



The development and validation of a nomogram for predicting brain metastases in lung squamous cell carcinoma patients: an analysis of the Surveillance, Epidemiology, and End Results (SEER) database

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Background: The incidence of brain metastasis (BM) in patients suffering from lung squamous cell carcinoma (LUSC) is lower than that in patients suffering from non-squamous cell carcinoma (NSCC) and there are few studies on BM of LUSC. The purpose of this investigation was to ascertain the risk factors of LUSC, as well as to establish a nomogram prognostic model to predict the incidence of BM in patients with LUSC.

Methods: Patients diagnosed with LUSC between 2010 and 2015 were identified from the Surveillance, Epidemiology, and End Results (SEER) database and the patient data were collated. All patients diagnosed from 2010–2012 were allocated into the training cohort, and the remaining patients diagnosed from 2013–2015 formed the test cohort. Using factors that were screened out through logistic regression analyses, the nomogram in the training cohort was established. It was then evaluated for discrimination and calibration using the test cohort. The performance of the nomogram was assessed by quantifying the area under the receiver operating characteristic (ROC) curve and evaluating the calibration curve.

Results: A total of 26,154 LUSC patients were included in the study. The training cohort consisted of 16,543 patients and there were 8611 patients in the test cohort. Age, marital status, insurance status, histological grade, tumor location, laterality, stage of the cancer, number of metastatic organs, chemotherapy, surgery, and radiotherapy were highly correlated with the incidence of BM. The area under the ROC curve (AUC) of the nomogram for the training cohort and the test cohort were 0.810 [95% confidence interval (CI): 0.796 to 0.823] and 0.805 (95% CI: 0.784 to 0.825), respectively. The slope of the calibration curve was close to 1.

Conclusions: The nomogram was able to accurately predict the incidence of BM. This may be beneficial for the early identification of high-risk LUSC patients and the establishment of individualized treatments.

Keywords: Lung squamous cell carcinoma (LUSC); predicting model; brain metastasis (BM); database

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Introduction

Lung cancer patients who present with brain metastasis (BM) usually have poor prognosis (1-3). These patients place a significant burden on their family caregivers, the nursing staff and the society (4-7). Only early diagnosis and timely therapy can improve their rate of survival (8-11). Currently, the NCCN/ACCP (National Comprehensive Cancer Network/American College of Chest Physicians) does not recommend the screening of asymptomatic patients with stage I non-small cell lung cancer (NSCLC) because stage I NSCLC are less likely to have BM (12,13). However, BM may occur in patients with early-stage NSCLC even in the absence of any neurological symptoms. Whether conventional brain imaging, such as magnetic resonance imaging (MRI) of the brain, should be used in all cases remains controversial (14). If BM can be predicted prior to brain MRI imaging with the use of a nomogram, asymptomatic early-stage NSCLC patients could avoid missed or delayed diagnoses. The incidence of BM is lower in patients with lung squamous cell carcinoma (LUSC) compared to those with non-squamous cell carcinoma (NSCC) and there are few studies on BM of LUSC (15,16). Efficient tools are urgently needed to facilitate the early diagnosis of BM in patients with LUSC (17).

The “seed and soil hypothesis” has been proposed to explain the pathogenesis of BM (18). Nomograms have shown favorable efficiency in predicting risks and outcomes in various cancers, such as breast cancer and intestinal cancer (19). Therefore, in this study, a new nomogram was established to predict the risk of BM in LUSC patients.

We present the following article in accordance with the TRIPOD reporting checklist (available at <http://dx.doi.org/10.21037/jtd-20-3494>).

Methods

Ethical statement

The study was conducted in accordance with the Declaration of Helsinki (as revised in 2013). The data were collected from the Surveillance, Epidemiology, and End Results (SEER) data library. Informed consent was not required as cancer is classified as an informed illness in the United States of America.

Patient selection

The SEER *Stat Software (version 8.3.4) was applied

to create the tabulation of all cases. Patients who were pathologically confirmed with LUSC from the SEER database during 2010–2015 were included. The exclusion criteria included the following: non-first primary LUSC; unknown brain metastases; incomplete data regarding gender, race, marital status, insurance status, grade of tumor, tumor location, laterality, and T or N cancer staging (as defined by the 7th edition of the American Joint Committee on Cancer (AJCC) staging system); bone, liver or lung metastasis; and unknown history of chemotherapy, surgery or radiotherapy. The patients diagnosed with LUSC during 2010–2012 and 2013–2015 exhibited comparable clinical pathological characteristics, therefore the training cohort included patients who were diagnosed during the period from 2010 to 2012, while the test cohort included patients who were diagnosed between 2013 and 2015.

Statistical analysis

The Pearson chi-square test or Fisher’s exact test was used to contrast the variables between the training cohort and the test cohort. Regression analysis was utilized to screen out risk elements of BM that were then further submitted to multivariate analysis. The odds ratio (OR) was expressed as the 95% confidence interval (CI). The nomogram was established using the training cohort data and then discrimination and calibration were evaluated using the test cohort. Variables (such as tumor location) with $P < 0.05$ were incorporated into the nomogram. The likelihood-ratio test was applied to select the proper model.

The discrimination performance of the nomogram was quantified by the area under the receiver operating characteristic (ROC) curve (AUC). By drawing the relationship between the observed and predicted incidences, the nomogram was calibrated with the Hosmer goodness-of-fit measurement. The bootstrap method (1,000 replicates) was employed for the internal verification of performance. Overall checkouts were two sided. Results were considered statistically different when $P < 0.05$. Statistical analysis was performed using the SPSS 3.5.1 software.

Results

Patient characteristics

The study cohort included 26,154 patients diagnosed during 2010–2015, with 17,543 patients in the training cohort and

8611 patients in the test cohort. Most patients were elderly whose average age was 67. The clinical characteristics in both cohorts are shown in *Table 1*. The median survival of LUSC patients with BM was evidently shorter than LUSC patients without BM. Similarly, patients with BM had significantly lower lung cancer-specific survival (LSCC) and overall survival (OS) compared to patients without BM (*Figure 1*).

Factors associated with BM

Univariate Cox regression analysis was conducted in the training cohort. As shown in *Table 2*, gender and race showed no significant relationship with BM. However, the factors most closely associated with BM were age at the

time of diagnosis, marital status, insurance, microanatomy grade, tumor location, laterality, number of organs with metastases, chemotherapy, surgery, and radiation. These factors were then submitted to multivariate Cox logistic regression analysis (*Table 3*). The outcomes demonstrated that age at the time of diagnosis, tumor grade, tumor stage, number of organs with metastases, chemotherapy, surgery, and radiation were independent factors for predicting BM. Therefore, these elements were used to establish the nomogram prognostic model. Tumor location ($P<0.07$) was also included (*Table 3*).

Construction of the nomogram

The nomogram was established by incorporating age at

Table 1 Patient demographics and clinical characteristics

Characteristics	All patients		Training group		Validation group	
	n=26,154	%	n=17,543	%	n=8,611	%
All cause	15,762		12,525		3,237	
LCSS	13,193		10,431		2,762	
Age at diagnosis (year)						
<50	603	2.3	435	2.5	168	2
50–70	11,940	45.7	8,090	46.1	3,850	44.7
>70	13,611	52	9,018	51.4	4,593	53.3
Gender						
Male	16,262	62.2	10,953	62.4	5,309	61.7
Female	9,892	37.8	6,590	37.6	3,302	38.3
Race						
White	21,915	83.3	14,745	84.1	7,170	83.3
Black	2,939	11.2	1,950	11.1	989	11.5
Other (American Indian/Alaskan Native, Asian/Pacific Islander)	1,300	5	848	4.8	452	5.2
Marital status						
Single	3,599	13.8	2,340	13.3	1,259	14.6
Married	13,083	50	8,833	50.4	4,250	49.4
Other	9,472	36.2	6,370	36.3	3,102	36
Insurance						
No	895	3.4	671	3.8	224	2.6
Yes	25,259	96.6	16,872	96.2	8,387	97.4

Table 1 (continued)

Table 1 (continued)

Characteristics	All patients		Training group		Validation group	
	n=26,154	%	n=17,543	%	n=8,611	%
Grade						
Grade I	804	3.1	526	3	278	3.2
Grade III	11,176	42.7	7,482	42.5	3,694	42.9
Grade III	13,948	53.3	9,384	53.5	4,564	53
Grade IV	226	0.9	151	0.9	75	0.9
Tumor location						
Upper lobe	14,690	56.2	9,873	56.3	4,817	55.9
Middle lobe	980	3.7	646	3.7	334	3.9
Lower lobe	7,812	29.9	5,217	29.7	2,595	30.1
Main bronchus	1,346	5.1	896	5.1	450	5.2
Other sites	1,326	5.1	911	5.2	415	4.8
Laterality						
Left	11,388	43.5	7,698	43.9	3,690	42.9
Right	14,659	56	9,779	55.7	4,880	56.7
Bilateral, single primary	107	0.4	66	0.4	41	0.5
T stage						
T1	5,236	20	3,484	19.9	1,752	20.3
T2	9,314	35.6	6,283	35.8	3,031	35.2
T3	6,121	23.4	4,149	23.7	1,972	22.9
T4	5,483	21	3,627	20.7	1,856	21.6
N stage						
N0	12,994	49.7	8,645	49.3	4,349	50.5
N1	2,896	11.1	1,995	11.4	901	10.5
N2	8,089	30.9	5,465	31.2	2,624	30.5
N3	2,175	8.3	1,438	8.2	737	8.6
Brain metastasis						
No	25,058	95.8	16,812	95.8	8,246	95.8
Yes	1,096	4.2	731	4.2	365	4.2
Number of organs with metastases besides the brain						
0	21,908	83.8	14,738	84	7,170	83.3
1	3,248	12.4	2,146	12.2	1,102	12.8
2	841	3.2	558	3.2	283	3.3
3	157	0.6	101	0.6	56	0.7

Table 1 (continued)

Table 1 (continued)

Characteristics	All patients		Training group		Validation group	
	n=26,154	%	n=17,543	%	n=8,611	%
Bone metastasis						
No	24,140	92.3	16,217	92.4	7,924	92
Yes	2,014	7.7	1,326	7.6	688	8
Liver metastasis						
No	25,102	96	16,841	96	8,261	95.9
Yes	1,052	4	702	4	350	4.1
Lung metastasis						
No	23,819	91.1	16,006	91.2	7,813	90.7
Yes	2,335	8.9	1,537	8.8	798	9.3
Chemotherapy						
No	15,158	58	10,213	58.2	4,945	57.4
Yes	10,996	42	7,330	41.8	3,666	42.6
Surgery						
No	16,344	62.5	10,751	61.3	5,593	65
Yes	9,810	37.5	6,792	38.7	3,018	35
Radiotherapy						
No	14,736	56.3	9,974	56.9	4,762	55.3
Yes	11,418	43.7	7,569	43.1	3,849	44.7

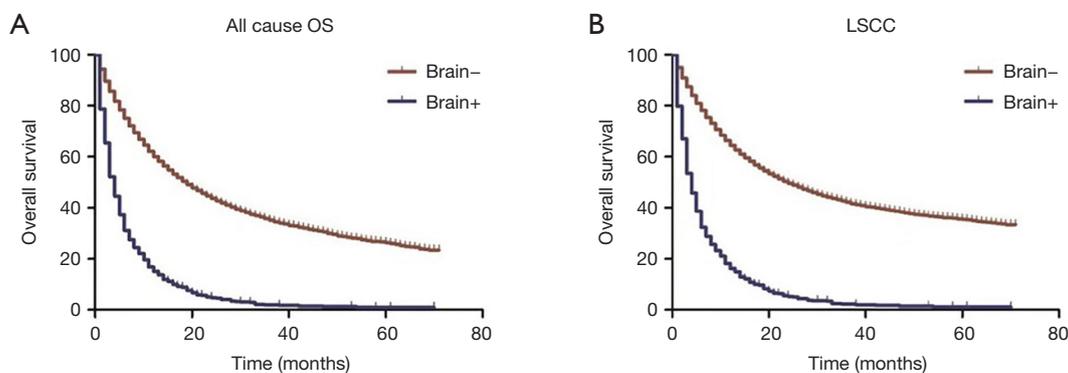


Figure 1 Kaplan-Meier survival curves. (A) Kaplan-Meier survival curves for LUSC patients showing overall survival. (B) Kaplan-Meier survival curves for LUSC patients showing lung cancer specific survival. Log-rank P values were both <0.0001. LUSC, lung squamous cell carcinoma.

diagnosis, grade, the location of tumor, T and N stage, number of metastases, chemotherapy, surgery, and radiation (Figure 2). The AUC of the nomogram was 0.810 (95% CI: 0.796 to 0.823) and 0.805 (95% CI: 0.784 to 0.825) in the

training cohort and the test cohort, respectively, suggesting a good predictive ability (Figure 3A,B). The calibration curve showed a high consistency between the observed and predicted survival (Figure 3C,D).

Table 2 Univariate Cox logistic regression analysis for both cohorts

Characteristics	Training group, no. (%)			Validation group, no. (%)		
	B- (16,812)	B+ (731)	P	B- (8,246)	B+ (365)	P
Age at diagnosis (year)			<0.0001			<0.0001
<50	394 (2.3)	41 (5.6)		152 (1.8)	16 (4.4)	
50–70	7,695 (45.8)	395 (54.0)		3,645 (44.2)	205 (56.2)	
>70	8,723 (51.9)	295 (40.4)		4,449 (54.0)	144 (39.5)	
Gender			0.584			0.660
Male	10,489 (62.4)	464 (63.5)		5,088 (61.7)	221 (60.5)	
Female	6,323 (37.6)	267 (36.5)		3,158 (38.3)	144 (39.5)	
Race			0.198			0.013
White	14,148 (84.2)	597 (81.7)		6,885 (83.5)	285 (78.1)	
Black	1,856 (11.0)	94 (12.9)		930 (11.3)	59 (16.2)	
Other (American Indian/Alaskan Native, Asian/Pacific Islander)	808 (4.8)	40 (5.5)		431 (5.2)	21 (5.8)	
Marital status			0.004			0.309
Single	2,216 (13.2)	124 (17.0)		1,197 (14.5)	62 (17.0)	
Married	8,461 (50.3)	372 (50.9)		4,068 (49.3)	182 (49.9)	
Other	6,135 (36.5)	235 (32.1)		2,981 (36.2)	121 (33.2)	
Insurance			0.008			0.004
No	629 (3.7)	42 (5.7)		205 (2.5)	19 (5.2)	
Yes	16,183 (96.3)	689 (94.3)		8,041 (97.5)	346 (94.8)	
Grade			<0.0001			<0.0001
Well differentiated; Grade I	511 (3.0)	15 (2.1)		272 (3.3)	6 (1.6)	
Poorly differentiated; Grade III	7,278 (43.3)	204 (27.9)		3,580 (43.4)	114 (31.2)	
Poorly differentiated; Grade III	8,883 (52.8)	501 (68.5)		4,323 (52.4)	241 (66.0)	
Undifferentiated; Grade IV	140 (0.8)	11 (1.5)		71 (0.9)	4 (1.1)	
Tumor location			0.008			0.013
Upper lobe	9,486 (56.4)	387 (52.9)		4,620 (56.0)	197 (54.0)	
Middle lobe	612 (3.6)	34 (4.7)		319 (3.9)	15 (4.1)	
Lower lobe	5,009 (29.8)	208 (28.5)		2,497 (30.3)	98 (26.8)	
Main bronchus	846 (5.0)	50 (6.8)		425 (5.2)	25 (6.8)	
Other sites	859 (5.1)	52 (7.1)		385 (4.7)	30 (8.2)	
Laterality			0.024			0.003
Left	7,389 (44.0)	309 (42.3)		3,542 (43.0)	148 (40.5)	
Right	9,364 (55.7)	415 (56.8)		4,669 (56.6)	211 (57.8)	
Bilateral, single primary	59 (0.4)	7 (1.0)		35 (0.4)	6 (1.6)	

Table 2 (continued)

Table 2 (continued)

Characteristics	Training group, no. (%)			Validation group, no. (%)		
	B- (16,812)	B+ (731)	P	B- (8,246)	B+ (365)	P
T stage			<0.0001			<0.0001
T1	3,446 (20.5)	38 (5.2)		1,730 (21.0)	22 (6.0)	
T2	6,049 (36.0)	234 (32.0)		2,933 (35.6)	98 (26.8)	
T3	3,927 (23.4)	222 (30.4)		1,881 (22.8)	91 (24.9)	
T4	3,390 (20.2)	237 (32.4)		1,702 (20.6)	154 (42.2)	
N stage			<0.0001			<0.0001
N0	8,476 (50.4)	169 (23.1)		4,273 (51.8)	76 (20.8)	
N1	1,920 (11.4)	75 (10.3)		860 (10.4)	41 (11.2)	
N2	5,091 (30.3)	374 (51.2)		2,440 (29.6)	184 (50.4)	
N3	1,325 (7.9)	114 (15.5)		673 (8.2)	64 (17.5)	
Number of organs with metastases besides the brain			<0.0001			<0.0001
0	14,320 (85.2)	418 (57.2)		6,985 (84.7)	185 (50.7)	
1	1,944 (11.5)	202 (27.6)		989 (12.0)	113 (31.0)	
2	473 (2.8)	85 (11.6)		233 (2.8)	50 (13.7)	
3	75 (0.4)	26 (3.6)		39 (0.5)	17 (4.7)	
Bone metastasis			<0.0001			<0.0001
No	15,662 (93.2)	555 (75.9)		7,654 (92.8)	269 (73.7)	
Yes	1,150 (6.8)	176 (24.1)		592 (7.2)	96 (26.3)	
Liver metastasis			<0.0001			<0.0001
No	16,212 (96.4)	629 (86.0)		7,961 (96.5)	300 (82.2)	
Yes	600 (3.6)	102 (14.0)		285 (3.5)	65 (17.8)	
Lung metastasis			<0.0001			<0.0001
No	15,447 (91.9)	559 (76.5)		7,551 (91.6)	262 (71.8)	
Yes	1,365 (8.1)	172 (23.5)		695 (8.4)	103 (28.2)	
Chemotherapy			<0.0001			<0.0001
No	9,836 (58.5)	377 (51.6)		4,771 (57.9)	174 (47.7)	
Yes	6,976 (41.5)	354 (48.4)		3,475 (42.1)	191 (52.3)	
Surgery			<0.0001			<0.0001
No	10,063 (59.9)	688 (94.1)		5,247 (63.6)	346 (94.8)	
Yes	6,749 (40.1)	43 (5.9)		2,999 (36.4)	19 (5.2)	
Radiotherapy			<0.0001			<0.0001
No	9,794 (58.3)	180 (24.6)		4,663 (56.5)	99 (27.1)	
Yes	7,018 (41.7)	551 (75.4)		3,583 (43.5)	266 (72.9)	

B-, absence of brain metastasis; B+, presence of brain metastasis.

Table 3 Multivariate Cox logistic regression analysis to discriminate risk elements of LUSC in the training cohort

Variables	β	Odds ratio	95% CI	P
Age at diagnosis	-0.033	0.97	0.96-0.98	<0.0001
Race				
Black	Reference			
White	0.085	1.09	0.86-1.38	0.480
Other	0.148	1.16	0.78-1.73	0.469
Gender				
Female	Reference			
Male	-0.113	0.89	0.76-1.05	0.183
Marital status				
Married	Reference			
Single	-0.016	0.98	0.78-1.24	0.894
Other	-0.164	0.85	0.71-1.02	0.073
Insurance				
No	Reference			
Yes	-0.023	0.98	0.69-1.38	0.897
Grade				
Grade I	Reference			
Grade II	-0.035	0.97	0.56-1.67	0.901
Grade III	0.495	1.64	0.96-2.81	0.071
Grade IV	0.896	2.45	1.06-5.68	0.037
Tumor location				
Lower lobe	Reference			
Upper lobe	-0.164	0.85	0.71-1.02	0.075
Middle lobe	0.238	1.27	0.85-1.89	0.241
Main bronchus	-0.302	0.74	0.53-1.03	0.076
Other sites	-0.055	0.95	0.68-1.32	0.748
Laterality				
Bilateral, single primary	Reference			
Left	-0.231	0.79	0.34-1.84	0.591
Right	-0.244	0.78	0.34-1.82	0.569
T stage				
T1	Reference			
T2	0.810	2.25	1.57-3.22	<0.0001
T3	0.830	2.29	1.59-3.30	<0.0001
T4	0.706	2.03	1.40-2.93	0.000

Table 3 (continued)

Table 3 (continued)

Variables	β	Odds ratio	95% CI	P
N stage				
N0	Reference			
N1	0.474	1.61	1.20–2.14	0.001
N2	0.479	1.61	1.32–1.98	<0.0001
N3	0.360	1.43	1.09–1.88	0.009
Number of metastases besides the brain	0.663	1.94	1.75–2.15	<0.0001
Chemotherapy				
No	Reference			
Yes	-0.614	0.54	0.46–0.64	<0.0001
Surgery				
No				
Yes	-1.492	0.22	0.16–0.31	<0.0001
Radiation				
No				
Yes	1.171	3.23	2.67–3.90	<0.0001

LUSC, lung squamous cell carcinoma; β is the regression coefficient.

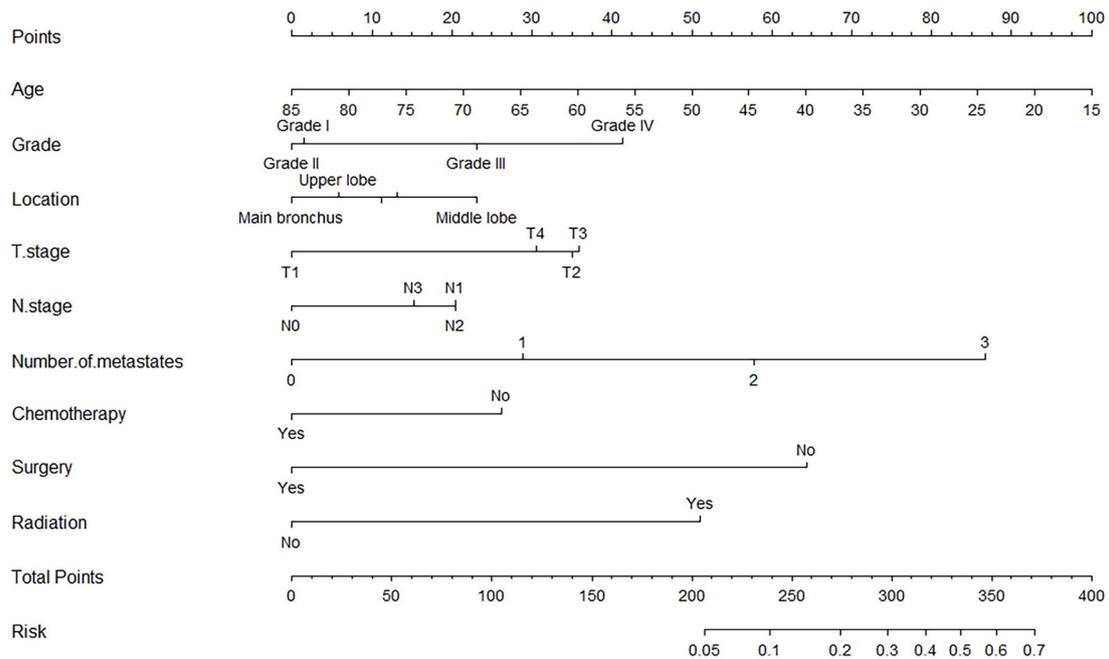


Figure 2 Nomogram predicting BM in patients with SCC. The first row illustrates the point allocation of every variable. Lines 2–10 represent the alternating quantity contained in the nomogram. For each patient, each variable of LUSC was given a score. The scores were totaled and illustrated in line 11. The bottom row presents the probability of BM. BM, brain metastasis; SCC, squamous cell carcinoma; LUSC, lung cancer specific survival.

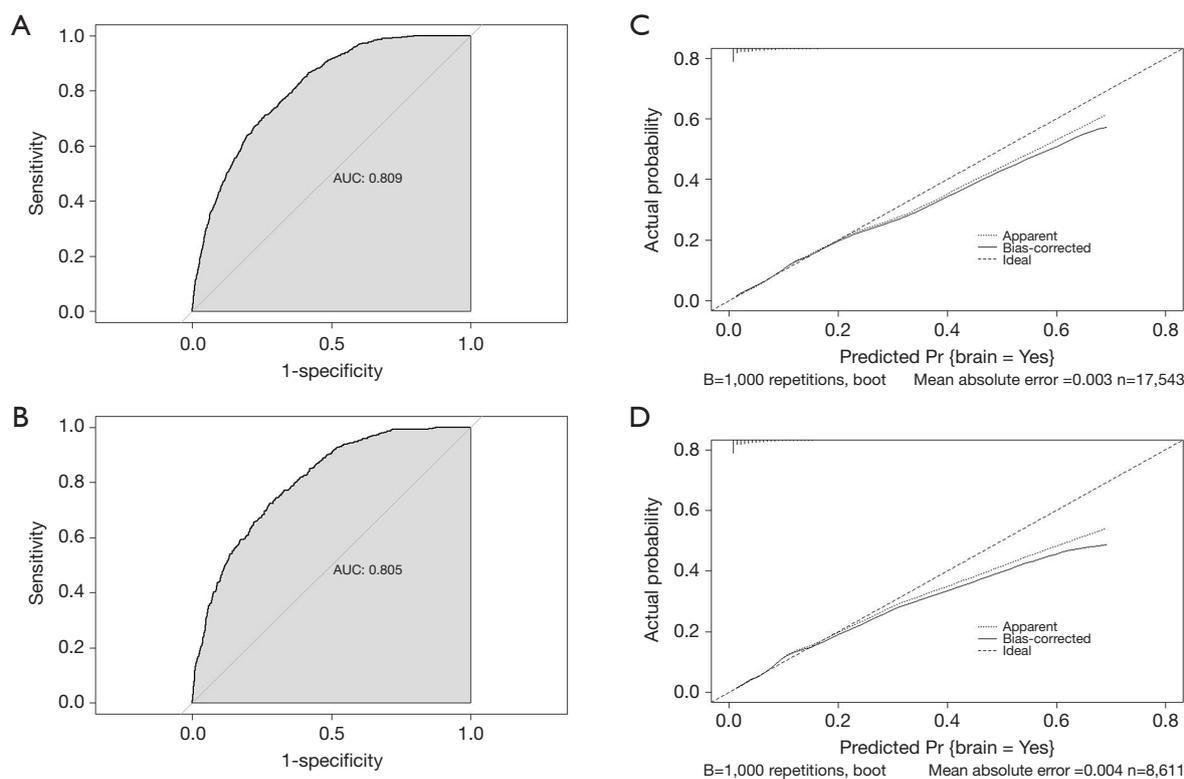


Figure 3 Discrimination and calibration of the nomogram. (A) and (B) ROC for discrimination. The AUC of the nomogram is 0.810 (95% CI: 0.796 to 0.823) and 0.805 (95% CI: 0.784 to 0.825) in the training cohort and the test cohort, respectively. (C) and (D) Calibration of the nomogram. The x-axis represents the BM probabilities predicted by the nomogram, while the y-axis represents the observed probabilities. The plummet lines express the frequency distribution of the forecast probability. The dotted line represents the match between two probabilities. The calibration curve shows good predictive ability of the nomogram. ROC, receiver operating characteristic; AUC, area under the curve; CI, confidence interval; BM, brain metastasis.

Discussion

Our research is the first to report a nomogram for predicting BM in patients with LUSC. This nomogram provides an efficient and convenient tool for early diagnosis of BM thereby allowing for the implementation of timely treatment strategies. Previous studies have demonstrated similar BM pathogenesis in patients with LUSC and lung adenocarcinoma (LUAD) (20-23). Muniz *et al.* found that the incidence of BM increased with age in patients with LUSC and LUAD (24), which is consistent with our results. Univariate and multivariate Cox logistic regression analyses demonstrated that age at diagnosis, tumor grade, location of tumor, tumor stage, number of organs with metastases, chemotherapy, surgery, and radiation were independent factors that could predict the incidence of BM. These factors were used to establish a prediction nomogram. The

predictive accuracy was validated, with AUC of 0.810 (95% CI: 0.796 to 0.823) and 0.805 (95% CI: 0.784 to 0.825) in the training cohort and the test cohort, respectively. These results support the potential use of this nomogram in clinical practice, especially in the establishment of individualized treatments.

However, due to the retrospective nature of this study, selection bias may exist. As new treatments continue to be developed for the improved survival of LUSC patients, this current nomogram should be renewed with additional clinical data, such as types of surgical treatment, radiotherapy dosage, and chemotherapy regimens. Furthermore, due to insufficient data, our nomogram could only be validated internally (19,25) and hence, validation with larger data cohorts and longer follow-up times will be required.

Conclusions

The nomogram could accurately predict the incidence of BM, which is helpful in the early identification of high-risk LUSC patients and the establishment of individualized treatments.

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Footnote

Reporting Checklist: The authors have completed the TRIPOD reporting checklist. Available at <http://dx.doi.org/10.21037/jtd-20-3494>

Conflicts of Interest: All authors have completed the ICMJE uniform disclosure form (available at <http://dx.doi.org/10.21037/jtd-20-3494>). 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 institutional review board approval was waived for this study as the SEER database is a public anonymized database. The author Zhang obtained access to the SEER database (accession number: 13013-Nov2019). The data issued in the SEER database do not require consent from patients as cancer is an informed illness in every state of the United States of America.

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