

Establishment and validation of prognostic nomograms to predict overall survival and cancer-specific survival for patients with osteosarcoma

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This study aimed to develop and validate nomograms predicting the survival of osteosarcoma patients from the SEER database and our hospital. Data of 1,066 osteosarcoma patients from the SEER database were randomly divided into a development cohort (n=800) and validation cohort one (n=266). Another cohort of 126 patients from our hospital was utilized as validation cohort two. Univariate and multivariate Cox analyses were performed to identify the independent prognostic factors for overall survival (OS) and cancer-specific survival (CSS). Nomograms predicting the 3- and 5-year OS and CSS probability were constructed and validated. The predictive performances of the established nomograms were evaluated by the concordance index (C-index) and the calibration plot. Variables of age, surgical stage, surgery, grade, tumor site, and tumor size were identified as independent prognosticators for OS and CSS in Cox analyses. The C-indexes for OS and CSS in the development cohort were 0.818 and 0.829. Comparatively, the C-indexes for OS and CSS were 0.843 and 0.834, 0.736 and 0.782 for validation cohort one and two, respectively. Calibration plots showed excellent consistency between nomogram prediction and actual survival. Nomograms based on the SEER database are of high accuracy and can serve as a reliable tool for individualized consultation and survival prediction in osteosarcoma patients.

Key words: nomogram, osteosarcoma, overall survival, cancer-specific survival

Osteosarcoma is the most common primary malignancy of bones which accounts for 35% of all primary malignant bone tumors, and it typically arises in children and adolescents younger than 24 years of age, with an estimated incidence of 0.34/100,000 per year [1, 2]. Prior to the introduction of chemotherapy, amputation was the main therapeutic measure for osteosarcoma patients but with low quality of life and unfavorable survival rates. In addition, 80 to 90 percent of osteosarcoma patients developed metastatic disease despite the achievement of local tumor control and finally died of metastases. It was surmised and subsequently demonstrated that the vast majority of osteosarcoma patients had developed the subclinical metastatic disease at the time of diagnosis, despite the absence of overt clinical symptoms [3]. Currently, neoadjuvant chemotherapy followed by surgical resection of the primary lesions has been established and demonstrated as the standard therapeutic strategy for newly diagnosed non-metastatic osteosarcoma [4]. With

multi-disciplinary treatments, the 5-year survival rate for these patients has risen from less than 20% to approximately 70% [5]. Nevertheless, available and effective options are still insufficient and the outcomes are still poor for metastatic and recurrent osteosarcoma patients [6, 7]. Better comprehension and identification of prognostic factors of osteosarcoma can provide us more information to select therapeutic interventions, which will contribute to improvement in the quality of life and prolonging survival in patients with osteosarcoma.

A series of factors have been reported in the literature to have predictive or prognostic values for the survival of patients with osteosarcoma. These predictive factors can be roughly divided into seven groups based on their characteristics, including 1) inflammation-related predictive variables, such as serum C-reactive protein (CRP), pre-treatment neutrophil-to-lymphocyte ratio (Pre-NLR), absolute lymphocyte count (ALC), Glasgow prognostic score (GPS), platelet-lymphocyte ratio (PLR), lymphocyte-monocyte ratio

(LMR), and neutrophil-platelet score (NPS) [8–10]; 2) nutrition-related predictive factors or scores, such as body mass index (BMI), prognostic nutritional index (PNI), Geriatric Nutritional Risk Index (GNRI) [11–13]; 3) aberrant expression of specific proteins, such as cell-cycle kinase inhibitor p27, cyclin-dependent protein kinase 9 (CDK9), motility-related protein-1 (MRP-1)/CD9 [14–16]; 4) upregulation or downregulation of microRNAs, non-coding RNA, and circular RNAs, such as serum microRNA-375, microRNA-17, long non-coding RNA HAGLROS, circulating hsa_circ_0081001 [17–21]; 5) abnormalities of constituents in the tumor microenvironment, such as tumor-infiltrating macrophages, CD8-positive cytotoxic lymphocytes, immune infiltrations, PD-L1 expression, dysregulation of M1/M2 macrophages ratio [22–25]; 6) features of the disease, such as TNM stage, AJCC stage, lymph node involvement, histopathologic features, metastasis, age at diagnosis, tumor grade, tumor site, surgical margins, pathological fractures, etc. [26–32]; 7) predictors from imaging examination, such as maximum and peak standardized uptake value (SUVmax, SUVpeak) on 18F-fluoro-2-deoxyglucose positron emission tomography/computed tomography (18FDG-PET/CT), kinetic parameters of dynamic contrast-enhanced magnetic resonance imaging (DCE-MRI), including the influx volume transfer constant [K(trans)], the relative extravascular extracellular space [v(e)], the relative vascular plasma space [v(p)] and the efflux rate constant [k(ep)] [33–35]. As reported in previous studies, these variables are significantly associated with the survival of osteosarcoma patients in statistical analyses. However, the survival of an osteosarcoma patient is determined by multiple factors, and therefore single predictor-based survival prediction is usually inaccurate and unreliable. On account of this fact, it is reasonable to construct a new evaluation system, which can integrate these pivotal prognostic factors together.

Nomogram is a newly developed prediction tool that has been widely used to estimate the survival probability in a variety of tumors, including lung cancer, breast cancer, and gastric cancer [36–38]. A predictive nomogram is an ocular, reliable, and effective tool based on multivariate regression models, which can provide intuitive and graphic calculating scales to estimate the survival probability of patients with osteosarcoma. The outstanding characteristics of nomogram in incorporating these key prognostic factors together significantly improve the accuracy and reliability in estimating medium- and long-term survival, as well as risk stratification of osteosarcoma patients.

The Surveillance, Epidemiology, and End Results (SEER) dataset is an open, updated database, which has collected anonymized clinical information of multiple malignancies from 1983 to now [39]. SEER database has covered approximately 30% of the overall US population and is composed of eighteen cancer registries [40]. In this study, we first established effective prognostic nomograms to estimate the 3- and 5-year overall survival (OS) and cancer-specific survival

(CSS) rates for patients with osteosarcoma based on the data from the SEER database. Then, we validated the developed nomograms internally and externally with data from the SEER database and our hospital.

Patients and methods

Data source. Two datasets, which respectively came from the SEER database (patients diagnosed with osteosarcoma from 2010 to 2015) and Cancer Hospital of China Medical University (patients diagnosed with osteosarcoma between 2010 to 2016) were utilized for this study. The SEER*Stat software (version 8.3.6; NCI, Bethesda, MD, USA) was used to extract the information of each osteosarcoma patient from the SEER database. All the data are anonymous, and therefore the requirement for informed consent was waived.

Inclusive and exclusive criteria for osteosarcoma patients. The inclusion criteria were as follows: 1) diagnosed with osteosarcoma as primary malignancy (the International Classification of Disease for Oncology [ICD-O] were 9180-9187, 9192-9194 and 9200); 2) diagnosed between the year of 2010 to 2015; 3) the primary lesion sites were limited to extremities (long or short bones of the lower or upper extremities) and axial bones (including skull, spine, ribs, sternum, or pelvis); 4) confirmation of histologic subtypes of osteosarcoma; 5) with detailed information for race, surgical stage, surgery, histologic grade, AJCC stage; 6) with known survival months, survival status, and cause of death.

The exclusion criteria included: 1) no information for race, surgery, and TNM stage; 2) unknown or unspecific AJCC stage; 3) no detailed histologic grade; 4) unknown tumor size or surgical stage. The study design and process of collecting patients were shown in a flow diagram (Figure 1).

Variables included in this study. The potential prognostic variables for this study mainly included age, gender, race, tumor site, surgical stage, surgery, histologic grade, histologic subtypes, AJCC stage, tumor size, survival months, survival status, and cause of death. The X-tile software (Yale University, New Haven, CT, USA) was used to identify the optimal cut-off values to categorized continuous variables of age and tumor size.

Statistical analysis. X-tile software was used to categorize continuous variables of age and tumor size. Categorical variables were presented as frequencies and percentages. Chi-square test or Fisher exact test was used to compare the differences of variables of cohorts. The primary endpoints for the current study were overall survival (OS) and cancer-specific survival (CSS). OS was defined as the time from diagnosis to death from any cause, and CSS was calculated from the time of diagnosis to death attributed to osteosarcoma only. A two-sided p-value <0.05 was considered as statistically significant.

Identification of prognosticators for survival. Univariate and multivariate Cox proportional hazards regression analyses were used to identify independent prognostic factors

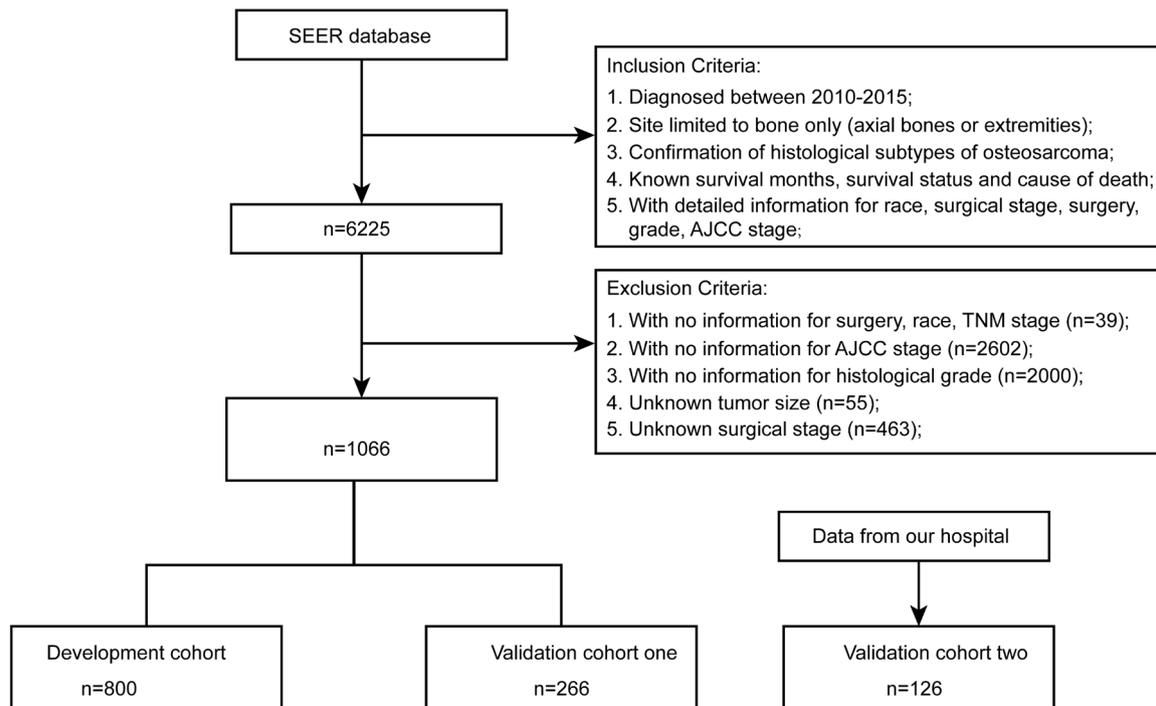


Figure 1. The flow chart of the inclusion and exclusion criteria of this study.

for OS and CSS from the potential prognostic factors. The hazard ratio and its corresponding 95% confidence interval of each variable were also calculated and presented.

Establishment and validation of predictive models for OS and CSS. The prognostic nomograms for 3- and 5-year OS and CSS were established based on the prognosticators determined in univariate and multivariate Cox analyses. Internal validation (data from the development cohort) and external validation (data from validation cohort one and two) were performed with 500 bootstrap resamples to prevent overfitting and to get a relatively unbiased estimation. Harrell's concordance-index (C-index) was utilized to evaluate the performance of these established predictive nomograms. Calibration curves were also constructed to assess the consistency between predicted and actual survival internally and externally.

Software for statistical analysis. Several pivotal softwares were utilized in our study, including SEER*Stat software (Version 8.3.6; NCI, Bethesda, USA), IBM SPSS Statistics 24 (IBM Corp, Armonk, NY, USA), and R software (version 3.6.0). SEER*Stat software was used to extract the information of each osteosarcoma patient from the SEER database. Chi-squared tests, Fisher exact tests, univariate and multivariate Cox analyses were conducted by SPSS. R software was used to establish nomograms and calibration plots with the help of some essential packages, such as rms, foreign, survival, etc.

Results

Baseline characteristics of osteosarcoma patients. A total of 6,225 osteosarcoma patients were preliminarily collected from the SEER database between 2010 to 2015. Based on the inclusion and exclusion criteria, an entire cohort of 1,066 osteosarcoma patients was finally enrolled for this study, and they were further divided into the development cohort and the validation cohort one. Another cohort of 126 osteosarcoma patients from our hospital was used as the validation cohort two (Figure 1).

The optimal cut-off values for age and tumor size identified by X-tile software were 26 and 61 years, and 9.6 and 20.5 cm, respectively (Figures 2A–2D). The details of the clinicopathological features of all patients included in this study are presented in Table 1. No significant differences in clinicopathological features were observed between the development cohort and the two validation cohorts.

Identification of prognostic factors for OS and CSS. The dataset from the development cohort was used to determine the independent prognostic factors for OS and CSS. In univariate analyses, variables of age, gender, tumor site, surgical stage, surgery, grade, histology, AJCC stage, node involvement, metastasis, tumor size were all significantly associated with OS (all $p < 0.05$). These factors were then selected to perform multivariate Cox analysis. In multivariate Cox analysis, variables of age, surgical stage, surgery, grade,

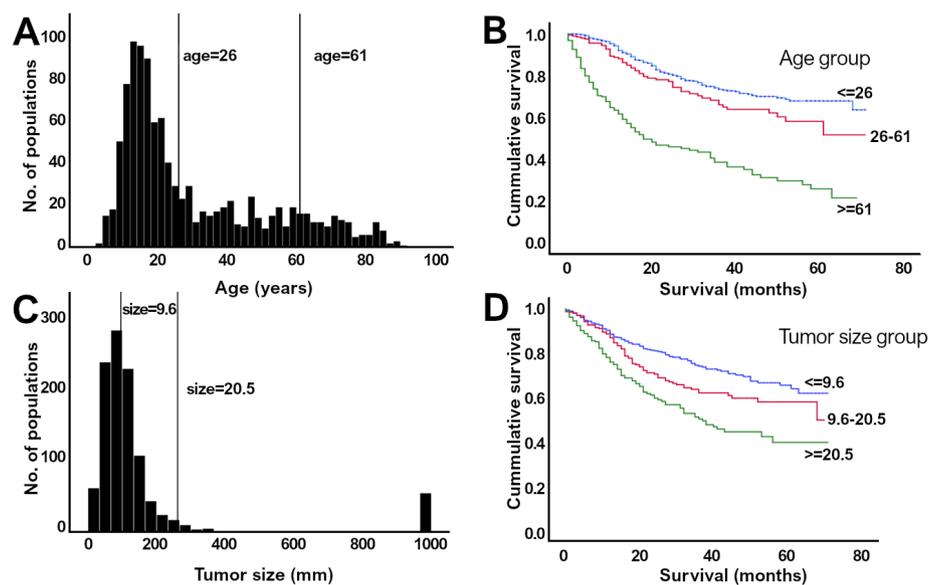


Figure 2. The optimal cut-off values of age at diagnosis (A), and tumor size (C) identified by the X-tile software. The survival curve analysis for age (B), and tumor size (D) based on the optimal cut-off values groupings.

Table 1. Baseline characteristics of all patients in this study.

Variables	Total cohort n=1192 No. (%)	Development cohort ^a n=800 No. (%)	Validation cohort one ^a n=266 No. (%)	Validation cohort two ^b n=126 No. (%)	p-value
Age					0.500
≤26	716 (60.1%)	478 (59.8%)	166 (62.4%)	72 (57.1%)	
26–61	316 (26.5%)	220 (27.5%)	60 (22.6%)	36 (28.6%)	
≥61	160 (13.4%)	102 (12.7%)	40 (15.0%)	18 (14.3%)	
Median (range)	16 (3–94)	17 (3–93)	17 (3–94)	18 (4–78)	
Gender					0.259
Male	656 (55.0%)	433 (54.1%)	145 (54.4%)	78 (61.9%)	
Female	536 (45.0%)	367 (45.9%)	121 (45.5%)	48 (38.1%)	
Race					0.192
Black	260 (21.8%)	160 (19.9%)	72 (27.1%)	28 (22.2%)	
White	881 (73.9%)	606 (75.8%)	182 (68.4%)	93 (73.8%)	
Other	51 (4.3%)	34 (4.3%)	12 (4.5%)	5 (4.0%)	
Tumor site					0.228
Axial	302 (25.3%)	207 (25.9%)	58 (21.8%)	37 (29.4%)	
Extremity	890 (74.7%)	593 (74.1%)	208 (78.2%)	89 (70.6%)	
Surgical stage					0.757
Regional	514 (43.1%)	337 (42.1%)	116 (43.6%)	61 (48.4%)	
Localized	400 (33.6%)	275 (34.4%)	87 (32.7%)	38 (30.2%)	
Distant	218 (23.3%)	188 (23.5%)	63 (23.7%)	27 (21.4%)	
Surgery					0.884
NO	137 (11.5%)	92 (11.5%)	32 (12.0%)	13 (10.3%)	
YES	1055 (88.5%)	708 (88.5%)	234 (88.0%)	113 (89.7%)	
Grade					0.456
Well differentiated	63 (5.3%)	41 (5.1%)	13 (4.9%)	9 (7.1%)	
Moderately differentiated	85 (7.1%)	60 (7.5%)	17 (6.4%)	8 (6.3%)	
Poorly differentiated	396 (33.2%)	261 (32.6%)	84 (31.6%)	51 (40.5%)	
Undifferentiated/anaplastic	648 (54.4%)	438 (54.8%)	152 (57.1%)	58 (46.0%)	

Table 1. Continued ...

Variables	Total cohort n=1192 No. (%)	Development cohort ^a n=800 No. (%)	Validation cohort one ^a n=266 No. (%)	Validation cohort two ^b n=126 No. (%)	p-value
Histology					0.112
Conventional osteosarcoma	783 (65.7%)	538 (67.3%)	161 (60.5%)	84 (66.7%)	
Chondroblastic osteosarcoma	185 (15.5%)	130 (16.3%)	36 (13.5%)	19 (15.1%)	
Fibroblastic osteosarcoma	36 (3.0%)	24 (3.0%)	10 (3.8%)	2 (1.6%)	
Telangiectatic osteosarcoma	29 (2.4%)	18 (2.3%)	9 (3.4%)	2 (1.6%)	
Osteosarcoma in Paget disease of bone	3 (0.3%)	3 (0.4%)	0 (0%)	0 (%)	
Small cell osteosarcoma	12 (1.0%)	6 (0.8%)	5 (1.9%)	1 (0.8%)	
Central osteosarcoma	60 (5.0%)	30 (3.8%)	20 (7.5%)	10 (7.9%)	
Intraosseous well differentiated osteosarcoma	2 (0.2%)	0 (0%)	2 (0.8%)	0 (0%)	
Parosteal osteosarcoma	63 (5.3%)	40 (5.0%)	17 (6.4%)	6 (4.8%)	
Periosteal osteosarcoma	12 (1.0%)	6 (0.8%)	5 (1.9%)	1 (0.8%)	
High grade surface osteosarcoma	7 (0.6%)	5 (0.6%)	1 (0.4%)	1 (0.8%)	
AJCC stage					0.788
I+II	930 (78.0%)	624 (78.0%)	205 (77.1%)	101 (80.2%)	
III+IV	262 (22.0%)	176 (22.0%)	61 (22.9%)	25 (19.8%)	
TNM_T					0.232
T1	481 (40.4%)	320 (40.0%)	102 (38.3%)	59 (46.8%)	
T2	639 (53.6%)	434 (54.3%)	141 (53.0%)	64 (50.8%)	
T3	31 (2.6%)	19 (2.4%)	10 (3.8%)	2 (1.6%)	
Tx	41 (3.4%)	27 (3.4%)	13 (4.9%)	1 (0.8%)	
TNM_N					0.113
N0	1161 (97.4%)	775 (96.9%)	260 (97.7%)	126 (100.0%)	
N1	31 (2.6%)	25 (3.1%)	6 (2.3%)	0 (0%)	
TNM_M					0.909
M0	957 (80.3%)	641 (80.1%)	213 (80.1%)	103 (81.7%)	
M1	235 (19.7%)	159 (19.9%)	53 (19.9%)	23 (18.3%)	
Tumor size					0.963
≤9.6	659 (55.3%)	437 (54.6%)	149 (56.0%)	71 (56.3%)	
9.6–20.5	430 (36.1%)	294 (36.8%)	93 (35.0%)	43 (34.1%)	
≥20.5	105 (8.6%)	69 (8.6%)	24 (9.0%)	12 (9.6%)	

Notes: ^adata from the SEER database; ^bdata from Cancer Hospital of China Medical University

tumor site, and tumor size were confirmed as independent prognostic factors for OS (Table 2).

Similar statistical procedures were conducted to identify the independent prognostic factors for CSS. In univariate analyses, variables of age, gender, tumor site, surgical stage, surgery, grade, histology, AJCC stage, node involvement, metastasis, tumor size were also significantly associated with CSS (all $p < 0.05$). Among them, the variables of age, surgical stage, surgery, grade, tumor site, and tumor size were further identified as independent prognostic factors for CSS (Table 3).

Establishment and validation of nomograms for OS and CSS. According to the results from multivariate Cox proportional hazards analyses, age, surgical stage, surgery, grade, tumor site, and tumor size were finally incorporated to develop the prognostic nomograms for predicting the 3- and 5-year OS and CSS in osteosarcoma patients (Figure 3). The detailed points of each selected variable were calculated and presented in Table 4.

The internal and external validations of the developed predictive prognostic nomograms were conducted based on the data from the SEER database and our hospital. Internal validation in the development cohort revealed that the C-index values for the nomograms of OS and CSS were 0.818 (95% CI 0.804–0.932) and 0.829 (95% CI 0.814–0.844), respectively. Comparatively, the calculated C-index values for OS and CSS in the external validation cohort one was 0.843 (95% CI 0.822–0.864) and 0.834 (95% CI 0.807–0.861), respectively. While in external validation cohort two, the corresponding C-index values for OS and CSS were 0.736 (95% CI 0.685–0.787) and 0.782 (95% CI 0.719–0.845), respectively. The C-index values from the three validation cohorts were all over 0.7, indicating the favorable accuracy of our prognostic nomograms. Furthermore, the calibration plots also showed excellent agreement between the nomogram estimated survival and actual survival in 3- and 5-year OS and CSS (Figures 4A–4L).

Table 2. Univariate and multivariate analyses of overall survival in the Development cohort.

Variables	Univariate analysis	Multivariate analysis	p-value
	p-value	HR (95%CI)	
Age	<0.001		
≤26		Reference	
26–61		2.046 (1.423–2.940)	<0.001
≥61		4.620 (3.313–6.440)	<0.001
Gender	0.001		
Male		Reference	
Female		0.814 (0.621–1.069)	0.139
Race	0.800		
Black		Reference	
White		1.309 (0.898–1.907)	0.161
Other		1.003 (0.572–1.755)	0.992
Tumor site	<0.001		
Axial		Reference	
Extremity		0.582 (0.567–1.077)	<0.001
Surgical stage	<0.001		
Regional		Reference	
Localized		0.871 (0.589–1.287)	0.487
Distant		2.442 (1.263–4.721)	0.007
Surgery	<0.001		
NO		Reference	
YES		0.404 (0.288–0.565)	<0.001
Grade	<0.001		
Well differentiated		Reference	
Moderately differentiated		1.977 (0.482–8.108)	0.343
Poorly differentiated		5.909 (1.754–19.897)	0.004
Undifferentiated/anaplastic		5.035 (1.512–16.768)	0.008
Histology	<0.001		
Conventional osteosarcoma		Reference	
Chondroblastic osteosarcoma		0.847 (0.575–1.247)	0.400
Fibroblastic osteosarcoma		0.909 (0.437–1.890)	0.799
Telangiectatic osteosarcoma		0.895 (0.280–2.881)	0.857
Osteosarcoma in Paget disease of bone		1.266 (0.285–5.609)	0.756
Small cell osteosarcoma		0.654 (0.199–2.148)	0.484
Central osteosarcoma		0.551 (0.262–1.156)	0.114
Intraosseous well differentiated osteosarcoma		0.869 (0.000–4.56E+78)	0.997
Parosteal osteosarcoma		0.854 (0.237–3.070)	0.808
Periosteal osteosarcoma		0.329 (0.000–2.31E+56)	0.991
High grade surface osteosarcoma		1.202 (0.165–8.727)	0.855
AJCC stage	<0.001		
I+II		Reference	
III+IV		3.652 (0.651–8.192)	<0.001
TNM_T	0.004		
T1		Reference	
T2		0.828 (0.498–1.375)	0.466
T3		0.997 (0.427–2.325)	0.994
Tx		0.954 (0.472–1.930)	0.897
TNM_N	0.04		
N0		Reference	
N1		0.916 (0.483–1.736)	0.789
TNM_M	<0.001		
M0		Reference	
M1		1.024 (0.417–2.506)	0.959
Tumor size	<0.001		
≤9.6		Reference	
9.6–20.5		1.658 (0.996–2.756)	0.051
≥21.4		2.143 (1.287–3.566)	0.003

Abbreviations: CI-confidence interval; HR-hazard ratio

Table 3. Univariate and multivariate analyses of cancer-specific survival in the Development cohort.

Variables	Univariate analysis	Multivariate analysis	p-value
	p-value	HR (95%CI)	
Age	<0.001		
≤26		Reference	
26–61		2.039 (1.361–3.055)	<0.001
≥61		3.144 (2.097–4.714)	<0.001
Gender	<0.001		
Male		Reference	
Female		0.743 (0.539–1.024)	0.070
Race	0.600		
Black		Reference	
White		1.506 (0.966–2.346)	0.070
Other		1.099 (0.570–2.120)	0.777
Tumor site	<0.001		
Axial		Reference	
Extremity		0.514 (0.615–1.358)	<0.001
Surgical stage	<0.001		
Regional		Reference	
Localized		1.041 (0.648–1.672)	0.867
Distant		2.720 (1.179–6.269)	0.018
Surgery	<0.001		
NO		Reference	
YES		0.337 (0.226–0.501)	<0.001
Grade	<0.001		
Well differentiated		Reference	
Moderately differentiated		2.207 (0.589–12.235)	<0.001
Poorly differentiated		5.307 (2.573–9.881)	<0.001
Undifferentiated/anaplastic		4.435 (2.014–7.495)	<0.001
Histology	<0.001		
Conventional osteosarcoma		Reference	
Chondroblastic osteosarcoma		0.739 (0.466–1.170)	0.197
Fibroblastic osteosarcoma		0.693 (0.277–1.733)	0.433
Telangiectatic osteosarcoma		1.117 (0.341–3.651)	0.854
Osteosarcoma in Paget disease of bone		1.313 (0.167–10.324)	0.795
Small cell osteosarcoma		0.827 (0.246–2.774)	0.757
Central osteosarcoma		0.525 (0.224–1.229)	0.137
Intraosseous well differentiated osteosarcoma		0.197 (0.000–3.895E+78)	0.909
Parosteal osteosarcoma		0.773 (0.160–3.737)	0.749
Periosteal osteosarcoma		0.199 (0.027–1.658)	0.142
High grade surface osteosarcoma		1.016 (0.032–6.326)	0.938
AJCC stage	<0.001		
I+II		Reference	
III+IV		3.716 (0.578–7.090)	<0.001
TNM_T	0.001		
T1		Reference	
T2		1.184 (0.665–2.105)	0.566
T3		1.399 (0.570–3.429)	0.463
Tx		1.033 (0.462–2.309)	0.936
TNM_N	0.09		
N0		Reference	
N1		0.876 (0.414–1.852)	0.730
TNM_M	<0.001		
M0		Reference	
M1		1.214 (0.458–3.211)	0.696
Tumor size	<0.001		
≤9.6		Reference	
9.6–20.5		1.826 (0.883–4.636)	0.003
≥21.4		3.942 (1.122–6.359)	0.017

Abbreviations: CI-confidence interval; HR-hazard ratio

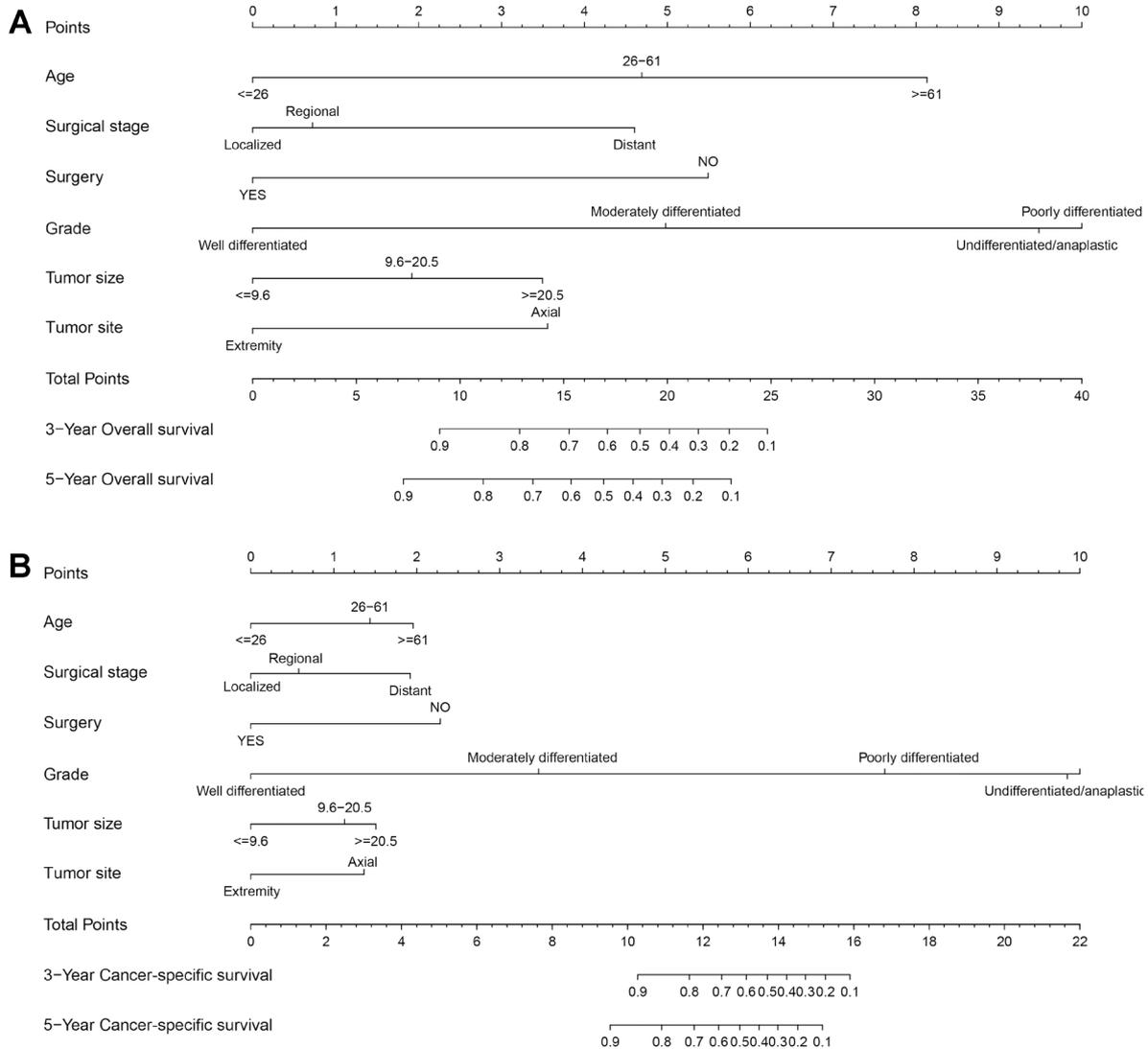


Figure 3. Nomograms to predict the overall survival (A) and cancer-specific survival (B) for patients with osteosarcoma. Firstly, by drawing a vertical line between each variable and the point scale, the points of this variable can be identified. Then, by projecting a vertical line from the total points scale, the sum of points of each variable included in the nomogram, to the OS and CSS scale, the individualized survival probability for an osteosarcoma patient can be obtained.

Discussion

Previous studies have suggested that multiple factors have prognostic or predictive values for the survival of osteosarcoma patients [41–43]. They can be roughly classified into seven groups based on their characteristics, as was mentioned in the introduction. Although these factors may form statistically significant associations with the survival of patients diagnosed with osteosarcoma, a single prognostic or predictive index may be inaccurate and unreliable in estimating an individual osteosarcoma patient’s medium- or long-term survival. In addition, studies that incorporate these factors into a comprehensive and representative index

or scoring system are scarce and thus impede its further application in individualized survival prediction. Fortunately, a nomogram is such a common statistical tool that can effectively integrate these prognostic factors together and provide satisfactory reliability, accuracy, and robustness in predicting an individual osteosarcoma patient’s survival probability [44]. Kim et al. constructed and internally validated a prognostic nomogram that could predict the 5-year probability of metastasis for AJCC stage II extremity osteosarcoma patients (n=141) and this nomogram showed obvious superiority in metastasis risk estimation compared with the AJCC staging system or tumor necrosis rate [45]. In 2014, the author published another nomogram to evaluate

Table 4. Points of each variable included in the nomograms.

Variables	Overall survival nomogram	Cancer-specific survival nomogram
Age (years)		
≤26	0	0
26–61	4.5	1.5
≥61	8.0	2.0
Surgical stage		
Regional	1.0	0.5
Localized	0	0
Distant	4.5	2.0
Surgery		
NO	5.5	2.0
YES	0	0
Grade		
Well differentiated	0	0
Moderately differentiated	5.0	3.5
Poorly differentiated	9.5	7.5
Undifferentiated/anaplastic	10.0	10.0
Tumor size (cm)		
≤9.6	0	0
9.6–20.5	2.0	1.0
≥21.4	3.5	1.5
Tumor site		
Extremity	0	0
Axial	3.5	1.5

the risk of metastasis in patients diagnosed with Enneking stage IIB extremity osteosarcoma (n=91) who received limb salvage surgery [46]. However, the two studies just focused on AJCC stage II or Enneking stage IIB extremity osteosarcoma patients, so the representativeness of the patients was limited. Moreover, several other limitations also existed in these studies, including without external validation based on other populations and relatively small sample sizes. Therefore, their results might not apply to other populations due to potential bias. Xia et al. also developed a nomogram that included an inflammatory response marker of neutrophil-to-lymphocyte ratio (NLR) to predict the survival of osteosarcoma patients who underwent curative surgery [9]. However, this study devised a nomogram with only three prognostic factors included, namely NLR, tumor stage, and initial metastases, and thus crippled its clinical and practical values for its inability to represent the overall condition of an osteosarcoma patient. Therefore, the above-mentioned nomograms should be validated in larger and different populations before its application in clinical practice. So, in our study, by integrating pivotal prognostic and determinant variables together, we first devel-

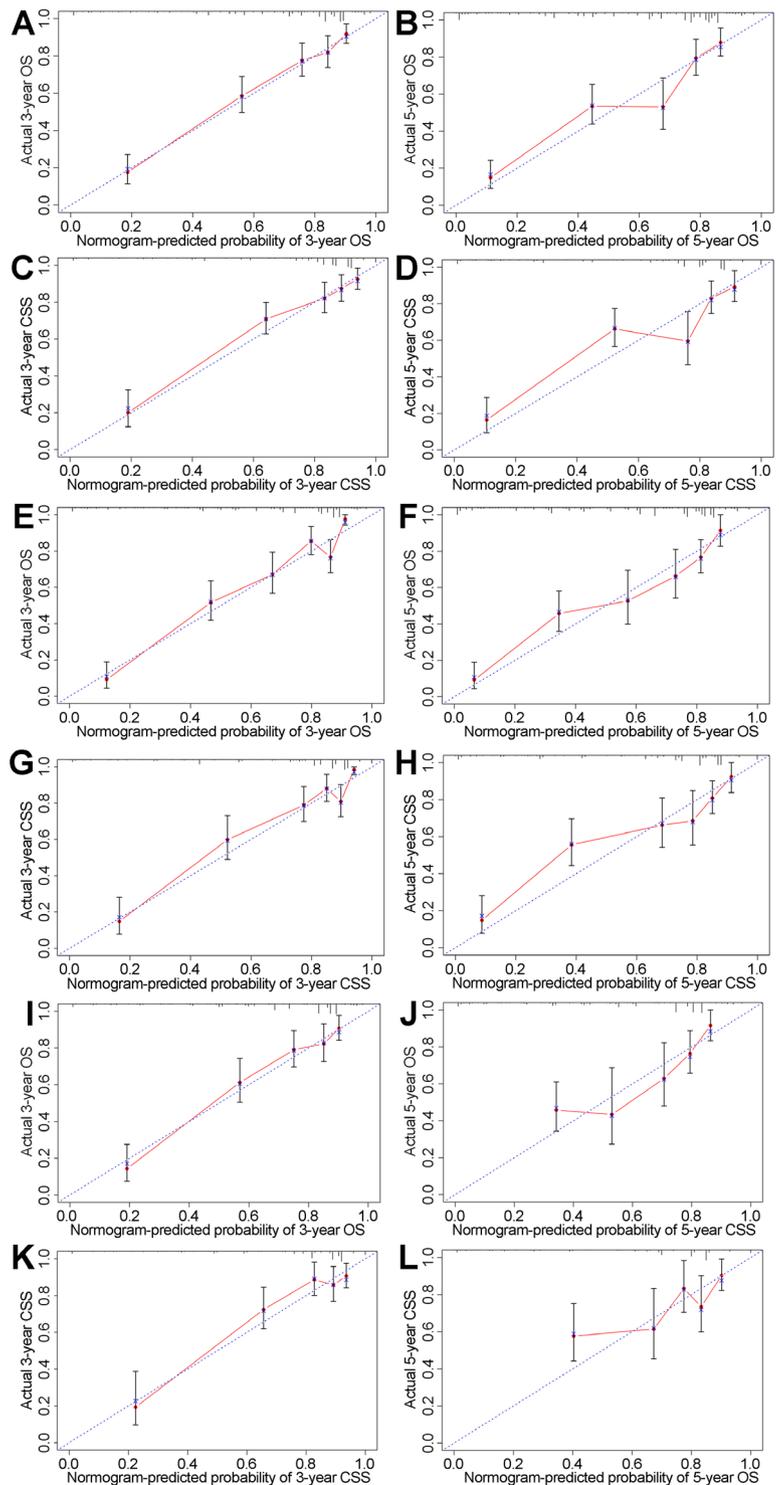


Figure 4. The twelve graphs show the calibration plots of internal validation and external validation for the 3- and 5-year overall survival (OS) and cancer-specific survival (CSS). A–D) The first four calibration curves present the internal validation for 3- and 5-year OS and CSS based on the data from the SEER database. E–H) The middle four calibration curves show the external validation for 3- and 5-year OS and CSS based on the data from the SEER database. I–L) The last four calibration curves also reveal the external validation, data come from our hospital, for 3- and 5-year OS and CSS.

oped and internally validated the prognostic nomograms to evaluate the 3- and 5-year OS and CSS rates for patients with osteosarcoma based on the data of a cohort from the SEER database. Then, we externally validated the established nomograms with the data of another cohort from the SEER database. Finally, we performed further external validation of the nomograms with the data from our hospital.

Based on results from univariate and multivariate Cox analyses, several clinicopathological factors were finally identified to be independently associated with the survival of patients with osteosarcoma, including patient age, surgical stage, surgery, tumor grade, tumor size, tumor site, and AJCC stage. In the present study, increasing age was significantly associated with worse survival status in osteosarcoma patients. Similar trends have been reported in previous studies, but the cut-off values for patient age may vary in different studies. Zheng et al. reported that patients with osteosarcoma older than 51 had worse survival time and Song et al. proposed that patients' age over 40 years old was an independent risk factor for poor survival [39, 40]. Similarly, another study based on the SEER database analyses and performed by Fu et al. also suggested that patients older than 25 years almost suffered 2–3 times increased risks to get poorer survival [47]. In our study, we identified the age of 26 and 61 as the best cut-off points for risk stratification by using the X-tile software and significant differences in OS and CSS were found between patients ≤ 26 years, 26–61 years, and ≥ 61 years. The distinction in cut-off values of age at disease onset can be explained to some extent that it was determined by the sample size and study populations in different studies. The surgical stage, or the extent of disease, was also an independent risk factor for worse survival outcomes. Patients who suffered metastatic lesions or accompanied by regional lymph nodes involvement usually had a poorer prognosis than those with a localized disease or without lymph nodes metastases [6, 39, 40, 47, 48]. Consistent with previous findings, our study also revealed that osteosarcoma patients with regional or distant disease had higher mortality than patients with localized disease.

Currently, neoadjuvant chemotherapy followed by surgical resection remains the standard modality for patients with non-metastatic osteosarcoma [4]. Our study also validated and confirmed the beneficial effects of surgery on the survival of patients with osteosarcoma who received treatment of chemotherapy based on the results of multivariate analyses. In our study, we also identified tumor grade as independent prognostic factors for the survival outcomes of osteosarcoma patients, which was in line with the results in previous studies. Patients with a poorly differentiated or undifferentiated histology grade were more prone to get worse OS and CSS than patients with well-differentiated or moderately differentiated histology grade. It was demonstrated by Wang et al. that high-grade osteosarcoma had significantly worse survival outcomes compared with low-grade osteosarcoma [49]. With respect to histology subtypes, the report from Jawad et al. demonstrated that Paget's osteosarcoma was

significantly related to poorer prognosis compared with other histological subtypes and this result was further confirmed by the report of Damron et al. [50, 51]. Our study also found that the periosteal osteosarcoma had a significantly better prognosis than all other histological subtypes of osteosarcoma, which was similar to the results reported by Zheng et al. [39].

In previous studies, axial tumor locations and large tumor sizes were deemed as the worst prognostic factors for the survival outcomes of patients with osteosarcoma [52–55]. Our study suggested that patients with axial tumors or larger tumors had decreased survival time compared with patients with extremity tumors or smaller tumors. Larger tumor sizes and axial tumor locations all impose several disadvantages for osteosarcoma patients, including increased risk for metastasis, soared difficulties for surgical resection, more difficulties to achieve adequate surgical margins, the rich blood supply to accelerate the process of elimination and metabolism of chemotherapeutic agents, and so on. Fujiwara et al. studied the association between proximity to the major vessels and the risk of local recurrence and survival in 226 patients with high-grade non-metastatic osteosarcoma [54]. Based on the examination of preoperative Magnetic Resonance Imaging (MRI), the vascular proximity was categorized into four types: type 1 with proximity >5 mm; type 2 with proximity ≤ 5 mm, >0 mm; type 3 with tumors attached to major vessels; type 4 with tumors surrounded major vessels. The results revealed that the limb salvage rate and the 5-year OS rate were larger in patients with longer tumor-to-vessel proximities than those with shorter proximities (92%, 88%, 51%, 0% vs. 82%, 77%, 57%, 67%). Moreover, the local recurrence rate was also higher in patients with shorter tumor-to-vessel proximities (types 1 to 3; 7%, 8%, and 22%). Furthermore, Bielack et al. and Duchman et al. also demonstrated that patients with larger tumors adjacent to trunk bones are more likely to develop metastases during chemotherapy [52, 56]. Besides, our study also found that the AJCC stage was also independently related to the prognosis of osteosarcoma patients. Patients with a higher AJCC stage were usually accompanied by a poorer prognosis and survival. In the present study, the AJCC stage was not included in the nomograms for it was a comprehensive evaluating system based on tumor size, lymph node involvement, and metastasis. If it was included for the construction of the nomogram, the weight coefficient for tumor size and surgical stage may alter for the reasons for repeated calculation of weight coefficient for them. The established nomograms for osteosarcoma patients in previous studies also did not incorporate the AJCC stage into nomograms [39, 40].

Different from previous studies, we performed external validation of the established nomograms with the data of a different cohort from our hospital. Previously, we have conducted a study to explore the predictive values of pre-treatment Naples prognostic score (NPS) in patients with osteosarcoma in a cohort of 133 patients [57]. We constructed two prediction models based on the multivariate Cox

analyses, namely the conventional prediction model (model A) based on prognostic factors of Enneking stage, metastasis, local recurrence, and NLR, and the combined prediction model (model B) based on prognosticators of Enneking stage, metastasis, local recurrence, NLR, and NPS. The results from our previous study demonstrated that NPS based nomograms got better time-dependent AUCs than conventional nomograms only established by clinical characteristics. The AUCs for model A were 0.802–0.878 and 0.773–0.840 for OS and PFS, while the AUCs for model B were 0.664–0.713 and 0.773–0.718 for OS and PFS, respectively. In this study, the C-indexes of the nomograms in the development set (0.818 for OS, 0.829 for CSS) and the two validation sets (validation set one, 0.843 for OS and 0.834 for CSS; validation set two, 0.736 for OS and 0.782 for CSS) were all over 0.70. By contrast, the predictive performances of the nomograms in this study and our previous study were both reliable and effective. However, the nomograms based on a large cohort from the SEER database seemed to be slightly better than the nomograms based on the small cohort from our hospital. This is within our anticipation because prediction models based on a larger cohort are more prone to harvest better prediction performances than those established from a relatively smaller cohort. Anyhow, the indexes (AUC or C-index) representing the predictive accuracy in our current study and previous study were both larger than 0.7, which indicates these nomograms were all reliable and effective.

Although the established nomograms in our study showed good performance in predicting the OS and CSS for osteosarcoma patients, several limitations should also be taken into consideration. First, the detailed data on chemotherapy and radiotherapy were incomplete and limited in the SEER database, leading to the loss of some important variables for nomogram construction and possibilities to cause some relevant bias. Second, some pivotal serological biomarkers were not included in our nomogram, such as lactate dehydrogenase (LDH) and alkaline phosphatase (ALP), for such variables were not available and cannot be extracted from the SEER database. Finally, other factors, like local recurrence and multiple metastases, were also important endpoints for osteosarcoma patients, but they were also unextractable in the SEER database, which impeded the process for further analysis. Despite these limitations, the nomograms also showed good predictive performance in external validation with the data from our hospital (C-index = 0.736).

In summary, we firstly established prognostic nomograms to predict the 3- and 5-year OS and CSS rates for patients with osteosarcoma based on the data from the SEER database. Then, the constructed nomograms were internally and externally validated with data from the SEER database and our hospital, which all showed good predictive performance for osteosarcoma patients. With this predictive tool, the 3- and 5-year survival of an individual osteosarcoma patient can be estimated, enabling oncologists to evaluate individualized survival probability and identify mortality risk stratification.

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