26,27). The mechanism between inflammation and tumor progression is not clear; however, a possible explanation is that in the early stages of cancer development, various cytokines produced by cancer cells may recruit inflammatory cells that form a microenvironment, promote tumor growth, rheumatic instability, and angiogenesis (28-30). Neutrophils may secrete circulating growth factors to promote cancer cell metastasis (31). Lymphocytes play an important role in inducing cell death and inhibiting tumor cell migration and proliferation. The interaction between platelets and tumor cells can trigger the subsequent metastasis of tumor cells (32). Thus, the NLR and PLR are important factors affecting the prognosis of LSCC patients. Our results also indicated that the NLR and PLR play important roles in predicting the risk of recurrence of LSCC.

Several studies have reported that the AGR has a superior prognostic value in LSCC patients (33,34). A low AGR value indicates a poor prognosis for LSCC patients (33). Alb and globulin are two important components of serum proteins and may be related to systemic inflammation. It has been reported that a low serum Alb level reflects poor nutritional status and is an independent predictor of poor survival in many cancers (35,36). Additionally, an increase in globulin value is associated with a chronic inflammatory response and cumulative exposure to various inflammatory cytokines (37). Thus, the cumulative effect of Alb and globulin may have good prognostic value for LSCC patients. However, apart from the studies of Chen *et al.* (34) and Zhou *et al.* (33), few studies have focused on the prognostic value of the AGR in LSCC patients. In our

prediction model, we found that the AGR was the most important factor for predicting the risk of recurrence of LSCC in patients. In future studies, clinicians should pay attention to the value of some basic clinical features, such as the AGR.

In the current study, a difference analysis was used to evaluate the data of the training set and the test set to ensure the reliability of the model. As described previously, the random-forest method was then used to develop a predictive model. Further, we used variables that are easily available and applicable in clinical practice to establish the model. However, this study had some limitations. First, the sample size of the females was small, which may be because the prevalence ratio of LSCC between males and females is 9.1:1 (38). Second, external validation is needed if the model is to be applied in clinical practice. Third, postoperative adjuvant treatment information such as radiotherapy and chemotherapy were not adjusted as confounders, which may affect the effect of the model.

Conclusions

A risk-prediction model to predict the recurrence risk in LSCC patients who have undergone a total laryngectomy was established, and inflammatory markers AGR, NLR, and PLR play an important role in the predictive model. This model may provide clinicians with a tool to predict recurrence risk in LSCC patients after a total laryngectomy, but external validation is needed before it is used in clinical practice.

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Footnote

Reporting Checklist: The authors have completed the TRIPOD reporting checklist. Available at <https://atm.amegroups.com/article/view/10.21037/atm-22-4802/rc>

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Conflicts of Interest: All authors have completed the ICMJE uniform disclosure form (available at <https://atm.amegroups.com/article/view/10.21037/atm-22-4802/coif>). 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 was conducted in accordance with the Declaration of Helsinki (as revised in 2013). The study was approved by the Ethics Committee of Shanxi Province Cancer Hospital (No. 202130) and individual consent for this retrospective analysis was waived.

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