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Development of prognostic models for HER2positive metastatic breast cancer in females: a retrospective population-based study



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Abstract

Background This study aimed to construct, evaluate, and validate nomograms for breast cancer-specific survival (BCSS) and overall survival (OS) prediction in patients with HER2- overexpressing (HER2+) metastatic breast cancer (MBC).

Methods The Surveillance, Epidemiology, and End Results (SEER) database was used to select female patients diagnosed with HER2 + MBC between 2010 and 2015. These patients were distributed into training and validation groups (7:3 ratio). Variables were screened using univariate and multivariate Cox regression analyses, and BCSS and OS nomograms were constructed to determine one-, three-, and five-year survival probabilities. The nomograms were evaluated and validated using the concordance index (C-index), time-dependent receiver operating characteristic (ROC) curves, calibration curves, and decision curve analysis. Stratification was evaluated using Kaplan–Meier curves and log-rank tests based on optimal total score cut-off values. We published web-based versions of these nomograms for clinical use.

Results A total of 2,151 eligible patients were randomized into training (n = 1,505) and validation (n = 646) groups. Independent prognostic factors of BCSS and OS included: age; marital status; race; oestrogen receptor status; surgery; chemotherapy; and bone, brain, liver, and lung metastases. The C-indices for the BCSS and OS training groups were 0.707 and 0.702, respectively. The ROC, calibration, and decision curves demonstrated the strength of the nomograms. According to cut-off values, patients were categorized into low-, intermediate-, and high-risk groups, with significant differences in survival outcomes between them.

Conclusion We constructed predictive nomograms and stratified risk to assess the prognosis of patients with HER2 + MBC, which could help inform therapeutic decisions.

Trial registration Not applicable.

Keywords HER2-positive breast cancer, Metastatic, Nomogram, Breast cancer-specific survival, Overall survival, SEER database

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Background

As one of the most frequent malignant tumours in women today, breast cancer has the highest incidence rate (31%) and the second highest mortality rate (15%) among all malignancies [1]. Breast cancer is classified into subtypes based on oestrogen receptor (ER), progesterone receptor (PR), and human epidermal growth factor receptor 2 (HER2) expression. Approximately 15–20% of primary breast cancers overexpress HER2; this cancer subtype (HER2+) is invasive and has a high risk of recurrence compared with subtypes that do not overexpress HER2, resulting in a poorer prognosis [2, 3].

Furthermore, distant metastases occur in approximately 29.4% of patients with HER2+breast cancer, with visceral metastasis being the most common [4]. HER2-targeted drugs are in development, including monoclonal antibodies, tyrosine kinase inhibitors, and antibody-coupled drugs, which have significantly improved patient prognosis [5]. However, HER2+metastatic breast cancer (MBC) currently has no cure, and resistance to HER2 therapy substantially decreases survival time [6]. Therefore, a prognostic model for patients with HER2+MBC is crucial.



Table 1 Baseline characteristics of the selected patients diagnosed with HER2-positive metastatic breast cancer

Characteristics	Total (<i>n</i> =2151)	Training group (<i>n</i> = 1505)	Validation group (n=646)	Р
Age, n(%)		,		0.954
20–39	272 (12.65)	191 (12.69)	81 (12.54)	
40–59	1011 (47.00)	710 (47.18)	301 (46.59)	
60–79	732 (34.03)	507 (33.69)	225 (34.83)	
80+	136 (6.32)	97 (6.45)	39 (6.04)	
Marital status, n(%)				0.169
Married	997 (46.35)	682 (45.32)	315 (48.76)	
Single	474 (22 04)	329 (21 86)	145 (22 45)	
Widowed/divorced/other	680 (31 61)	494 (32 82)	186 (28 79)	
Bace n(%)	000 (01101)	13 (02.02)	100 (2011 5)	0.676
Black	356 (16 55)	253 (16.81)	103 (15 94)	0.070
White	1578 (73 36)	1096 (72.82)	482 (74 61)	
Other	217 (10.09)	156 (10.37)	61 (9.44)	
Histologic type, p(%)	217 (10.05)	150 (10.57)	01 (9.44)	0.675
	1843 (85.68)	1285 (85 28)	558 (86 38)	0.075
	61 (2.9.4)	45 (200)	16 (2.49)	
Mixed	01 (2.04)	43 (2.99)	20 (2.46)	
Other	00 (S.72) 1 (7 (7 7 ()	00 (5.99)	20 (3.10)	
	107 (7.70)	115 (7.64)	52 (8.05)	1 000
Grade, n(%)	42 (2.00)	20 (1 00)	12 (2.01)	1.000
1	43 (2.00)	30 (1.99)	13 (2.01)	
II	/28 (33.84)	510 (33.89)	218 (33.75)	
	13/4 (63.88)	961 (63.85)	413 (63.93)	
	6 (0.28)	4 (0.27)	2 (0.31)	
ER Status, n(%)				0.243
Positive	1325 (61.60)	915 (60.80)	410 (63.47)	
Negative	826 (38.40)	590 (39.20)	236 (36.53)	
PR Status, n(%)				0.776
Positive	939 (43.65)	654 (43.46)	285 (44.12)	
Negative	1212 (56.35)	851 (56.54)	361 (55.88)	
T stage, n(%)				0.432
1	227 (10.55)	158 (10.50)	69 (10.68)	
2	718 (33.38)	506 (33.62)	212 (32.82)	
3	407 (18.92)	296 (19.67)	111 (17.18)	
4	799 (37.15)	545 (36.21)	254 (39.32)	
N stage, n(%)				0.935
0	379 (17.62)	268 (17.81)	111 (17.18)	
1	1063 (49.42)	747 (49.63)	316 (48.92)	
2	288 (13.39)	198 (13.16)	90 (13.93)	
3	421 (19.57)	292 (19.40)	129 (19.97)	
Bone metastasis, n(%)				0.635
Yes	1212 (56.35)	843 (56.01)	369 (57.12)	
No	939 (43.65)	662 (43.99)	277 (42.88)	
Brain metastasis, n(%)				0.922
Yes	165 (7.67)	116 (7.71)	49 (7.59)	
No	1986 (92.33)	1389 (92.29)	597 (92.41)	
Liver metastasis, n(%)				0.205
Yes	853 (39.66)	610 (40.53)	243 (37.62)	
No	1298 (60.34)	895 (59.47)	403 (62.38)	
Lung metastasis, n(%)				0.082
Yes	708 (32.91)	478 (31.76)	230 (35.60)	
No	1443 (67.09)	1027 (68.24)	416 (64.40)	
Surgery, n(%)				0.925

Table 1 (continued)

Characteristics	Total	Training group	Validation group	Р
	(<i>n</i> =2151)	(<i>n</i> =1505)	(n=646)	
Yes	869 (40.40)	609 (40.47)	260 (40.25)	
No	1282 (59.60)	896 (59.53)	386 (59.75)	
Radiotherapy, n(%)				0.862
Yes	740 (34.40)	516 (34.29)	224 (34.67)	
No/Unknown	1411 (65.60)	989 (65.71)	422 (65.33)	
Chemotherapy, n(%)				0.809
Yes	1728 (80.33)	1207 (80.20)	521 (80.65)	
No/Unknown	423 (19.67)	298 (19.80)	125 (19.35)	

A nomogram is a common clinical predictive tool in the cancer field and is used to visualize predictive models that accurately assess the prognosis of individual patients by integrating various disease characteristics [7]. Various nomograms that predict the prognosis of MBC have been developed for other subtypes, such as HER2-negative and triple-negative MBC [8, 9]. However, a prognostic nomogram for individuals with HER2+MBC does not exist. As a result, we developed a prognostic model to forecast breast cancer-specific survival (BCSS) and overall survival (OS) in patients with HER2+MBC using clinical data from the Surveillance, Epidemiology, and End Results (SEER) database, aiming to provide a convenient Web-based version of the program for easy use by clinicians.

Materials and methods

Source of data

Patient data were gathered from the SEER database (http://seer.cancer.gov/), the official source of information on cancer incidence and survival in the United States. This database provides demographic, clinicopathological, and survival data. We utilised the SEER Research Data, 17 Registries (2000–2020) version released in November 2022 and collected data from patients diagnosed with breast cancer using SEER*Stat software (version 8.4.2). The 17 cancer registries cover approximately 26.5% of the U.S. population, with a broad geographic distribution representing the East and West coasts as well as the Midwest and South regions of the U.S. The patient population is remarkably heterogeneous, encompassing a wide range of races and ethnicities, genders, ages, socioeconomic statuses, and healthcare accessibility.

Patients

This retrospective analysis evaluated data from patients with breast cancer diagnosed between 2010 and 2015. The criteria for inclusion comprised the following: (1) female sex; (2) age \geq 20 years; (3) diagnosed with breast cancer based on site and morphology (site recode ICD-O-3/WHO 2008); (4) microscopically confirmed cancer; (5) breast cancer was the first primary and only malignant

cancer identified; (6) HER2+; and (7) with distant metastasis. The exclusion criteria were: (1) those with unavailable information; (2) breast cancer was confirmed by death certificate or autopsy only; and (3) patients with stage T0, Tx, and Nx.

Patients who fit these requirements were enrolled, and a 7:3 randomization process was used to classify them into training and validation groups. Figure 1 illustrates the study screening procedures. The SEER database is openly accessible; thus, this study did not require informed consent.

Predictors

We extracted and reclassified variables from the SEER database, including age; marital status; race; histological type; grade; ER and PR status; American Joint Committee on Cancer (AJCC) 7th edition tumour (T) and node (N) stages; surgery; radiotherapy; chemotherapy; and bone, brain, liver, and lung metastases.

Outcomes

The primary outcomes were BCSS and OS. The period from diagnosis to breast cancer-related death was defined as BCSS and that from diagnosis to either all-cause death or the date of the last follow-up as OS.

Statistical analyses

The training and validation groups were randomly assigned (7:3 ratio) using R software (version 4.3.2; R Core Team, Vienna, Austria), and the two groups' baseline characteristics were compared using the chi-squared test. To determine independent prognostic factors influencing BCSS and OS, sixteen variables were analysed using univariate Cox regression; those with *p*-values < 0.05 were included in the multivariate Cox analysis.

Subsequently, using the 'rms' and 'survival' packages in R, we constructed nomograms based on the independent prognostic factors for predicting one-, three-, and five-year BCSS and OS for HER2+MBC. The concordance indices (C-index) and time-dependent receiver operating characteristic (ROC) curves with the areas under the curves (AUCs) were employed for assessment

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	BCSS			os			BCSS			S		
	HR	95%CI	Р	Ħ	95%CI	٩	HR	95%CI	٩	Ħ	95%CI	μ
Age												
20–39	Ref			Ref			Ref			Ref		
40–59	1.456	1.150–1.843	0.002	1.418	1.134–1.773	0.002	1.249	0.980– 1.584	0.073	1.221	0.973-1.533	0.085
60-79	2.102	1.654–2.671	< 0.001	2.119	1.690–2.657	< 0.001	1.699	1.322- 2.183	< 0.001	1.735	1.369–2.199	< 0.001
80+	4.573	3.352-6.238	< 0.001	4.576	3.411-6.138	< 0.001	2.791	1.965– 3.964	< 0.001	2.864	2.054-3.994	< 0.001
Marital status	J. C			y° C			Jeu			Jeu		
Single	1.229	1.032–1.464	0.021	1.247	1.057-1.471	0.009	1.133	0.946– 1 358	0.176	1.147	0.967–1.361	0.116
Widowed/divorced/other	1.637	1.411-1.900	< 0.001	1.623	1.408–1.870	< 0.001	1.250	1.068– 1.462	0.005	1.240	1.068–1.440	0.005
Race								1				
Black	Ref			Ref			Ref			Ref		
White	0.770	0.650-0.911	0.002	0.733	0.625-0.859	< 0.001	0.722	0.606– 0.861	< 0.001	0.686	0.583-0.809	< 0.001
Other	0.712	0.549–0.924	0.010	0.660	0.515-0.846	0.001	0.800	0.614- 1.041	0.097	0.735	0.571-0.946	0.017
Histologic type												
IDC	Ref			Ref			Ref			Ref		
ILC	1.350	0.957–1.904	0.087	1.250	0.892-1.754	0.195	/	/	~	/	/	/
Mixed	0.885	0.625-1.255	0.494	0.846	0.603-1.186	0.331	/	/	~	/	/	/
Other	1.052	0.820-1.352	0.687	1.075	0.850-1.359	0.547	/	/	/	/		/
Grade												
_	Ref			Ref			Ref			Ref		
_	1.063	0.660-1.712	0.801	1.080	0.687-1.697	0.739	/	/	/	/	/	/
=	1.209	0.757-1.933	0.427	1.192	0.764-1.860	0.440	/	/	~	/	/	/
	2.461	0.833-7.276	0.103	2.210	0.755-6.467	0.148	/	/	/	/	/	/
ER Status												
Positive	Ref			Ref			Ref			Ref		
Negative	1.337	1.171-1.528	< 0.001	1.336	1.177–1.517	< 0.001	1.419	1.193– 1.688	< 0.001	1.442	1.223–1.701	< 0.001
PR Status												
Positive	Ref			Ref			Ref			Ref		
Negative	1.257	1.100-1.436	0.001	1.243	1.095–1.411	0.001	1.110	0.936– 1.316	0.231	1.092	0.929–1.284	0.286

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Table 2 (continued)												
Variables	Univari	ate COX analysis						Mu	Iltivariate CO	X analysis		
	BCSS			SO			BCSS			SO		
	Ħ	95%CI	Р	Ħ	95%CI	Ρ	Ħ	95%CI	Р	Ħ	95%CI	Р
T stage												
	Ket			Ket			Ket			Ket		
Т2	1.143	0.889–1.470	0.296	1.143	0.901-1.450	0.273	1.226	0.951– 1.579	0.115	1.225	0.964–1.558	0.097
T3	1.206	0.921–1.579	0.173	1.202	0.931-1.552	0.159	1.143	0.870- 1.500	0.337	1.139	0.880–1.474	0.324
Т4	1.758	1.377–2.244	< 0.001	1.742	1.382–2.196	< 0.001	1.507	1.177– 1.929	0.001	1.477	1.169–1.867	0.001
N stage												
NO	Ref			Ref			Ref			Ref		
N1	0.878	0.734-1.051	0.156	0.872	0.735-1.033	0.112	/	/	/	/	/	/
N2	0.846	0.667-1.074	0.169	0.851	0.680-1.066	0.160	/	/	/	/	/	/
N3	0.928	0.751-1.147	0.491	0.911	0.745-1.114	0.365	/	/	/	/	/	/
Bone metastasis												
Yes	Ref			Ref			Ref			Ref		
No	0.854	0.747–0.976	0.020	0.864	0.761-0.981	0.024	0.726	0.630-	< 0.001	0.743	0.649–0.851	< 0.001
Brain metastasis								0000				
Yes	Ref			Ref			Ref			Ref		
No	0.534	0.427-0.667	< 0.001	0.554	0.447-0.687	< 0.001	0.545	0.435– 0.684	< 0.001	0.582	0.465-0.727	< 0.001
Liver metastasis												
Yes	Ref			Ref			Ref			Ref		
No	0.693	0.607–0.791	< 0.001	0.733	0.646-0.831	< 0.001	0.625	0.545-0.717	< 0.001	0.661	0.580-0.753	< 0.001
Lung metastasis												
Yes	Ref 0661		500 0 1	Ref 0640		100 0 1	Ref 0 771		100.01	Ref 0.701	, , , , , , , , , , , , , , , , , , ,	100.01
2						- 00.0 /	17/0	0.832	0000	5		- 00.0
Surgery												
res No	кет 1.959	1.701-2.254	< 0.001	кет 1.941	1.699–2.218	< 0.001	кет 1.450	1.251–	< 0.001	кет 1.484	1.284–1.717	< 0.001
								1.681				
Radiotherapy	9 - U									J - C		
res No/Unknown	кет 1.140	0.992-1.311	0.065	кет 1.180	1.033-1.348	0.015	Ket /		_	кет 0.927	0.803-1.070	0.301
Chemotherapy												
Yes No/Unknown	Ref 2.683	2.307–3.120	< 0.001	Ref 2.621	2.268–3.028	< 0.001	Ref 2.318	1.952-	< 0.001	Ref 2.219	1.882–2.615	< 0.001
								2.752				

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of discrimination capacity; a higher AUC value indicated better discrimination [10]. Model-predicted survival was compared with observational survival using calibration curves [11]. Decision curve analysis (DCA) was applied to evaluate the clinical practicability of each model.

We also utilized the 'nomogramFormula' package in R to calculate the total points for all patients. The optimal cut-off values of the total points were then calculated using X-tile software (version 3.6.1) (https://medicine.y ale.edu/lab/rimm/research/software/), and the patients were stratified into low-, intermediate-, and high-risk groups depending on this value. Kaplan–Meier survival curves and log-rank tests were performed to evaluate the nomograms' capacity to stratify BCSS and OS risk. Lastly, we used the 'DynNom' package in R and shinyapps.io (https://www.shinyapps.io/) to develop and publish our Web-based dynamic nomograms. All statistical analyses were performed using R software, and *p*-values<0.05 were considered statistically significant.

Results

Patient characteristics

A total of 2,151 patients were assessed and randomly assigned to training (n=1,505) or validation (n=646)groups. The two groups did not differ when considering their baseline characteristics (Table 1). Most patients were 40-59 years old (47.00%), followed by the 60-79 years (34.03%), 20-39 years (12.65%), and >80 years (6.32%) groups. Furthermore, 46.35% and 22.04% of the patients were married and single, respectively, and 1,578 (73.36%) were Caucasian. Invasive ductal carcinoma was the predominant pathological type (85.68%), and histological grades I-IV accounted for 2.00%, 33.84%, 63.88%, and 0.28%, respectively. Overall, 61.60% and 43.65% of the tumours were ER- and PR-positive, respectively. Stages T1 to T4 accounted for 10.55%, 33.38%, 18.92%, and 37.15% of cases, respectively. Stages N0 to N3 accounted for 17.62%, 49.42%, 13.39%, and 19.57% of cases, respectively. Bone, brain, liver, and lung metastases occurred in 56.35%, 7.67%, 39.66%, and 32.91% of patients, respectively. Finally, 40.40%, 34.40%, and 80.33% underwent surgery, radiotherapy, and chemotherapy, respectively.

Univariate and multivariate Cox analyses for prognosis

In the training group, 12 variables—including age; marital status; race; ER status; PR status; T stage; surgery; chemotherapy; and bone, brain, liver, and lung metastases—were significantly correlated with both BCSS and OS in the univariate Cox analysis (all p < 0.05); radiotherapy only correlated with OS. Subsequently, ten variables were identified as independent prognostic factors in the multivariate Cox analysis, including age; marital status; race; ER status; surgery; chemotherapy; and bone, brain, liver, and lung metastases (Table 2).

Nomogram construction and validation

Based on the variables determined above, nomograms were constructed for patients with HER2+MBC to predict one-, three-, and five-year BCSS and OS (Fig. 2). Take the following case as an example: a 36-year-old married white patient with ER-positive (ER+) HER2+MBC that had metastasized to the bone but not to the brain, liver, or lung was treated with chemotherapy but not surgery. In this case, the total nomogram scores for BCSS and OS were 70.57 and 68.34, respectively. Thus, the one-, three-, and five-year BCSS probabilities were 93%, 82%, and 72%, respectively, and the OS probabilities were 92%, 80%, and 69%, respectively.

The BCSS and OS C-indices in the training group were 0.707 (95% confidence interval [CI]: 0.690–0.724) and 0.702 (95% CI: 0.686–0.718), respectively; in the validation group, they were 0.731 (95% CI: 0.707–0.755) and 0.725 (95% CI: 0.725–0.748), respectively. We also calculated the AUC values to assess the models' discrimination abilities. For the BCSS nomogram, the one-, three-, and five-year BCSS AUCs in the training group were 0.823, 0.756, and 0.743, respectively; in the validation group, they were 0.841, 0.814, and 0.780, respectively (Fig. 3a, b). For the OS nomogram, the one-, three-, and five-year OS AUCs in the training group were 0.811, 0.754, and 0.742, respectively; in the validation group, they were 0.834, 0.813, and 0.777, respectively (Fig. 3c, d).

Furthermore, both the training and validation groups' calibration curves comparing the one-, three-, and fiveyear predicted and observed BCSS and OS showed a high degree of consistency (Fig. 4). The DCA enables clinicians to select the optimal model. The benefit curves of the BCSS and OS nomograms were greater than those of the reference strategies, indicating that the models have practical clinical applications (Fig. 5).

Risk stratification and web-based dynamic nomogram development

To stratify risk, the optimal cut-off values of the patients' total nomogram scores were calculated with X-tile software. The BCSS total score cut-off values were 151.4 and 254.1 for categorizing patients into low-risk (score ≤ 151.4), intermediate-risk ($151.4 < \text{score} \leq 254.1$), and high-risk (score ≥ 254.1) groups (Fig. 6a, b). The OS cut-off values were 148.7 and 244.4 for categorizing patients into low-risk (score ≤ 148.7), intermediate-risk ($148.7 < \text{score} \leq 244.4$), and high-risk (score ≥ 244.4) groups (Fig. 6c, d).

Kaplan–Meier survival curves were created based on the above results. The median BCSS for the low-, intermediate-, and high-risk groups in the training group



Fig. 2 Nomograms for predicting survival in patients with HER2 + MBC. The nomograms for predicting one-, three-, and five-year BCSS (a) and OS (b). HER2 + MBC, human epidermal growth factor receptor 2-overexpressing metastatic breast cancer; BCSS, breast cancer-specific survival; OS, overall survival



Fig. 3 Time-dependent ROC curves of the nomograms. ROC curves of the nomogram for predicting one-, three-, and five-year BCSS in the training group (a) and validation group (b). ROC curves of the nomogram for predicting one-, three-, and five-year OS in the training group (c) and validation group (d). ROC, receiver operating characteristic; BCSS, breast cancer-specific survival; OS, overall survival

were 95, 33, and 5 months, respectively, and 95, 26, and 5 months in the validation group, respectively (Fig. 7a, b). The median OS for the low-, intermediate-, and high-risk groups in the training group were 82, 28, and 4 months, respectively, and 78, 24, and 6 months in the validation group, respectively (Fig. 7c, d). The survival outcomes significantly differed in the Kaplan–Meier survival curves for both BCSS and OS (p<0.001).

Finally, we established and published Web-based dynamic nomograms for BCSS (Fig. 8a) and OS (Fig. 8b) for patients with HER2+MBC, accessible at:

 (https://her2mbcnomo.shinyapps.io/DynNomapp/); and.

2) (https://her2mbcnomo.shinyapps.io/modelos/).

Discussion

Patients with a HER2+status commonly have a poor prognosis owing to the high invasiveness and recurrence rate associated with this cancer subtype. HER2+MBC remains incurable; therefore, a reliable prognostic model is crucial. The study by Wang et al., including 1174 patients, demonstrated that the histological subtype was a key factor in HER2+MBC prognosis but did not

construct a model that could individualise and accurately predict BCSS and OS [12]. The cohort study by Fan et al. included 34,819 patients with HER2+breast cancer and reported a nomogram to forecast BCSS. Although the C-index of this nomogram was as high as 0.853 (95% CI: 0.845–0.861), it did not explicitly target the high-risk MBC group [13]. Lyu et al. constructed a BCSS prognostic model based on a subgroup of 1204 HER2+patients with breast cancer with bone metastases, which possessed strong accuracy, with C-indexes of 0.74 and 0.77 for the training and validation groups, respectively. However, this nomogram excluded other distant metastases and, therefore, was not applicable to all metastatic cases [14]. Lin et al. included 1680 patients with HER2+MBC as a training cohort and constructed the first nomogram for predicting the probability of OS based on independent prognostic factors of demographic and clinicopathologic characteristics. In the training and validation cohorts, the C-index was 0.70 (95% CI, 0.68-0.72) and 0.68 (95% CI, 0.65–0.72), respectively [15]. In our study, 1505 patients were included in the training group, and a total of 10 demographic and clinicopathologic characteristics and treatment modalities were demonstrated as independent



Fig. 4 Calibration curves of the nomograms. Calibration curves for predicting one- (**a**, **d**), three- (**b**, **e**) and five-year (**c**, **f**) BCSS in the training and validation groups, respectively. Calibration curves for predicting one- (**g**, **j**), three- (**h**, **k**) and five-year (**i**, **l**) OS in the training and validation groups, respectively. BCSS, breast cancer-specific survival; OS, overall survival

prognostic factors. Based on these findings, we developed nomograms for BCSS and OS that had higher predictive accuracy and better clinical applicability than the previous models. Furthermore, based on the nomograms' total scores, we identified the optimal cut-off values for low-, intermediate-, and high-risk stratification, which could help clinicians accurately recognize high-risk patients and promptly adjust their therapeutic schedule. Finally, we published easy-to-use Web versions of the nomograms for convenience and clinical application.

Age, marital status, and race were key demographic variables used in the models in this study. Age was a major predictor of survival in patients with MBC, consistent with previous studies [16]. Younger patients with breast cancer are more likely to have larger tumour diameters, more axillary lymph node involvement, poorer pathologic staging, and more distant metastases than older patients, leading to a poorer prognosis, which is contrary to our study's results [17, 18]. We found that increased age negatively affected BCSS and OS, possibly because older individuals often undergo nonstandard treatments owing to their poor physical condition that results in a lower tolerance for surgery, chemotherapy, and radiotherapy [19]. Furthermore, immune senescence, defined as a decline in immunity with age, can lead to poor prognosis in older adults owing to decreased defences against tumours, even if tumour aggressiveness decreases with age [20–22]. Consequently, age



Fig. 5 DCA of the nomograms. DCA of the nomogram for predicting one- (a, d), three- (b, e) and five-year (c, f) BCSS in the training and validation groups, respectively. DCA of the nomogram predicting one- (g, j), three- (h, k) and five-year (i, l) OS in the training group and validation group, respectively. DCA, decision curve analysis; BCSS, breast cancer-specific survival; OS, overall survival

remains an important independent outcome predictor for patients with HER2+MBC.

Breast cancer is a systemic disease, and the influence of sociopsychological factors, such as marital status, on its prognosis cannot be underestimated. Similar to previous studies, we found that married patients exhibited an increased survival rate compared with single and widowed patients. Married patients, with the support of their spouses and children, are potentially more accepting of surgery, chemotherapy, and radiation therapy and adhere more strictly to their clinicians' treatment strategies. Moreover, the severity of depressive symptoms is significantly associated with elevated levels of circulating pro-inflammatory factors. Married patients may receive



Fig. 6 The optimal cut-off values for the nomograms. Optimal cut-off values for the patients' total scores based on the BCSS and OS nomograms using X-tile analysis (**a**, **c**). The histograms for describing the distribution of patients based on the cut-off values (**b**, **d**). BCSS, breast cancer-specific survival; OS, overall survival

social support from their families, reducing the risk of depressive symptoms and improving their prognosis [23–25]. We also identified race as an independent prognostic predictor of HER2+MBC. Black race has been associated with poor prognosis and low survival of MBC [26, 27]. The primary reasons for racial differences are socioeconomic support and tumour characteristics [27].

Clinicopathological features, such as ER status and bone, brain, liver, and lung metastases, are closely related to prognosis. Our study found that ER positivity was a protective prognostic factor, consistent with previous studies that suggested an interaction between ER-negative (ER–) and HER2+statuses, leading to increased breast cancer invasiveness. Thus, patients with ER+and HER2+cancer have a less-invasive subtype and a wider range of therapeutic options, such as hormonal therapies [28, 29]. The ER status also affects the metastasis site. According to Cletus et al., patients with ER+breast cancer had a higher rate of bone metastasis than those with ER– subtypes, and patients with ER– breast cancers had a higher probability of visceral metastasis, especially liver metastasis. Consistent with these results, we identified bone metastasis as an independent prognostic factor, although it had the least impact on patient prognosis compared with the other identified factors. In contrast, patients with brain metastasis had a substantial decrease



Fig. 7 Kaplan–Meier survival curves after risk stratification. Kaplan–Meier survival curves of low-, intermediate-, and high-risk groups for the BCSS nomogram in the training (a) and validation groups (b). Kaplan–Meier survival curves of low-, intermediate-, and high-risk groups for the OS nomogram in the training (c) and validation groups (d). BCSS, breast cancer-specific survival; OS, overall survival

in BCSS and OS [26, 30]. Lastly, the Tumour Node Metastasis staging system is widely used for disease risk assessment, and the T stage was a meaningful prognostic factor in this study. However, it was not included in the models because we discovered that the prognosis for stage T2 was shown to be worse than that of stage T3, inconsistent with previous studies [16].

We found that surgery and chemotherapy prolonged BCSS and OS in patients with HER2+MBC. The National Comprehensive Cancer Network guidelines recommend targeted therapy combined with chemotherapy for HER2+metastatic disease; however, surgical treatment of patients with MBC is controversial. Surgery improves the prognosis initially by reducing the tumour load via primary tumour resection, blocking the source of distant tumour metastasis [31, 32]. Moreover, surgical resection of tumour lesions decreases the number of immunosuppressive factors released by the tumour, restores immune activity, and assists in systemic therapy [33]. However, surgery dramatically reduces the levels of anti-angiogenic factors and growth factor inhibitors, accelerating tumour recurrence, and postoperative complications and delays in adjuvant therapy can affect prognosis [34-36].

In contrast, the MF07-01 study reported improved 40-month survival in patients with stage IV breast cancer treated with locoregional therapy, and a French multicentre retrospective study found that locoregional treatment prolonged OS in patients with MBC, especially those with HER+cancer [37, 38]. Furthermore, a randomized controlled trial in India and a prospective phase III trial (ABCSG-28) demonstrated that surgery improved OS in patients with stage IV breast cancer [39, 40]. In this study, surgical treatment improved BCSS and OS.

The advent of anti-HER2 therapy has dramatically improved the prognosis of patients with HER2+MBC, with treatment options including monoclonal antibodies against HER2, tyrosine kinase inhibitors, and antibodydrug conjugates (ADCs) [41]. The CLEOPATRA study validated dual HER2-blocking therapy with docetaxel combined with pertuzumab and trastuzumab as the preferred first-line treatment option for HER2+MBC [42]. The PHENIX and PHOEBE studies together validated the superior efficacy of the second-line pyrotinib and capecitabine combination [43, 44]. Lapatinib also showed a trend toward better outcomes in patients with brain metastases when no new anti-HER2 agents were

a Dynamic Nomogram

Age	Survival plot Predicted Survival Numerical Summary Model Summary	
20-39	Estimated Survival Proba	bility
	1.00	
Marital.status		
Married 👻		
Race	0.73	
White •		
ER status	6 0.00	
Positive		
	0.25-	
vas		
	0.00	
Brain.metastasis	o 50 Follow Up Time	500
No		
Liver.metastasis		
No		
No		
Surgery		
No		
Chemotherapy		
Yes 👻		
Predicted Survival at this Follow Up:		
ime		
0 3		
0 14 28 42 55 72 54 68 112 123 131		
Alpha blending (transparency)		
Predict		
Predict		
Predict Press Guit to exit the application		

plot Predicted Survival Numerical Summary

95% Confidence Interval for Survival Probability

b Dynamic Nomogram



Model S

Fig. 8 A screenshot of the Web-based dynamic nomograms. The nomograms for BCSS (a) and OS (b) in patients with HER2 + MBC. HER2 + MBC, human epidermal growth factor receptor 2-overexpressing metastatic breast cancer; BCSS, breast cancer-specific survival; OS, overall survival

available [45]. The EMILIA study established trastuzumab emtansine (T-DM1) as the second-line treatment for HER2+MBC [46]. Moreover, the DESTINY-03 study showed trastuzumab deruxtecan (T-DXd) significantly improve progression-free survival compared with T-DM1, making it a preferred option for second-line treatment [47]. However, the SEER database mainly provides clinical characteristics and survival data and does not include detailed information on HER2-targeted therapies and specific chemotherapeutic regimens, which may limit the accuracy and completeness of our model. While current studies suggest that trastuzumab and pertuzumab in combination with chemotherapy are commonly preferred first-line treatment options for patients with HER2+MBC [48, 49], it is worth noting that patients receiving chemotherapy does not mean that they also receive HER2-targeted therapy. Therefore, it is crucial to obtain relevant treatment data regarding anti-HER2 agents. We expect that future experiments will further refine relevant explorations in this area.

There are some limitations to our study. First, this study adopted a retrospective design, relying on the strength of the SEER database, which potentially led to selection bias. Second, the SEER database contains insufficient information regarding factors potentially affecting prognosis, such as HER2-targeted treatment plans, detailed chemotherapy plans, and Ki-67 expression. Third, the SEER database only provided information on distant metastatic sites and molecular types since 2010, and partial treatment data are still incomplete after 2015. Therefore, only data from 2010 to 2015 were selected for our analyses, so the most recent treatment regimen may have had an impact on patient outcomes. Lastly, an independent database was not used to externally validate the nomograms. Further large prospective studies and the integration of more relevant variables from real-world patient data, especially systemic treatment regimens, are required to enhance and refine our predictive models.

Conclusion

In summary, we constructed two nomograms to predict individual survival among female patients with distant metastatic HER2+breast cancer using risk stratification. Moreover, we published the models online for general use by clinicians to accurately predict one-, three-, and five-year BCSS and OS and distinguish high-risk patients in this population to optimize clinical decision-making.

Abbreviations

Antibody-drug conjugate
American Joint Committee on Cancer
American Joint Committee on Cancer
Areas under the curves
Breast cancer-specific survival
Confidence interval
Concordance index
Decision curve analysis
Oestrogen receptor
Oestrogen receptor-negative
Oestrogen receptor-positive
Human epidermal growth factor receptor 2
Human epidermal growth factor receptor 2-overexpressing
Metastatic breast cancer
Node stage
Overall survival
Progesterone receptor
Receiver operating characteristic
Surveillance, Epidemiology, and End Results
Trastuzumab emtansine
Trastuzumab daruvtacan
lumour stage

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Author contributions

YC contributed to the conceptualization of the study, methodology, visualization, validation of the data, and writing the original draft. YQ contributed to the conceptualization of the study, methodology, validation of the data, and writing the original draft. HS was responsible for the software and visualization. SY was responsible for the software and visualization. JL contributed to data curation. WW contributed to the review and editing of the manuscript, supervision, and funding acquisition. All authors have approved the final version of the manuscript. Authors YC and YQ share first authorship.

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Data availability

The SEER database (http://seer.cancer.gov/), which is publicly accessible, provided the data analyzed in this study.

Declarations

Ethics approval and consent to participate

Our study was approved by the Ethics Review Board of The Affiliated Lihuili Hospital, Ningbo University, Ningbo (registration number: KY2024ML045). Informed consent was not required as the data were collected from a publicly available repository.

Consent for publication

Not applicable.

Competing interests

The authors declare no competing interests.

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