Establishing and Assessing a Nomogram Based on the Clinical Features of Mucosal Melanoma of the Head and Neck

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Abstract

Objectives: The rarity of mucosal melanoma of the head and neck (MMHN) and the lack of prospective clinical trials has resulted in the limited knowledge of its clinical features and prognosis. We aimed to understand the clinical characteristics and developed a nomogram to better predict the prognosis of patients with MMHN.

Methods: Based on a total of 300 patients with nonmetastatic MMHN, multivariable Cox regression was performed to analyze independent prognostic factors. The overall survival (OS), disease-free survival (DFS), distant metastasis-free survival (DMFS), and locoregional relapse-free survival (LRRFS) of MMHN patients could be monitored using the nomogram. To facilitate clinical application, an online dynamic nomogram was established.

Results: Multivariate analysis identified primary tumor site, T stage and N stage as independent risk factors for survival. This factor-based nomogram had prognostic value for OS, DFS, DMFS, and LRRFS. Receiver operating characteristic analysis indicated the high diagnostic accuracy of the nomogram (AUC > 0.7). Kaplan–Meier survival curves indicated that the risk score of the nomogram effectively stratified MMHN patients with poor survival into a high-risk group (all $P < 0.001$).

Conclusions: The nomogram is conducive to stratifying MMHN patients into clinically meaningful taxonomies and subsequently providing individualized treatment.

Introduction

Mucosal melanoma of the head and neck (MMHN) is a highly malignant tumor accounting for more than 50% of all mucosal melanomas¹ and is mainly found in the nasal cavity, paranasal sinuses and oral cavity². Compared with cutaneous and acral melanomas, mucosal melanoma has aggressive clinical characteristics and a worse prognosis^{3–5}. Despite the application of various clinical treatments, including surgery, radiotherapy, chemotherapy and immunotherapy, the local control and long-term prognosis of MMHN remain quite dismal and seem to show no improvement trends in recent years^{6,7}. Due to its rarity, there is no effective prognostic tool available for MMHN. Therefore, more accurate prognostic tools are needed in addition to improving the treatment.

Considering the rarity of MMHN and the paucity of optimal treatment modalities following diagnostic tools, in this study, we included the largest single-center sample size, aiming to further explore MMHN by analyzing the clinical features and treatment outcomes. A nomogram serves as a new reliable tool for predicting the prognosis of cancers^{8–10}. Therefore, we combined the TNM staging system and clinical features to develop a nomogram to accurately predict the survival outcomes of MMHN patients to facilitate clinical decision making.

Materials And Methods

Patients

From March 1986 to November 2019, three hundred patients diagnosed with MMHN in our hospital were included in the primary cohort. MMHN patients from March 1986 to December 2014 were chosen as the validation cohort. The inclusion criteria for all patients were previously untreated, nonmetastatic, and newly histologically confirmed stage III-IVB MMHN. All patients were restaged according to the 8th edition American Joint Committee on Cancer (AJCC) staging system for MMHN¹¹. The TNM classification was based on surgical documents, pathological features and imaging findings. The exclusion criteria were as follows: (1) distant metastases before the treatment, secondary malignancy, or both; (2) pregnancy or lactation; and (3) incomplete previous medical records, auxiliary examinations, and follow-up information. The Ethics Committee at Sun Yat-sen University Cancer Center in China approved our study protocol.

Follow-up and Endpoints

Our main endpoint was overall survival (OS), and the secondary endpoints were disease-free survival (DFS), distant metastasis-free survival (DMFS), and locoregional relapse-free survival (LRRFS). OS was defined as the time from the diagnosis of melanoma to the date of death or the last known follow-up, whichever occurred first; DMFS was defined as the time to distant metastasis, death, or patient censoring, whichever occurred first; DFS was defined as the time to failure, death from any cause, or patient censoring, whichever occurred first; and LRRFS was defined as the time to local/regional relapse, death, or patient censoring, whichever occurred first. The median follow-up time was 32 months (range: 1-262 months). After treatment, the patients were evaluated once every 3 months during the first 3 years and every 6 months thereafter.

Statistical Analysis

The patients were classified into 2 groups based on age (< 60 years vs. ≥ 60 years). Variables satisfying $P < 0.1$ in univariate Cox regression analysis were included in the multivariable analysis. $P < 0.05$ in multivariable Cox regression analysis was used to select independent prognostic variables of survival. All independent prognostic factors were used to create a predictive nomogram (by the rms package in R). The concordance index (c-index) values with 95% confidence intervals (CIs) were evaluated to assess the accuracy of the nomogram in the primary and validation cohorts. Calibration plots for OS, DMFS, and DFS at three and five years were generated by comparing the predicted OS, DMFS, and DFS with the actual OS, DMFS, and DFS. Moreover, the predictive precision and discrimination of the nomogram were further analyzed by the Akaike information criterion (AIC), c-index, and area under the curve (AUC) of the receiver operating characteristic (ROC) curve. Survival curves were generated using the Kaplan–Meier method. Differences in survival between risk groups were analyzed by the log-rank test. The statistical analysis was performed with R software (R version 3.6.1) (<http://www.r-project.org>) and IBM SPSS software version 25.0.

Results

Patient Characteristics and Follow-up

From March 1986 to November 2019, 300 patients (primary cohort) and 182 patients (validation cohort) with MMHN were enrolled in this study. The median age was 57 (range, 19–87) years with a male-to-female ratio of 1.54:1 for the primary cohort. The median age was 58 (range, 25–80) years with a male-to-female ratio of 1.54:1 for the validation cohort. The baseline patient characteristics of the primary cohort and validation cohort are listed in Table 1.

Table 1

Comparison of the different characteristics between patients in the primary and validation cohorts.

Characteristic	Number of patients (%)		<i>P</i> -value
	Primary patients (n = 300)	Validation cohort (n = 182)	
Gender			0.944
Male	182 (60.7)	111 (61.0)	
Female	118 (39.3)	77 (39.0)	
Age (years old)			0.373
< 60	174 (58.0)	98 (53.8)	
≥ 60	126 (42.0)	84 (46.2)	
Smoking			0.064
No	220 (73.3)	119 (65.4)	
Yes	80 (26.7)	63 (34.6)	
Primary site			0.753
Others	44 (14.7)	23 (12.6)	
Nasal cavity	159 (53.0)	100 (54.9)	
Paranasal sinus	24 (8.0)	11 (6.0)	
Oral cavity	73 (24.3)	48 (26.4)	
8th T stage			0.966
T3	121 (40.3)	73 (40.1)	
T4a	129 (43.0)	77 (42.3)	
T4b	50 (16.7)	32 (17.5)	
8th N stage			0.604
N0	224 (74.7)	132 (72.5)	
N1	76 (25.3)	50 (27.5)	
Surgery			0.933

Abbreviations:

All continuous variables were changed to categorical variables. Pearson χ^2 test was used to compute the *P*-value.

No	21 (7.0)	13 (7.1)	
Yes	170 (56.7)	100 (54.9)	
Reoperation	109 (36.3)	69 (37.9)	
Radiotherapy			< 0.001
No	179 (59.7)	129 (70.9)	
2D	22 (7.3)	22 (12.1)	
IMRT	99 (33.0)	31 (17.0)	
Chemotherapy			0.377
No	131 (43.7)	87 (47.8)	
Yes	169 (56.3)	95 (52.2)	
Immunotherapy			0.659
No	217 (72.3)	135 (74.2)	
Yes	83 (27.7)	47 (25.8)	
Abbreviations:			
All continuous variables were changed to categorical variables. Pearson χ^2 test was used to compute the <i>P</i> -value.			

The median follow-up time was 32.0 months (range, 1-262 months). A total of 172 patients experienced disease failure; 75 (7.7%), 52 (3.8%), and 109 (14.6%) patients developed local recurrence, regional recurrence, and distant metastasis, respectively; and 188 (14.4%) patients died.

Univariate and Multivariate Cox Regression Analyses

The significant variables related to OS were primary tumor location, T stage, N stage, and immunotherapy in univariate analysis. The significant variables related to DFS, DMFS, and LRRFS were primary tumor location, T stage, and N stage in univariate analysis. We incorporated the above factors into multivariate Cox regression analysis. Eventually, T stage, N stage, and primary tumor site were independent prognostic factors. The results of univariate and multivariate Cox analyses are summarized in Table 2, 3 and Fig. 1.

Table 2
Univariate Cox regression analysis of OS and DFS in the primary cohort.

Variable	OS		DFS	
	HR (95% CI)	P	HR (95% CI)	P
Gender				
Male	Reference		Reference	
Female	0.964 (0.717 to 1.297)	0.811	1.051 (0.766 to 1.442)	0.759
Age (years old)				
< 60	Reference		Reference	
≥ 60	1.246 (0.934 to 1.660)	0.134	1.203 (0.881 to 1.644)	0.245
Smoking				
No	Reference		Reference	
Yes	1.020 (0.747 to 1.395)	0.899	0.787 (0.591 to 1.049)	0.102
Primary site				
Others	Reference		Reference	
Nasal cavity	2.127 (1.256 to 3.603)	0.005	1.510 (0.996 to 2.288)	0.052
Paranasal sinus	6.113 (3.101 to 12.052)	< 0.001	3.066 (1.733 to 5.425)	< 0.001
Oral cavity	2.586 (1.465 to 4.564)	0.001	2.217 (1.413 to 3.479)	0.001
8th T stage				
T3	Reference		Reference	
T4a	2.801 (1.968 to 3.989)	< 0.001	2.150 (1.596 to 2.896)	< 0.001
T4b	11.025 (7.301 to 16.647)	< 0.001	6.179 (4.227 to 9.034)	< 0.001
8th N stage				
N0	Reference		Reference	
N1	1.542 (1.113 to 2.237)	0.009	1.637 (1.223 to 2.191)	0.001

Variable	OS		DFS	
	HR (95% CI)	P	HR (95% CI)	P
Surgery				
No	Reference		Reference	
Yes	0.707 (0.409 to 1.220)	0.212	0.612 (0.386 to 0.969)	0.036
Radiotherapy				
No	Reference		Reference	
Yes	1.178 (0.879 to 1.580)	0.273	1.070 (0.828 to 1.383)	0.603
Chemotherapy				
No	Reference		Reference	
Yes	0.863 (0.647 to 1.151)	0.317	1.022 (0.792 to 1.318)	0.869
Immunologic/targeted therapy				
No	Reference		Reference	
Yes	0.694 (0.495 to 0.974)	0.035	0.873 (0.657 to 1.159)	0.348

Table 3
Univariate Cox regression analysis of DMFS and LRRFS in the primary cohort.

Variable	DMFS		LRRFS	
	HR (95% CI)	P	HR (95% CI)	P
Gender				
Male	Reference		Reference	
Female	0.964 (0.717 to 1.297)	0.811	1.036 (0.793 to 1.355)	0.794
Age (years old)				
< 60	Reference		Reference	
≥ 60	1.246 (0.641 to 1.164)	0.134	1.070 (0.822 to 1.391)	0.616
Smoking				
No	Reference		Reference	
Yes	0.864 (0.625 to 1.236)	0.336	0.819 (0.609 to 1.100)	0.185
Primary site				
Others	Reference		Reference	
Nasal cavity	1.695 (1.084 to 2.649)	0.021	1.795 (1.128 to 2.858)	0.014
Paranasal sinus	3.736 (2.032 to 6.867)	< 0.001	3.710 (1.995 to 6.897)	< 0.001
Oral cavity	1.9844 (1.217 to 3.235)	0.006	2.631 (1.601 to 4.323)	< 0.001
8th T stage				
T3	Reference		Reference	
T4a	2.544(1.838,3.520)	< 0.001	1.945 (1.434 to 2.639)	< 0.001
T4b	10.314(6.861,15.502)	< 0.001	5.816 (4.002 to 8.452)	< 0.001
8th N stage				
N0	Reference		Reference	
N1	1.597 (1.180 to 2.161)	0.002	1.665 (1.231 to 2.253)	0.001

Variable	DMFS		LRRFS	
	HR (95% CI)	P	HR (95% CI)	P
Surgery				
No	Reference		Reference	
Yes	0.758 (0.455 to 1.262)	0.286	0.664 (0.409 to 1.078)	0.098
Radiotherapy				
No	Reference		Reference	
Yes	1.359 (1.037 to 1.224)	0.026	1.007 (0.771 to 1.316)	0.957
Chemotherapy				
No	Reference		Reference	
Yes	1.022 (0.792 to 1.318)	0.869	0.934 (0.718 to 1.215)	0.611
Immunologic/targeted therapy				
No	Reference		Reference	
Yes	0.858 (0.634 to 1.161)	0.320	0.801 (0.594 to 1.079)	0.144

Establishing and Validating a Nomogram

All factors, including primary tumor site, T stage, and N stage, were included in the nomogram. By summarizing the score of each variable and positioning the total scores on the score scale, a nomogram was established to predict the 3- and 5-year OS, DFS, DMFS, and LRRFS in the primary cohort (Figs. 2A–D).

The c-indexes of the nomogram for predicting OS, DFS, DMFS, and LRRFS were almost over 0.7 in all cohorts, which indicated that the performance of the model was satisfactory.

In the calibration plot, the X-axis is the nomogram prediction of OS, DFS, DMFS, and LRRFS, and the Y-axis is the observed values of OS, DFS, DMFS, and LRRFS calculated by the Kaplan-Meier method. The solid line is the ideal reference line, which represents the consistency between the predicted survival rate and the observed survival rate. The calibration plots for the OS, DFS, DMFS, and LRRFS probabilities at 3 and 5 years showed significant correspondence between the predictions and observations in all cohorts (Figs. 3 and 4). Moreover, ROC curve analysis was also used to assess the predictive capacity of the nomogram, and the AUC for all cohorts indicated excellent validity (AUC > 0.70) (Fig. 5).

Nomogram for Risk Stratification

Consequently, stratification was established according to the nomogram for OS, DFS, DMFS, and LRRFS. Based on the total scores calculated by the nomogram, we set a threshold for the total score (33% and 66%). The primary and validation cohorts were divided into low-risk [total score: <50 (OS); <42 (DFS); <40 (DMFS); <45 (LRRFS)], intermediate-risk [total score: 50–85 (OS); 42–64 (DFS); 40–60 (DMFS); 45–69 (LRRFS)] and high-risk [total score: > 85 (OS); >64 (DFS); >60 (DMFS); >69 (LRRFS)] groups. According to the Kaplan–Meier survival curves, there were obvious differences among the different risk groups (all P-values < 0.001; Fig. 6).

Establishment of a web server for Easy Application of the nomogram

We have created a web server of our nomogram (Fig. 7), which can be easily accessed at https://liling.shinyapps.io/MMHN_OS/, https://liling.shinyapps.io/MMHN_DFS/, https://liling.shinyapps.io/MMHN_DMFS/, and https://liling.shinyapps.io/MMHN_LRRFS/. OS, DFS, DMFS, and LRRFS in NPC patients can be predicted conveniently by selecting corresponding clinical features and reading generated figures and tables.

Discussion

MMHN is an aggressive malignancy whose comprehensive treatment modalities are complex and clinical outcomes are still unsatisfactory¹². The AJCC 8th edition staging system lacks the ability to predict prognosis¹³. MMHN patients in the same stage can have obviously different survival rates¹⁴. Integrating other predictive factors into the TNM staging system will add prognostic value to this system and guide more promising treatment decisions.

Our results revealed that the primary tumor site was an independent prognostic factor. Consistent with the findings of several previous studies, our results showed that worse outcomes existed when the primary tumor site was in the paranasal cavity than in the oral or nasal cavities^{15–17}. There may be several plausible reasons for the poorer survival of patients with a primary tumor site in the paranasal sinuses. Due to the complex anatomical location of the paranasal sinus, paranasal sinus melanoma is found at later stages and has a wide range of invasion, so the prognosis of these patients may be poor. T stage and N stage were also independent prognostic factors, similar to the findings of other articles^{18–20}.

The study has the following advantages. First, the survival rate of a patient can be calculated simply and visually. For example, suppose a patient with T3 (0 points) N0 (0 points) paranasal sinus MM (50 points) comes to the clinic; he/she would have a total of 50 points, yielding an estimated 3-year OS rate of 28.57%. Second, the nomogram can further divide patients into high-, intermediate- and low-risk groups, and the survival rate of each group can effectively distinguish the prognosis. Doctors can give individualized treatment to patients according to their corresponding groups. The intensity of treatment

can be appropriately strengthened for patients in the high-risk group. For example, immunologic/targeted therapy can improve the survival rate of MMHN²¹. Finally, in addition to the traditional nomogram, we also created a dynamic nomogram that can predict the prognosis of patients through simple operation on a web page (Fig. 7).

Our study has several limitations. First, this study was a retrospective study with inevitable selection bias. However, this retrospective study was worth carrying out because it is of great significance to lay the foundation for further prospective studies.

Second, this study involved patients from the same hospital; thus, the findings may lack applicability for patients from other regions and institutions. Finally, the sample size was not large enough, so there were no significant differences in some prognostic factors, especially the treatment modalities. Nonetheless, our study included 300 Chinese MMHN patients, which currently constitutes the largest sample from a single center to our knowledge.

In conclusion, our nomogram combines primary site with T stage and N stage to predict the survival outcomes of MMHN patients, which can help doctors easily distinguish high- and low-risk patients clinically for individualized treatment.

Abbreviations

AJCC/UICC: American Joint Committee on Cancer/Union for International Cancer Control; CI: confidence interval; OS: overall survival; DMFS: distant metastasis-free survival; HR: hazard ratio; IMRT: intensity-modulated radiotherapy; LRRFS: locoregional relapse-free survival.

Declarations

Acknowledgements

Not applicable

Authors' contributions

Conceptualization: LXL, LC, QQX. Data analysis: QQX, QJL, LC, XYS. Original

draft writing: QQX, QJL, LC, JXS. Review and editing: LXL, LC, QQX, QJL, XYS,

JXS. All authors read and approved the final manuscript.

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Availability of data and materials

The databases analyzed during the current study are available.

Ethics approval and consent to participate

The Hospital Ethics Committee at Sun Yat-sen University Cancer Center in China

approved the study, which analyzed anonymous information as well as waived the demand for informed consent.

Consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

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Figures

Figure 1

Multivariate Cox regression analysis of OS (A), DFS (B), DMFS (C), and LRRFS (D) in the primary cohort.

Figure 2

Developing a nomogram. Nomogram was based on the NPC patients' data of primary site, 8th T stage, and 8th N stage for 3-, 5-year OS (A), DFS (B), DMFS (C), and LRRFS (D) in the primary cohort.

Figure 3

Calibration plots of the nomogram to predict 3- and 5-year OS and DFS in the primary (A, C, E, G) and validation cohorts (B, D, F, H). Nomogram-estimated 3- or 5-year OS (A-D) and DFS (E-H) were plotted on the x-axis; the observed OS and DFS were plotted on the y-axis. Dashed lines along the 45-degree line represented that the predicted probabilities are equal to the actual probabilities.

Figure 4

Calibration plots for 3- and 5-year DMFS and LRRFS in the primary (A, C, E, G) and validation cohorts (B, D, F, H). Nomogram-estimated 3- or 5-year DMFS (A-D) and LRRFS (E-H) were plotted on the x-axis; the observed DMFS and LRRFS were plotted on the y-axis. Dashed lines along the 45-degree line represented that the predicted probabilities are equal to the actual probabilities.

Figure 5

ROC curves by the nomogram to predict 3-, 5-year OS (A, B), DFS (C, D), DMFS (E, F), and LRRFS (G, H).

Figure 6

Kaplan-Meier curves for OS (A, B), DFS (C, D), DMFS (E, F), and LRRFS (G, H) of patients in low-, intermediate- and high-risk groups. The risk groups were stratified according to the 33% and 66% of total risk scores in the primary (A, C, E, G) and validation cohorts (B, D, F, H).

Dynamic Nomogram

Figure 7

A web server for easy application of the nomogram