



Establishment and validation of a prognostic model for gastric cancer patients of Hebei Province in China

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Background: Hebei Province is a high-risk area for gastric cancer in China, and there is currently no survival prediction model for gastric cancer patients in Hebei Province. This study aimed to build the best survival prediction model for gastric cancer patients in Hebei Province.

Methods: The development dataset included 1,993 hospitalized gastric cancer patients from the Hebei Cancer Registration Project during 2016 and 2017. Three tree-based machine learning methods [survival trees (ST), random survival forests (RSF), and gradient boosting machines (GBM)] and Cox, were used to develop the models by ten-fold cross validation with 200 iterations. California Chinese hospitalized gastric cancer patients were used as external test models. In addition, we compared the multivariable group with the Tumor Node Metastasis (TNM) group.

Results: The 3- and 5-year cancer-specific survival (CSS) rates of the development dataset were 57.07% and 44.48%, respectively. For predicting the 3-year CSS rates of gastric cancer patients of multivariable group, the C-indexes in train datasets were 0.75, 0.72, 0.79 and 0.76 for Cox, ST, RSF and GBM. Multivariable group performed better than TNM group. The predictive ability of Cox and RSF were superior to ST and GBM. A nomogram was established to predict the 3- and 5-year CSS rates of gastric cancer patients.

Conclusions: The nomogram was useful for facilitating clinicians to predict the survival of gastric cancer patients, and identifying high-risk patients so as to adopt more reasonable treatment plans.

Keywords: Gastric cancer; prognostic model; machine learning; Hebei Province; China

Submitted Aug 11, 2023. Accepted for publication Nov 30, 2023. Published online Dec 13, 2023.

doi: 10.21037/cco-23-85

View this article at: <https://dx.doi.org/10.21037/cco-23-85>

Introduction

Background

Gastric cancer is one of the top ten common malignant tumors worldwide (1). According to Global Cancer Statistics 2020 (GLOBOCAN 2020) (1), 43.9% of new cases and

48.6% of deaths occurred in China. Hebei Province, located in North China, is the only province that includes all landforms in China. The incidence and mortality of gastric cancer of Hebei Province are higher than that of China (2). In recent decades, the incidence and mortality of gastric cancer patients are declining in many developed countries,

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but it still causes a huge disease burden (3). More than 90% of early gastric cancer patients survive for 5 years or more after receiving surgery (4,5). However, the prognosis of late gastric cancer patients is poor, with a 5-year survival rate of about 20% for stage III patients and about 7% for stage IV patients (6). The Tumor Node Metastasis (TNM) staging system, established by the American Joint Committee on Cancer (AJCC) and Union for International Cancer Control (UICC), is a practical method for evaluating the prognosis of gastric cancer patients and is widely used worldwide (6). However, TNM staging system, which is only based on tumor infiltration, tumor size, lymph node metastasis, and distant metastasis, has inevitable limitations. The patients within the same stage may have different survival outcomes, and patients in different stages may also have the same survival outcome (7). In addition, many studies have shown that age, gender, tumor size, and therapy are closely related to the survival of gastric cancer patients (7-9). The occurrence and development of gastric cancer are very complex. Many clinical and pathological features can influence prognosis, most of which are multidimensional and non-linear relationships (10). Therefore, it is necessary to integrate prognostic factors selected from comprehensive clinical information and establish a more accurate survival prediction model for gastric cancer patients.

Highlight box

Key findings

- The multivariable group outperformed Tumor Node Metastasis (TNM) group. Cox and random survival forest performed better than survival tree and gradient boosting machine. The nomogram may facilitate clinicians to predict survival of gastric cancer patients in Hebei Province.

What is known and what is new?

- Machine learning has been widely applied. It's unknown whether existing models are suitable for the prognosis of gastric cancer patients in Hebei Province.
- Three tree-based machine learning and Cox, were used to build gastric cancer patients prognostic models. Additionally, California Chinese hospitalized gastric cancer patients in Surveillance, Epidemiology, and End Results (SEER) database were used as external test dataset. We compared multivariable group with TNM group.

What is the implication, and what should change now?

- This study fills in the blank of prognostic models for hospitalized gastric cancer patients in Hebei Province. We will try to compare more models to establish a more comprehensive prognostic model.

Rationale and knowledge gap

In order to obtain better models for predicting patients' survival, many survival prediction tools have been developed, including parametric, semi-parametric, and non-parametric methods. The parametric method requires that the survival time meet the normal distribution, and the accelerated failure time (AFT) model is commonly used (11). Most survival data are non-normal distribution, so parametric method is not commonly used. The semi-parametric method is based on the Cox proportional hazard (PH) regression model, which has been applied in many studies to predict the survival of gastric cancer (8,12-18). However, due to the need to comply with the PH assumption, the use of Cox is limited. In order to eliminate these constraints, a series of non-parametric methods have been developed for cancer survival prediction, which can consider the interaction effects among variables. In the non-parametric methods, the most common methods were tree-based machine learning methods [including survival trees (ST), random survival forests (RSF), gradient boosting machines (GBM), etc.] (19-30). The ST is a machine learning method that constructs a tree structure by splitting nodes by maximizing survival differences until all terminal nodes containing only the minimum number of unique events (31,32). Both RSF and GBM are combined of a large number of ST. RSF uses the bootstrap method to extract sub samples from the original samples to construct a ST, averaging the cumulative risk function of each ST and ultimately obtaining the total cumulative risk function (33). GBM is a machine learning method based on gradient descent boosting. The fundamentals of GBM are training a new ST according to the negative gradient information of the loss function based on the current ST, and combining the trained newborn ST with the existing ST (34).

Machine learning has been widely applied in data analysis such as medical care, and is an effective tool for improving clinical strategies. Banerjee *et al.*, Qiu *et al.*, Du *et al.*, van Zutphen *et al.* and Lin *et al.* showed that RSF performs better than Cox in predicting the survival of thyroid cancer, glioma, oropharyngeal cancer, colorectal cancer and liver cancer (35-40). Samara's research shows that RSF performs better than linear support vector machine (SVM), Adaboost, Bagging and other machine learning algorithms in predicting the survival of glioblastoma (41). At present, there is no research on the survival prediction of gastric cancer patients in Hebei Province, and it is unknown whether the existing survival prediction models are suitable for the survival prediction of gastric cancer patients in

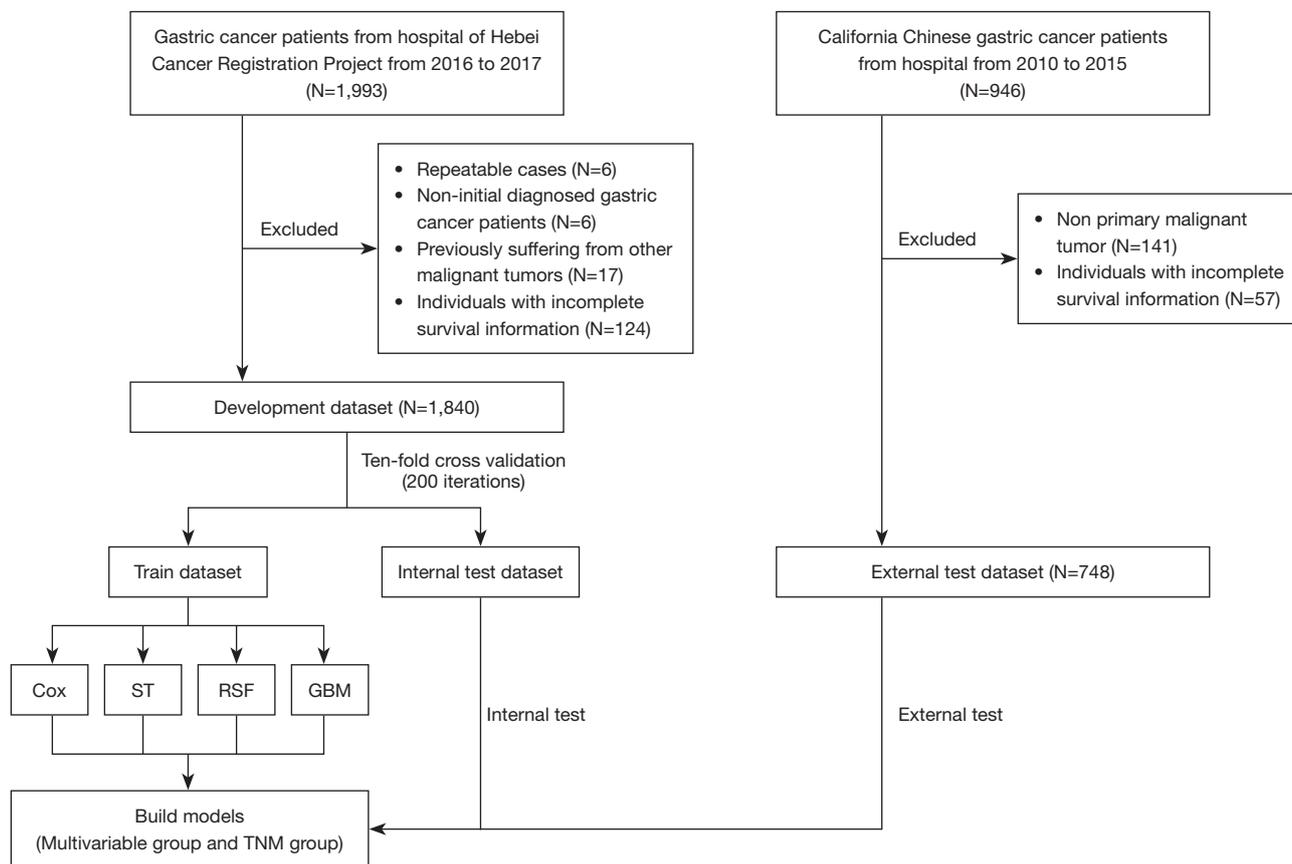


Figure 1 Flowchart of study design and patient selection. ST, survival trees; RSF, random survival forests; GBM, gradient boosting machines; TNM, Tumor Node Metastasis.

Hebei Province.

Objective

This study was based on the information of hospitalized gastric cancer patients from the Hebei Cancer Registration Project during 2016 and 2017. Three tree-based machine learning methods (ST, RSF, and GBM) and the Cox PH regression model, were used to build gastric cancer patients survival prediction models. Additionally, California Chinese gastric cancer patients from hospitals in the Surveillance, Epidemiology, and End Results (SEER) database were used as external test dataset. In addition, we compared the multivariable group with TNM group. This study aimed to build the best survival prediction model, identify high-risk gastric cancer patients as soon as possible, and provide reference for clinical doctors to develop specific treatment plans for patients and allocate medical resources reasonably.

This manuscript is written in accordance with the TRIPOD reporting checklist (available at <https://cco.amegroups.com/article/view/10.21037/cco-23-85/rc>).

Methods

Data source

Data of development dataset were recruited from Hebei Cancer Registration Project. We included 1,993 hospitalized gastric cancer patients who were diagnosed between 1 January 2016 to 31 December 2017. The following patients were excluded: (I) repeatable cases (N=6); (II) non-initial diagnosed cases (N=6); (III) previously suffered from other malignant tumors (N=17); (IV) individuals with incomplete survival information (N=124) (Figure 1).

Data of external test dataset were obtained from the “Incidence - SEER Research Plus Data, 17 Registries, Nov 2021 Sub (2000-2019)” of SEER database. To exclude

differences among hospital source data and population data, as well as differences between different races, we selected the hospitalized gastric cancer patients of California Chinese from 2010 to 2015. A total of 946 patients were included. The exclusion criteria are as follows: (I) non-initial diagnosed cases (N=141); (II) individuals with incomplete survival information (N=57) (*Figure 1*).

Predictors and outcomes

This study included 13 predictors, including gender, age, area, marital status, tumor site, grade, TNM stage, T (tumor) stage, N (node) stage, M (metastasis) stage, surgery, radiotherapy, and chemotherapy. The predictors of TNM stage, T stage, N stage, M stage were referring to AJCC Clinical Stage, 7th edition.

The outcome variables included “survival months” and “3-year cancer-specific survival (CSS) status” or “5-year CSS status”.

Data of development dataset were collected and strictly controlled by trained cancer registration professionals. The survival outcomes were obtained by active and passive follow-up. Passive follow-up was mainly obtained through all-causes-of-death survey database, rehospitalization information, outpatient information and medical insurance information. Active follow-up was obtained by trained professionals trimonthly. The deadline date of follow-up was December 31, 2022.

Establishment and validation of the model

The development dataset was used to train and internally test the model with ten-fold cross validation of 200 iterations. The external test dataset was used for external test. Four models (including Cox, ST, RSF and GBM) were used to establish the survival prediction model for gastric cancer in Hebei Province. Additionally, we compared the multivariable group (including multiple variables) with TNM group (including AJCC 7th edition T stage, N stage, and M stage) (*Figure 1*).

Evaluation indicators of the model

The Harrell’s consistency index (C-index) and area under the receiver operating characteristic curve (AUC) were used to evaluate the model’s differentiation. The calibration curve was used to evaluate the consistency between the predicted values and the actual observed values. The 45-degree

straight gray line of calibration curve represents the perfect match between the observed (y-axis) and predicted (x-axis) survival probabilities. A closer distance between two curves indicates higher consistency.

Statistical analysis

Software R (version 4.1.2) was used for data analysis. “randomForestSRC” package was used to impute missing values. “survival”, “rpart”, “randomForestSRC” and “gbm” packages were used to develop the models. The value of $P < 0.05$ indicates a statistically significant difference.

Ethical statement

The study was conducted in accordance with the Declaration of Helsinki (as revised in 2013). For external test dataset from the SEER database, the authors obtained authorization to access the SEER Research Data supported by the National Cancer Institute with approval number 11241-Nov2021. Because public and anonymous data from the SEER database were used, informed patient consent was not required. For development dataset from Hebei Cancer Registration Project, the Ethics Committee of the Fourth Hospital of Hebei Medical University/the Tumor Hospital of Hebei Province has confirmed that no ethical approval is required. Individual consent for this retrospective analysis was waived.

Results

Demographic characteristics of gastric cancer patients

After inclusion and exclusion, 1,840 patients and 748 patients with gastric cancer were included in the development dataset and external test dataset, respectively. The 1-, 3-, and 5-year CSS rates of the development dataset were 82.61%, 57.07%, and 44.48%, respectively. The 1-, 3-, and 5-year CSS rates of the external test dataset were 68.58%, 47.99%, and 42.91%, respectively. The detail of demographic characteristics of the imputation data were shown in *Table 1*.

Figure 2 shows the proportion of stages of gastric cancer patients in the development dataset and external test dataset. The proportion of stage I, II, III, and IV of gastric cancer patients in the development dataset were 19.35%, 19.24%, 49.45%, and 11.96%, respectively. The proportion of stage I, II, III, and IV of gastric cancer patients in the external test dataset were 28.07%, 16.58%, 26.07%, and 29.28%, respectively. The proportion of stage III and IV

Table 1 Demographic characteristics of gastric cancer patients in development dataset and external test dataset

Variables	Development dataset, n (%)	External test dataset, n (%)	P value [†]
Total	1,840 (100.00)	748 (100.00)	
Gender			<0.001
Male	1,442 (78.37)	413 (55.21)	
Female	398 (21.63)	335 (44.79)	
Age (years)			<0.001
<55	373 (20.27)	131 (17.51)	
55–64	692 (37.61)	119 (15.91)	
65–74	626 (34.02)	183 (24.47)	
≥75	149 (8.10)	315 (42.11)	
Areas			<0.001
Urban	649 (35.27)	726 (97.06)	
Rural	1191 (64.73)	22 (2.94)	
Marital status			<0.001
Married	1,820 (98.91)	532 (71.12)	
Unmarried	6 (0.33)	75 (10.03)	
Others [‡]	14 (0.76)	141 (18.85)	
Tumor site			<0.001
Cardia	910 (49.46)	93 (12.43)	
Overlapping	31 (1.68)	53 (7.09)	
Other site	899 (48.86)	602 (80.48)	
Grade			0.494
G1–G2 [§]	519 (28.21)	221 (29.55)	
G3–G4 [¶]	1,321 (71.79)	527 (70.45)	
TNM stage			<0.001
I	356 (19.35)	210 (28.07)	
II	354 (19.24)	124 (16.58)	
III	910 (49.46)	195 (26.07)	
IV	220 (11.96)	219 (29.28)	
T stage			<0.001
T1	267 (14.51)	217 (29.01)	
T2	201 (10.92)	92 (12.30)	
T3–T4	1,372 (74.57)	439 (58.69)	
N stage			<0.001
N0	671 (36.47)	389 (52.01)	
N1	298 (16.20)	192 (25.67)	
N2	372 (20.22)	70 (9.36)	
N3	499 (27.12)	97 (12.97)	

Table 1 (continued)

Table 1 (continued)

Variables	Development dataset, n (%)	External test dataset, n (%)	P value [†]
M stage			<0.001
M0	1,626 (88.37)	538 (71.93)	
M1	214 (11.63)	210 (28.07)	
Surgery			<0.001
No	347 (18.86)	321 (42.91)	
Yes	1,493 (81.14)	427 (57.09)	
Radiotherapy			<0.001
No	1,790 (97.28)	592 (79.14)	
Yes	50 (2.72)	156 (20.86)	
Chemotherapy			<0.001
No	709 (38.53)	376 (50.27)	
Yes	1,131 (61.47)	372 (49.73)	

[†], P value was derived from the Chi-square test performed to compare the development dataset and external test dataset; [‡], others included divorced, separated and widowed; [§], G1–G2 represented well differentiated or moderately differentiated; [¶], G3–G4 represented poorly differentiated or undifferentiated. TNM, Tumor Node Metastasis; T stage, tumor stage; N stage, node stage; M stage, metastasis stage.

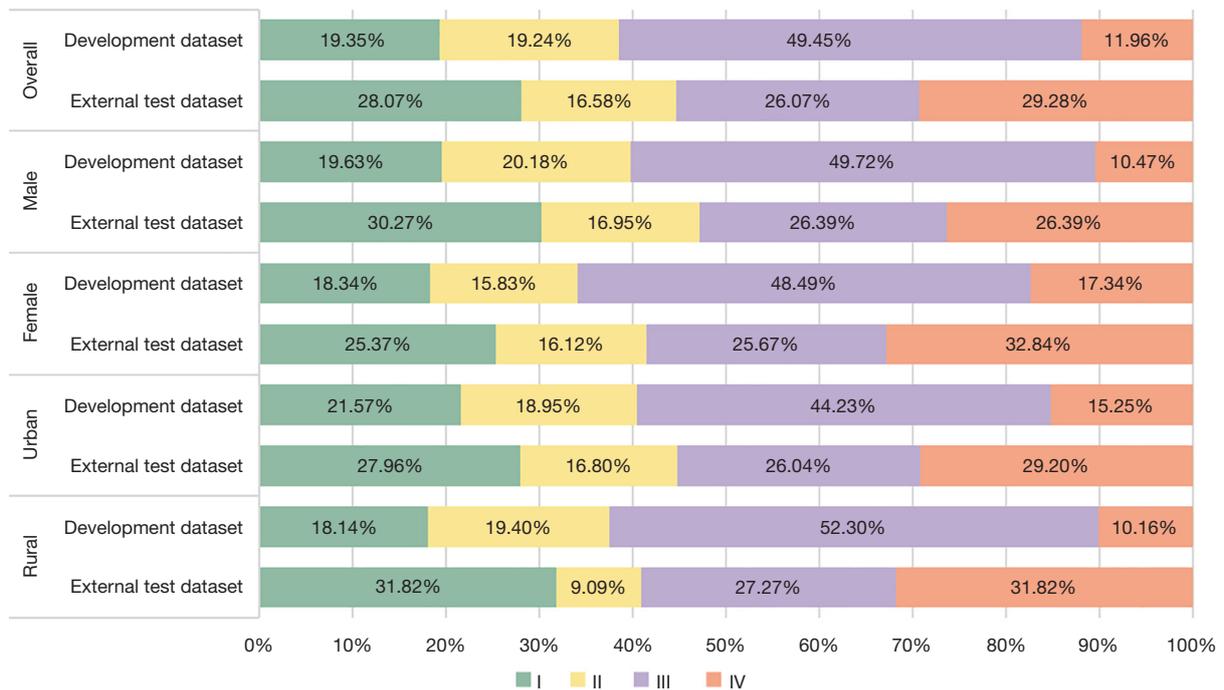


Figure 2 The proportion of stages of gastric cancer patients in the development dataset and external test dataset.

of overall patients in the development dataset (61.41%), was higher than that in the external test dataset (55.35%). The proportion of stage III and IV of female patients

(65.83%) was higher than that of male patients (60.19%) in the development dataset. The proportion of stage III and IV of rural patients (62.46%) was higher than that of

Table 2 The C-index of different models of 3- and 5-year CSS in gastric cancer patients

Cohort	Three-year CSS cohort, mean (95% CI)			Five-year CSS cohort, mean (95% CI)		
	Train dataset	Internal test dataset	External test dataset	Train dataset	Internal test dataset	External test dataset
Multivariable group						
Cox	0.75 (0.71–0.79)	0.73 (0.63–0.84)	0.75 (0.70–0.80)	0.71 (0.68–0.74)	0.71 (0.62–0.80)	0.76 (0.72–0.81)
ST	0.72 (0.69–0.76)	0.70 (0.59–0.80)	0.73 (0.68–0.77)	0.68 (0.65–0.71)	0.68 (0.60–0.77)	0.73 (0.68–0.77)
RSF	0.79 (0.75–0.82)	0.79 (0.68–0.90)	0.77 (0.72–0.81)	0.74 (0.71–0.77)	0.71 (0.62–0.80)	0.76 (0.72–0.81)
GBM	0.76 (0.72–0.79)	0.79 (0.69–0.90)	0.78 (0.74–0.83)	0.72 (0.69–0.75)	0.71 (0.62–0.80)	0.78 (0.73–0.82)
TNM group						
Cox	0.71 (0.67–0.75)	0.71 (0.59–0.82)	0.66 (0.61–0.71)	0.67 (0.64–0.71)	0.68 (0.58–0.78)	0.68 (0.63–0.73)
ST	0.69 (0.65–0.73)	0.70 (0.58–0.81)	0.60 (0.54–0.65)	0.66 (0.63–0.69)	0.66 (0.57–0.75)	0.68 (0.63–0.73)
RSF	0.72 (0.68–0.75)	0.70 (0.58–0.81)	0.68 (0.62–0.73)	0.68 (0.64–0.71)	0.68 (0.58–0.77)	0.67 (0.62–0.73)
GBM	0.71 (0.68–0.75)	0.70 (0.59–0.81)	0.67 (0.61–0.72)	0.67 (0.64–0.71)	0.68 (0.58–0.77)	0.67 (0.62–0.72)

CSS, cancer-specific survival; CI, confidence interval; ST, survival trees; RSF, random survival forests; GBM, gradient boosting machines; TNM, Tumor Node Metastasis.

urban patients (59.48%) in the development dataset. The proportion of stage III and IV of female patients (58.51%) was higher than that of male patients (52.78%) in the external test dataset. The proportion of stage III and IV of rural patients (59.09%) was higher than that of urban patients (55.24%) in the external test dataset.

Variables selection

According to Kaplan-Meier curves and log-rank tests on the above variables, it can be concluded that the ten variables (including TNM stage, gender, age, tumor site, grade, T stage, N stage, M stage, surgery, and radiotherapy) met the PH hypothesis (Figures S1,S2).

Considering that TNM stage was obtained by combining T stage, N stage and M stage, we excluded TNM stage from predictors. In the multivariable group of ST, RSF and GBM, we included 12 predictors (including gender, age, area, marital status, tumor site, grade, T stage, N stage, M stage, surgery, radiotherapy, and chemotherapy). In the multivariable group of Cox, we only included nine variables which met the PH assumption (including gender, age, tumor site, grade, T stage, N stage, M stage, surgery, and radiotherapy).

Model performance

Table 2 and Figure 3 showed the C-index of different models

for predicting 3- and 5-year CSS of gastric cancer patients in the multivariable group and TNM group. In the 3-year CSS cohort, the Cox [C-index: 0.75, 95% confidence interval (CI): 0.71–0.79], RSF (C-index: 0.79, 95% CI: 0.75–0.82), and GBM (C-index: 0.76, 95% CI: 0.72–0.79) were greater than ST (C-index: 0.72, 95% CI: 0.69–0.76) of train dataset of multivariable group. In the multivariable group, except for ST, the C-index of other models were higher than 0.70. The Cox (C-index: 0.71, 95% CI: 0.67–0.75), RSF (C-index: 0.72, 95% CI: 0.68–0.75), and C-index of GBM (C-index: 0.71, 95% CI: 0.68–0.75) were greater than ST (C-index: 0.69, 95% CI: 0.65–0.73) of train dataset of TNM group. The C-index of Cox, RSF and GBM were greater than ST in every cohort and group.

In the 5-year CSS cohort, the C-index of Cox (C-index: 0.71, 95% CI: 0.68–0.74), RSF (C-index: 0.74, 95% CI: 0.71–0.77), and GBM (C-index: 0.72, 95% CI: 0.69–0.75) were greater than ST (C-index: 0.68, 95% CI: 0.65–0.71) of train dataset of multivariable group. The Cox (C-index: 0.67, 95% CI: 0.64–0.71), RSF (C-index: 0.68, 95% CI: 0.64–0.71), and C-index of GBM (C-index: 0.67, 95% CI: 0.63–0.69) of train dataset of TNM group. All C-index of TNM group were less than 0.70 in 5-year CSS cohort.

Whether in the 3-year CSS cohort or in the 5-year CSS cohort, the C-index of multivariable group was greater than that of the TNM group (Figure 3).

Figure 4 depicted the receiver operating characteristic

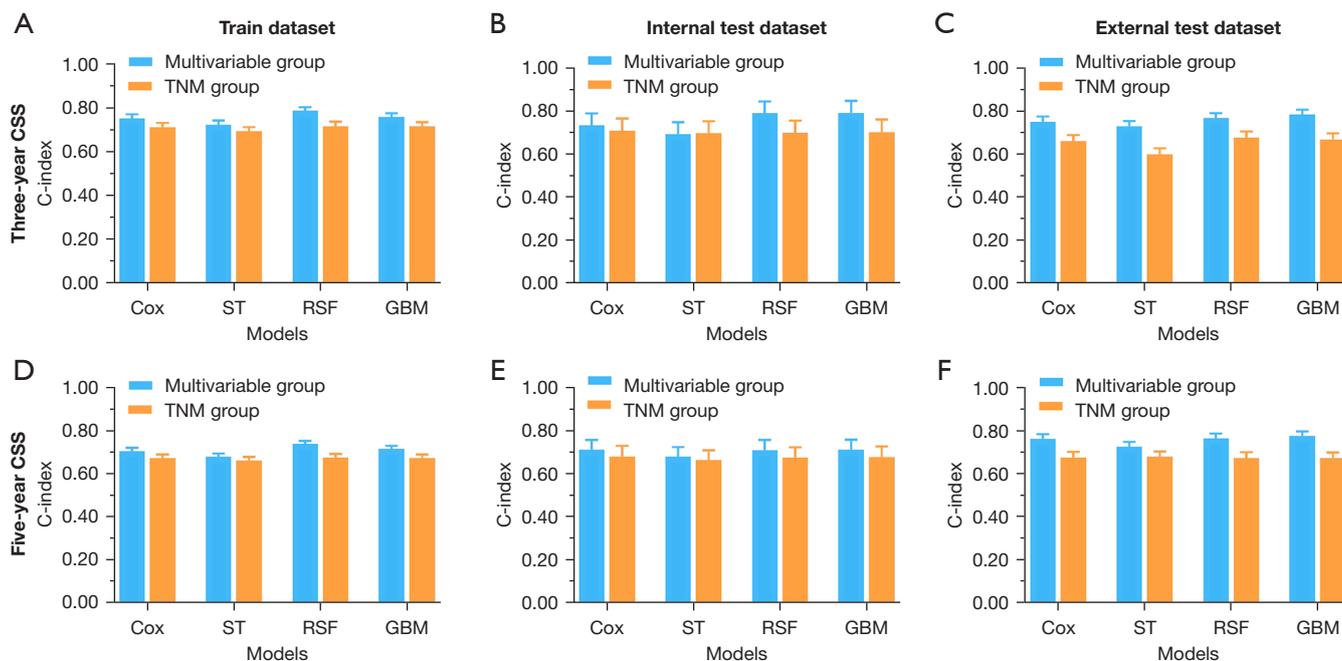


Figure 3 The histograms to compare the C-index of different models of 3- and 5-year CSS in gastric cancer patients. (A) Histogram of C-index in 3-year CSS cohorts of train dataset. (B) Histogram of C-index in 3-year CSS cohorts of internal test dataset. (C) Histogram of C-index in 3-year CSS cohorts of external test dataset. (D) Histogram of C-index in 5-year CSS cohorts of train dataset. (E) Histogram of C-index in 5-year CSS cohorts of internal test dataset. (F) Histogram of C-index in 5-year CSS cohorts of external test dataset. CSS, cancer-specific survival; C-index, the Harrell's consistency index; TNM, Tumor Node Metastasis; ST, survival trees; RSF, random survival forests; GBM, gradient boosting machines.

(ROC) curves of different models for predicting the 3-year CSS of gastric cancer in the multivariable group and TNM group. In the train dataset of multivariable group, the AUC values of Cox, ST, RSF, and GBM were 0.81 (95% CI: 0.79–0.83), 0.78 (95% CI: 0.76–0.80), 0.86 (95% CI: 0.84–0.87), and 0.82 (95% CI: 0.80–0.84), respectively. In the train dataset of TNM group, the AUC values of Cox, ST, RSF, and GBM were 0.76 (95% CI: 0.74–0.79), 0.74 (95% CI: 0.72–0.76), 0.77 (95% CI: 0.75–0.79), and 0.77 (95% CI: 0.74–0.79), respectively. In every dataset or every group, the AUC value of ST was lower than that of other three models. In every dataset, the AUC values of multivariable group were higher than those of the TNM group.

Figure S3 depicted the ROC curves of different models for predicting the 5-year CSS of gastric cancer in the multivariable group and TNM group. In the train dataset of multivariable group, the AUC values of Cox, ST, RSF and GBM are 0.72 (95% CI: 0.70–0.75), 0.70 (95% CI: 0.67–0.72), 0.81 (95% CI: 0.79–0.83) and 0.74 (95% CI: 0.71–0.77), respectively. In the train dataset of TNM group, the AUC values of Cox, ST, RSF, and GBM were 0.69

(95% CI: 0.67–0.72), 0.68 (95% CI: 0.65–0.71), 0.69 (95% CI: 0.66–0.72), and 0.69 (95% CI: 0.67–0.72), respectively. In every dataset or every group, the AUC value of ST was lower than that of other three models. In every dataset, the AUC values of multivariable group were higher than those of the TNM group.

Figure 5 depicted the calibration curves of different models for predicting the 3-year CSS of gastric cancer patients in the multivariable group and TNM group. In multivariable group, the consistency of RSF and Cox were superior than that of GBM and ST; the predicted value of GBM was higher than the actual value, and the predicted value of ST was lower than the actual value. In the TNM group, the performance of the calibration curves for the train dataset and internal test dataset were similar to that in the multivariable group. The consistency of calibration curve of every model in the external test dataset was slightly inferior to that in train dataset and internal test dataset. In the 3-year CSS, the consistency of the calibration curves of all models in the multivariable group were better than that in the TNM group.

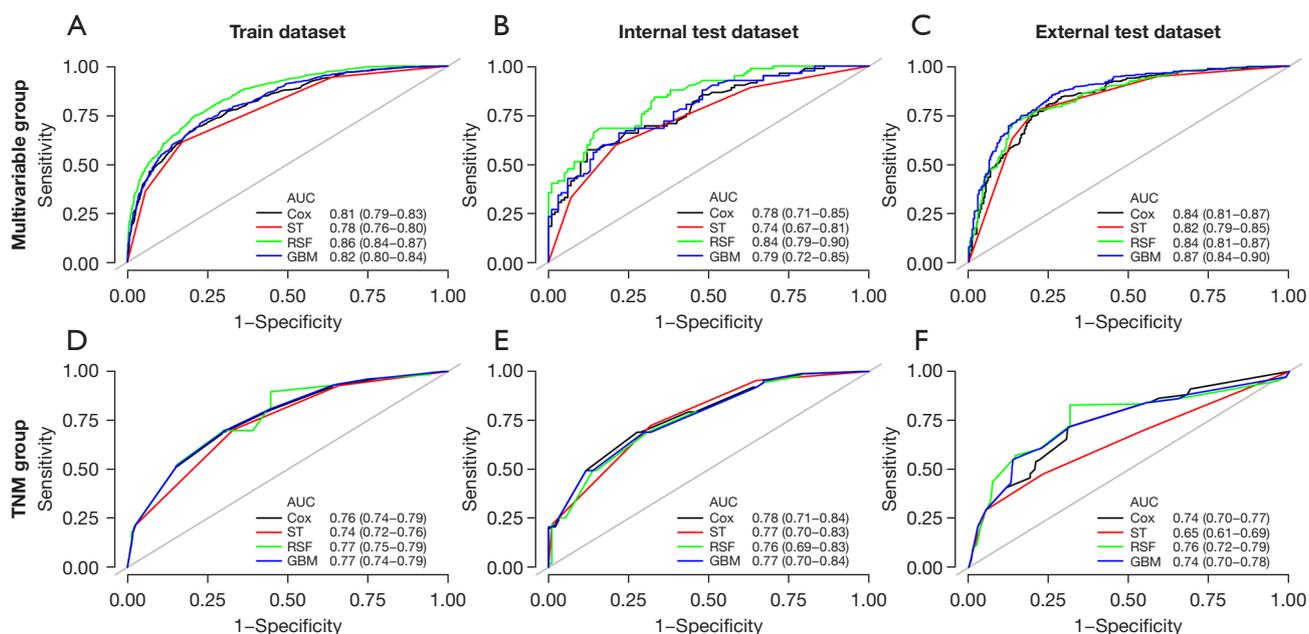


Figure 4 The ROC curve of different models of 3-year CSS in gastric cancer patients. (A) ROC curves in 3-year CSS in multivariable group of train dataset. (B) ROC curves in 3-year CSS in multivariable group of internal test dataset. (C) ROC curves in 3-year CSS in multivariable group of external test dataset. (D) ROC curves in 3-year CSS in TNM group of train dataset. (E) ROC curves in 3-year CSS in TNM group of internal test dataset. (F) ROC curves in 3-year CSS in TNM group of external test dataset. AUC, area under the ROC curve; ROC, receiver operating characteristic; ST, survival trees; RSF, random survival forests; GBM, gradient boosting machines; TNM, Tumor Node Metastasis; CSS, cancer-specific survival.

Figure S4 plotted the calibration curves of different models for predicting the 5-year CSS of gastric cancer patients in the multivariable group and TNM group. The results were similar to those of the 3-year CSS cohort. In the multivariable group, the consistency of Cox and RSF were superior to that of GBM and ST. In the TNM group, the performance of the calibration curves of the train dataset and internal test dataset were similar to that of the multivariable group, with RSF and Cox having better consistency than GBM and ST. The calibration curve consistency of each model in the external test dataset was poor. In the 5-year CSS cohort, the consistency of the calibration curves of all models in the multivariable group was better than that of the TNM group.

Outcome of Cox

From above analysis, we can achieve that the predictive performance of Cox and RSF were superior to that of ST and GBM, and the predictive performance of Cox was similar to that of RSF. Due to the fact that the current

application of RSF wasn't as simple as Cox, in order to better apply the model to practice, we visualized the Cox's results by drawing a forest plot and constructed a nomogram for clinicians to predict patients' survival.

Figure S5 drawn a forest plot of influencing factor of gastric cancer patients' survival based on the Cox of development dataset in multivariable group. The results show that age, tumor site, T stage, N stage, M stage, and surgery situation were the influencing factors for the survival prognosis of gastric cancer patients ($P < 0.05$). Poorer CSS was associated with elder in age (≥ 75 years with hazard ratio (HR) = 1.48, 95% CI: 1.15–1.89, $P = 0.002$) compared to < 55 years; having overlapping tumor sites (HR = 1.68, 95% CI: 1.11–2.55, $P = 0.014$) compared to gastric cardia; T3–T4 stage (HR = 2.91, 95% CI: 2.07–4.09, $P < 0.001$) compared to T1 stage; N2 stage (HR = 1.68, 95% CI: 1.36–2.06, $P < 0.001$) and N3 stage (HR = 2.22, 95% CI: 1.83–2.70, $P < 0.001$) compared to N0 stage; M1 stage (HR = 1.85, 95% CI: 1.53–2.24, $P < 0.001$) compared to M0 stage. Improved CSS was associated with other tumor sites (HR = 0.79, 95% CI: 0.69–0.91, $P = 0.001$) compared to gastric

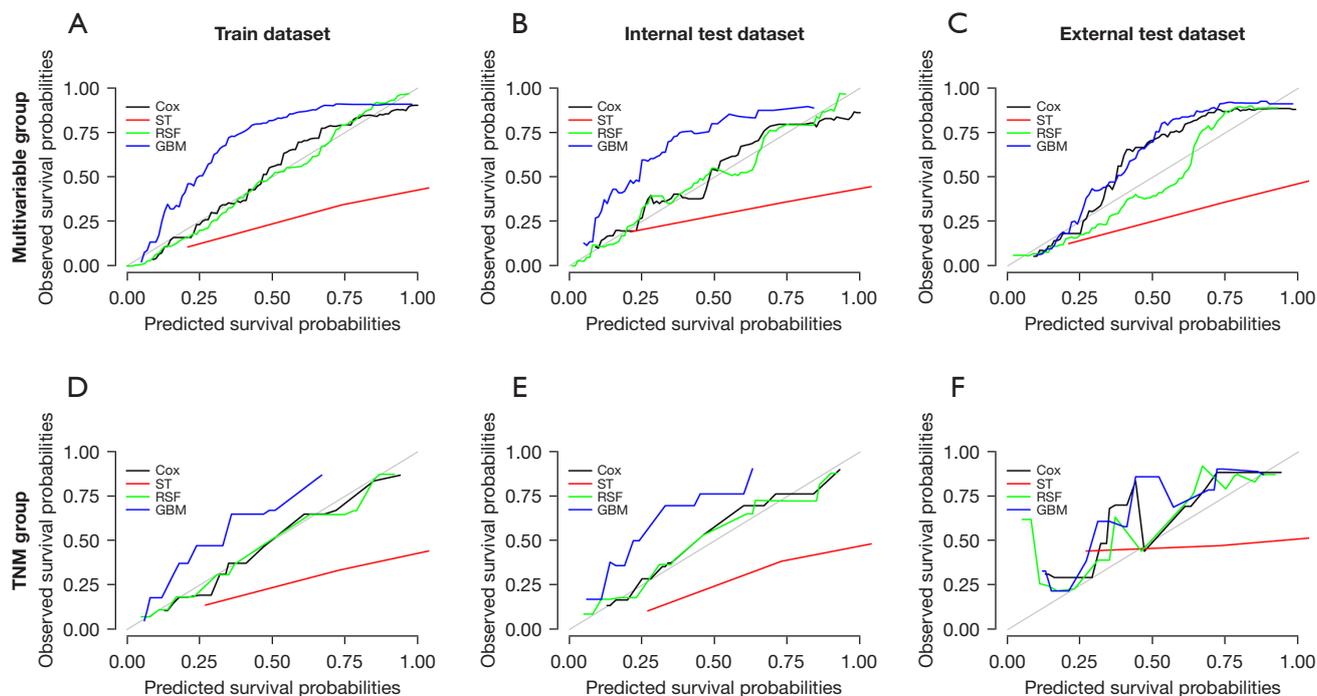


Figure 5 The calibration curve of different models of 3-year CSS in gastric cancer patients. (A) Calibration curves in 3-year CSS in multivariable group of train dataset. (B) Calibration curves in 3-year CSS in multivariable group of internal test dataset. (C) Calibration curves in 3-year CSS in multivariable group of external test dataset. (D) Calibration curves in 3-year CSS in TNM group of train dataset. (E) Calibration curves in 3-year CSS in TNM group of internal test dataset. (F) Calibration curves in 3-year CSS in TNM group of external test dataset. ST, survival trees; RSF, random survival forests; GBM, gradient boosting machines; TNM, Tumor Node Metastasis; CSS, cancer-specific survival.

cardia; and underwent surgery (HR =0.42, 95% CI: 0.35–0.49, $P < 0.001$) compared to didn't undergo surgery.

Figure 6 depicted a nomogram established based on multivariable Cox PH regression model to predict the 3- and 5-year CSS of gastric cancer patients based on development dataset. By bringing the variables of the patient in nomogram, the scores of each variable can be obtained. Finally, the scores of each variable were added up to obtain the 3- and 5-year CSS of the patient. For example, we brought the variables of patient number 36189409 of external test dataset into the nomogram, we obtained the 3- and 5-year CSS were 0.264 (0.150–0.466) and 0.180 (0.087–0.374), respectively.

Sensitivity analysis

To eliminate the bias caused by different variables of different models, we built a same variables group for sensitivity analysis. In the same variables group, the variables included in the ST, RSF, and GBM were consistent with

that in Cox. The variables included gender, age, tumor site, grade, T stage, N stage, M stage, surgery, and radiotherapy. Table 3 and Figures S6,S7 respectively showed the C-indexes, ROC curves and calibration curves of different models that predicted the 3- and 5-year CSS of gastric cancer patients in the same variables group. From the obtained results, it can be seen that the predicting performance of the same variables group were similar to those of the multivariable group.

Table 3 presents the C-indexes of different models for predicting 3- and 5-year CSS of gastric cancer patients in same variables group. In the 3-year CSS cohort, the C-index of Cox (C-index: 0.75, 95% CI: 0.72–0.79), RSF (C-index: 0.77, 95% CI: 0.74–0.80), and GBM (C-index: 0.75, 95% CI: 0.72–0.79) in the train dataset were greater than that of ST (C-index: 0.73, 95% CI: 0.69–0.76). In the 5-year CSS cohort, the C-index of Cox (C-index: 0.71, 95% CI: 0.67–0.74), RSF (C-index: 0.72, 95% CI: 0.70–0.75), and GBM (C-index: 0.71, 95% CI: 0.68–0.74) in the train dataset were greater than that of ST (C-index: 0.68, 95% CI: 0.65–0.71).

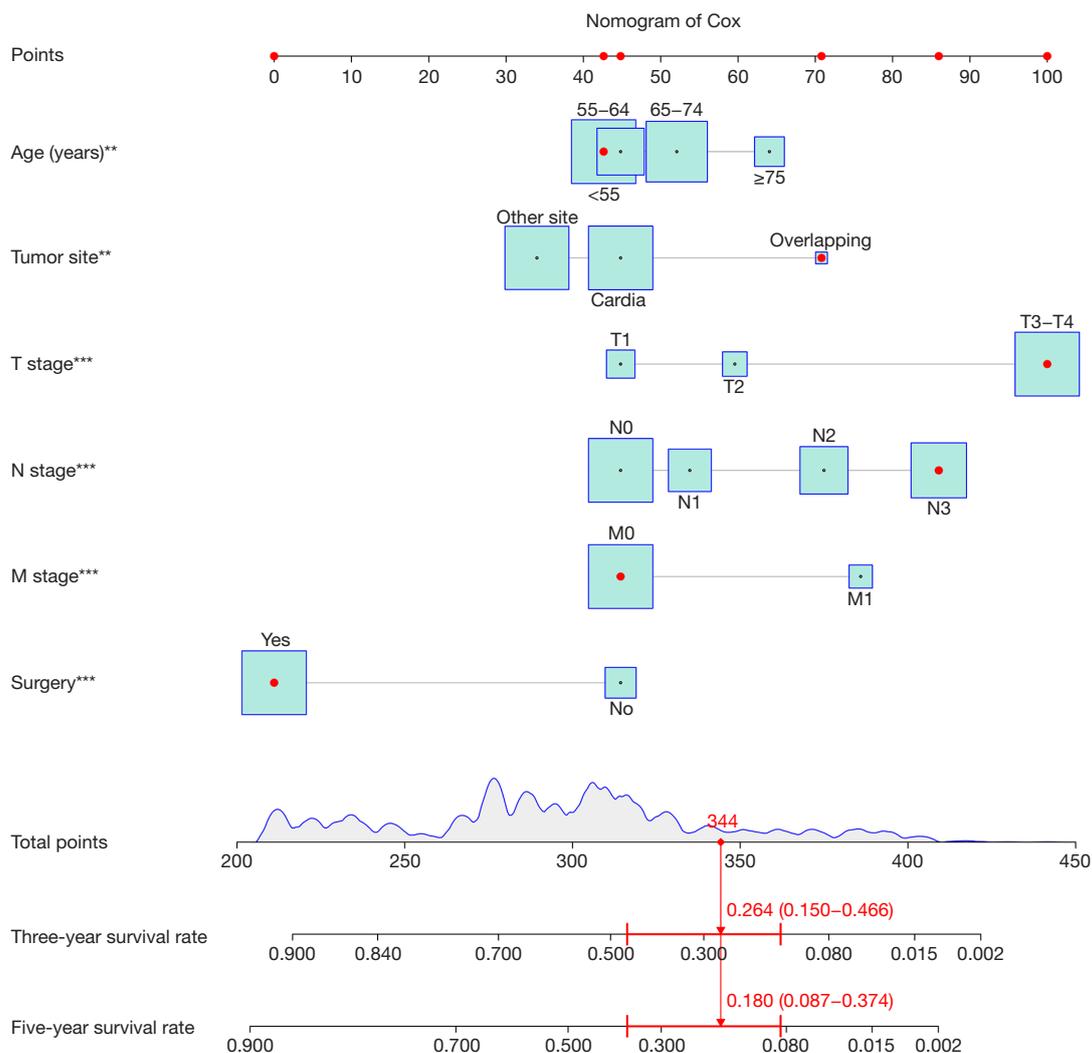


Figure 6 A nomogram for predicting the specific survival rate of gastric cancer patients based on Cox proportional hazard regression model. T stage, tumor stage; N stage, node stage; M stage, metastasis stage. **, P<0.01; ***, P<0.001.

The C-index of Cox, RSF, and GBM were greater than ST in every dataset.

Figure S6 draws the ROC curves of different models that predict the 3- and 5-year CSS of gastric cancer in the same variables group. In the train dataset of the 3-year CSS cohort, the AUC values of Cox, ST, RSF, and GBM were 0.81 (95% CI: 0.79–0.83), 0.78 (95% CI: 0.76–0.80), 0.84 (95% CI: 0.83–0.86), and 0.82 (95% CI: 0.80–0.84), respectively. In the train dataset of the 5-year CSS cohort, the AUC values of Cox, ST, RSF, and GBM were 0.73 (95% CI: 0.70–0.76), 0.70 (95% CI: 0.67–0.73), 0.78 (95% CI: 0.75–0.80), and 0.73 (95% CI: 0.71–0.76), respectively. Among the four models in every dataset, the AUC value of

ST was lower than other three models.

Figure S7 plots the calibration curves of different models for predicting 3- and 5-year CSS of gastric cancer in the same variables group. Whether in 3- or 5-year CSS cohorts, the consistency of RSF and Cox were better than those of GBM and ST. The predicted value of GBM was higher than the actual value, while the predicted value of ST was lower than the actual value.

Discussion

Key findings

This study developed four models (Cox, ST, RSF and

Table 3 The C-index of different models of 3- and 5-year CSS of gastric cancer patients in same variables group

Models	Train dataset, mean (95% CI)	Internal test dataset, mean (95% CI)	External test dataset, mean (95% CI)
Three-year CSS cohort			
Cox	0.75 (0.72–0.79)	0.70 (0.58–0.82)	0.75 (0.70–0.80)
ST	0.73 (0.69–0.76)	0.68 (0.56–0.80)	0.73 (0.69–0.78)
RSF	0.77 (0.74–0.80)	0.76 (0.65–0.86)	0.76 (0.71–0.81)
GBM	0.75 (0.72–0.79)	0.77 (0.67–0.87)	0.77 (0.73–0.82)
Five-year CSS cohort			
Cox	0.71 (0.67–0.74)	0.71 (0.61–0.80)	0.76 (0.72–0.80)
ST	0.68 (0.65–0.71)	0.68 (0.59–0.76)	0.73 (0.68–0.77)
RSF	0.72 (0.70–0.75)	0.70 (0.61–0.80)	0.76 (0.71–0.80)
GBM	0.71 (0.68–0.74)	0.71 (0.62–0.80)	0.76 (0.72–0.81)

CSS, cancer-specific survival; CI, confidence interval; ST, survival trees; RSF, random survival forests; GBM, gradient boosting machines.

GBM) to predict the survival of gastric cancer patients. Whether in 3-year CSS cohort or 5-year CSS cohort, the C-index or AUC of Cox, RSF, and GBM were greater than ST in every dataset. The performance of calibration curves of Cox and RSF were better than those of GBM and ST in both the multivariable group and TNM group. The performance of the calibration curves of all models in the multivariable group were better than that in the TNM group. In addition, we also established a nomogram to predict the 3- and 5-year CSS of gastric cancer patients.

Explanations of findings

The proportion of stage III and IV hospitalized gastric cancer patients in the Hebei Cancer Registration Project (61.41%) was higher than that of Chinese California gastric cancer patients (55.35%). Previous studies have shown that the prognosis of patients with stage III and IV gastric cancer is much worse than that of patients with stage I and II gastric cancer (42,43). The survival rate of hospitalized gastric cancer patients in Hebei Province (1-, 3-, and 5-year CSS were 82.61%, 57.07%, and 44.48%, respectively) were higher than that of Chinese California hospitalized gastric cancer patients (1-, 3-, and 5-year CSS were 68.58%, 47.99%, and 42.91%, respectively), which may be due to Hebei Province was a high-risk area for gastric cancer and had relatively advanced gastric cancer diagnosis and treatment technologies. In addition, it may be related to treatment, tumor biology or follow-up time. Further

exploration is still needed.

The C-index or AUC of Cox, RSF, and GBM were greater than ST in every dataset. The performance of calibration curves of Cox and RSF were better than those of GBM and ST. The results of this study were similar to our previous results of osteosarcoma (44). It indicated that Cox and RSF not only outperform ST and GBM in the survival prognosis of osteosarcoma, but also in the survival prognosis of gastric cancer, which may be extrapolated to other cancers in future studies. ST splitting nodes by maximizing survival differences among nodes using log-rank testing. However, the prediction error is large, resulting in low prediction accuracy (31,32). Both RSF and GBM are combined of a large number of ST. The fundamentals of GBM are training a new ST according to the negative gradient information of the loss function based on the current ST, and combining the trained newborn ST with the existing ST (34). In this study, The C-index and AUC of GBM are similar to that of Cox or RSF. However, the consistency of calibration curve of GBM performs poorer, which means it needs to be improved. RSF uses the bootstrap method to extract sub-samples from the original samples to construct a ST, averaging the cumulative risk function of each ST and ultimately obtaining the total cumulative risk function (33). RSF considers all possible connections among outcome variables and predictors, as well as all possible interactions among variables. Therefore, it approximates the data generation mechanism in observed value, obtains the predicted value that is closest to the actual value (37). RSF may be an alternative non-

parametric method to Cox.

Comparison with similar researches

The survival prediction model constructed based on multiple variables was superior to that based on T stage, N stage, and M stage. This may be due to the fact that the TNM staging system only considers tumor infiltration, lymph node metastasis, and tumor metastasis, without considering biological differences among different patients. This leads to different survival outcomes for patients in the same stage, and same survival outcome for patients in different stages. Therefore, the TNM staging system is not sufficient to predict the prognosis of tumors (45-47). The results of Cox PH regression model showed that age, tumor site, T stage, N stage, M stage, and surgery situation were the influencing factors for the survival prognosis of gastric cancer patients. The survival prognosis of gastric cancer patients aged ≥ 75 years old is significantly worse than those aged < 55 years old, which is similar to previous research results (42). This may be because that compared to young patients, elderly patients have relatively poor physical condition and more basic diseases. They are prone to recurrence after receiving treatment such as surgery or chemotherapy, so the survival prognosis is poor (48-51). The survival prognosis of patients with overlapping sites were significantly worse than those with cardiac sites; the survival prognosis of patients with T3-T4 stage were significantly worse than those with T1 stage; the survival prognosis of patients with N2 stage and N3 stage were significantly worse than those with N0 stage; the survival prognosis of patients with M1 stage were significantly worse than those with M0 stage; the survival prognosis of patients who underwent surgery were better than those who didn't undergo surgery, and these research results were consistent with previous studies (52-56). The results of this study showed that there was no statistical difference in survival prognosis between different genders, which was consistent with previous research results (57,58).

Strengths and limitations

The advantages of this study were as follow: firstly, in this study, we set hospitalized gastric cancer patients of the Hebei Cancer Registration Project as the development dataset to develop models, and used Chinese California hospitalized gastric cancer patients as the external test dataset to external validate models. This external test

dataset can exclude differences between hospital source data and population data, as well as differences among different races. The external test dataset proves that our model is not only suitable for hospitalized gastric cancer patients in Hebei Province, but also for Chinese California hospitalized gastric cancer patients. Such differences between development dataset and external test dataset can serve as a valuable tool for effective external validation. At the same time, all models were established using ten-fold cross validation with 200 iterations, which further increased the reliability of the model. Secondly, in this study, we compared multivariable group with TNM group. In our study, we used a TNM group, composed of three variables (T stage, N stage, and M stage), which were more detailed and had more accurate predictive effects than TNM stage. Thirdly, we set up same variables group (Cox, ST, RSF and GBM include the same variables) for sensitivity analysis to avoid bias caused by different models include different variables. Fourthly, we use the "random forest" multiple imputation method to impute the missing data, which can reduce the information bias caused by excluding samples due to the missing of some variables. Fifthly, we developed a nomogram based on Cox PH regression model to predict the CSS rate of gastric cancer patients. By bringing various variables of the patient in nomogram, we can obtain the 3- and 5-year CSS of the patient. This prognostic tool not only allows doctors and patients to know the patients' probability of survival, but also provides recommendations for clinical doctors' treatment decisions and methods.

This study still has some limitations: firstly, the variables included in this study are limited. Previous studies have shown that some other variables, such as patient nutritional status (42,59), Eastern Cooperative Oncology Group (ECOG) performance (60), radiomics (61), and *Helicobacter pylori* infection (62), may be related to the survival of gastric cancer patients. In future research, we will attempt to incorporate these factors into the model. Secondly, due to technical reasons, we only used three tree-based machine learning methods (ST, RSF, and GBM) to compare with the Cox model.

Implications and actions needed

This study fills in the blank of prognostic models for hospitalized gastric cancer patients in Hebei Province. The proposed nomogram can be used to calculate the 3- and 5-year CSS of gastric cancer patients based on the clinical information. It may be utilized practically to help

clinicians to obtain individualized survival prediction and provide better treatment allocation. In the future, we will try to compare more models [such as SVM, artificial neural networks, XGBoost (eXtreme gradient boosting), etc.] to establish a more comprehensive and excellent prognostic model.

Conclusions

The performance of the multivariable group was superior to TNM group. Cox and RSF have better predictive effects than ST and GBM. The nomogram was useful for facilitating clinicians to predict the survival of gastric cancer patients, and identifying high-risk patients so as to adopt more reasonable treatment plans.

Acknowledgments

We are very grateful to the staff in Hebei Province cancer registries for their kind work in data collection and follow up. Anyone else who contributed to the manuscript but does not qualify for authorship has been acknowledged with their permission.

Funding: None.

Footnote

Reporting Checklist: The authors have completed the TRIPOD reporting checklist. Available at <https://cco.amegroups.com/article/view/10.21037/cco-23-85/rc>

Data Sharing Statement: Available at <https://cco.amegroups.com/article/view/10.21037/cco-23-85/dss>

Peer Review File: Available at <https://cco.amegroups.com/article/view/10.21037/cco-23-85/prf>

Conflicts of Interest: All authors have completed the ICMJE uniform disclosure form (available at <https://cco.amegroups.com/article/view/10.21037/cco-23-85/coif>). The authors have no conflicts of interest to declare.

Ethical Statement: The authors are accountable for all aspects of the work in ensuring that questions related to the accuracy or integrity of any part of the work are appropriately investigated and resolved. The study was conducted in accordance with the Declaration of Helsinki (as revised in 2013). For external test dataset from the SEER

database, the authors obtained authorization to access the SEER Research Data supported by the National Cancer Institute with approval number 11241-Nov2021. Because public and anonymous data from the SEER database were used, informed patient consent was not required. For development dataset from Hebei Cancer Registration Project, the Ethics Committee of the Fourth Hospital of Hebei Medical University/the Tumor Hospital of Hebei Province has confirmed that no ethical approval is required. Individual consent for this retrospective analysis was waived.

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Cite this article as: Hao Y, Li D, Liang D, Liu Y, Zhang S, He Y. Establishment and validation of a prognostic model for gastric cancer patients of Hebei Province in China. *Chin Clin Oncol* 2023;12(6):63. doi: 10.21037/cco-23-85

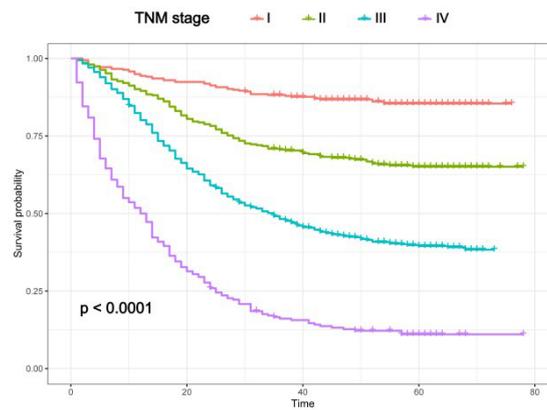


Figure S1 Kaplan-Meier curve of different TNM stages. TNM, Tumor Node Metastasis.

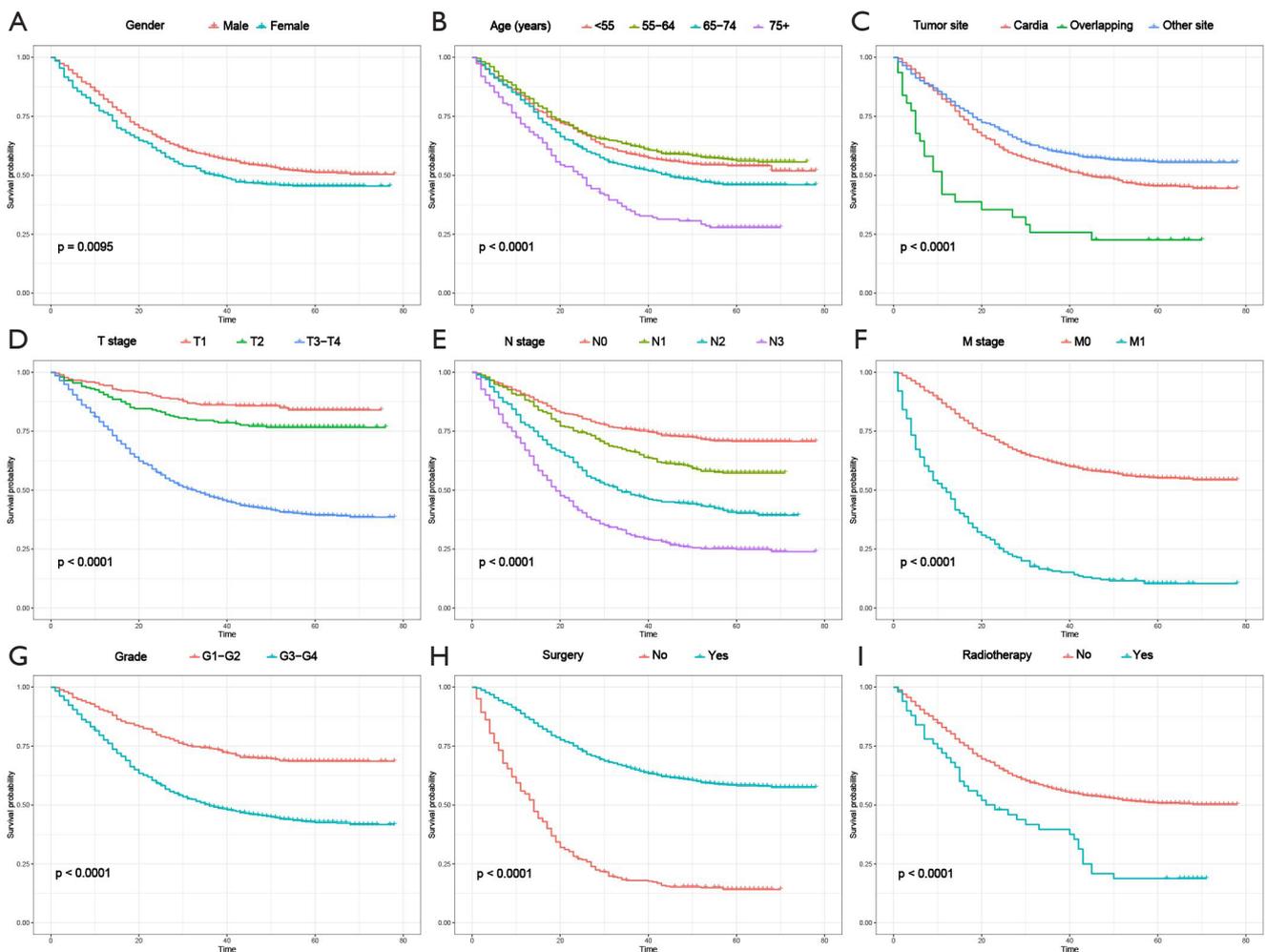


Figure S2 Kaplan-Meier curves of meaningful variables by log-rank test. (A) Kaplan-Meier curves of different genders; (B) Kaplan-Meier curves of different age; (C) Kaplan-Meier curves of different tumor sites; (D) Kaplan-Meier curves of different T stages; (E) Kaplan-Meier curves of different N stages; (F) Kaplan-Meier curves of different M stages; (G) Kaplan-Meier curves of different grades; (H) Kaplan-Meier curves of different surgery situations; (I) Kaplan-Meier curves of different radiotherapy situations. G1–G2: well-differentiated or moderately differentiated; G3–G4: poorly differentiated or undifferentiated. T stage, tumor stage; N stage, node stage; M stage, metastasis stage.

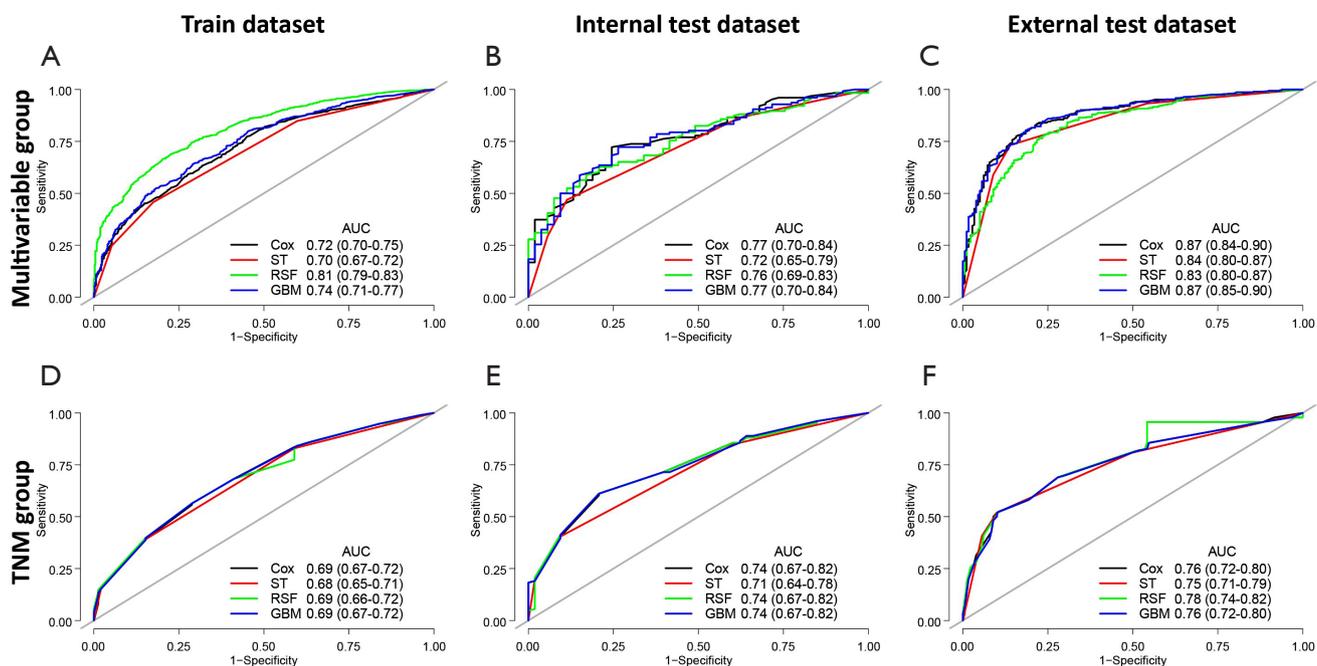


Figure S3 The ROC curve of different models of 5-year CSS in gastric cancer patients; (A) ROC curves in 5-year CSS in multivariable group of train dataset; (B) ROC curves in 5-year CSS in multivariable group of internal test dataset; (C) ROC curves in 5-year CSS in multivariable group of external test dataset; (D) ROC curves in 5-year CSS in TNM group of train dataset; (E) ROC curves in 5-year CSS in TNM group of internal test dataset; (F) ROC curves in 5-year CSS in TNM group of external test dataset. AUC, area under the ROC curve; ROC, receiver operating characteristic; ST, survival trees; RSF, random survival forests; GBM, gradient boosting machines; TNM, Tumor Node Metastasis; CSS, cancer-specific survival.

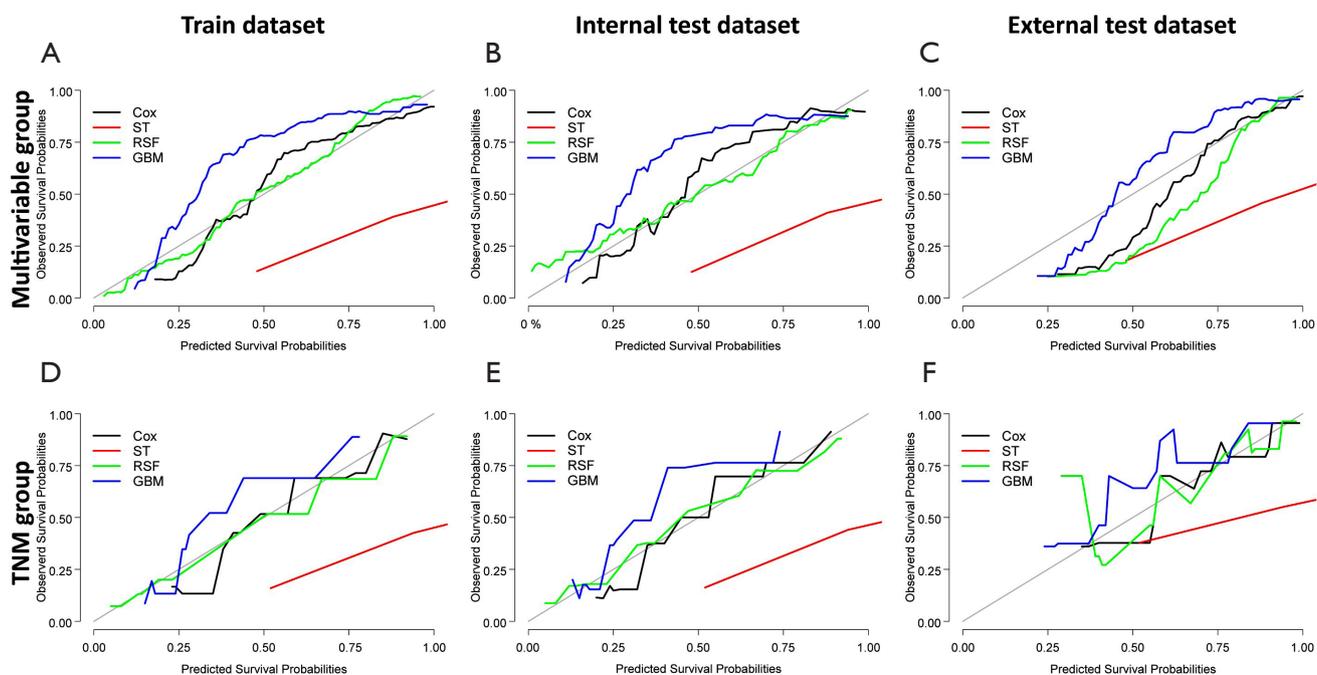


Figure S4 The calibration curve of different models of 5-year CSS in gastric cancer patients; (A) Calibration curves in 5-year CSS in multivariable group of train dataset; (B) calibration curves in 5-year CSS in multivariable group of internal test dataset; (C) calibration curves in 5-year CSS in multivariable group of external test dataset; (D) calibration curves in 5-year CSS in TNM group of train dataset; (E) calibration curves in 5-year CSS in TNM group of internal test dataset; (F) calibration curves in 5-year CSS in TNM group of external test dataset. ST, survival trees; RSF, random survival forests; GBM, gradient boosting machines; TNM, Tumor Node Metastasis; CSS, cancer-specific survival.

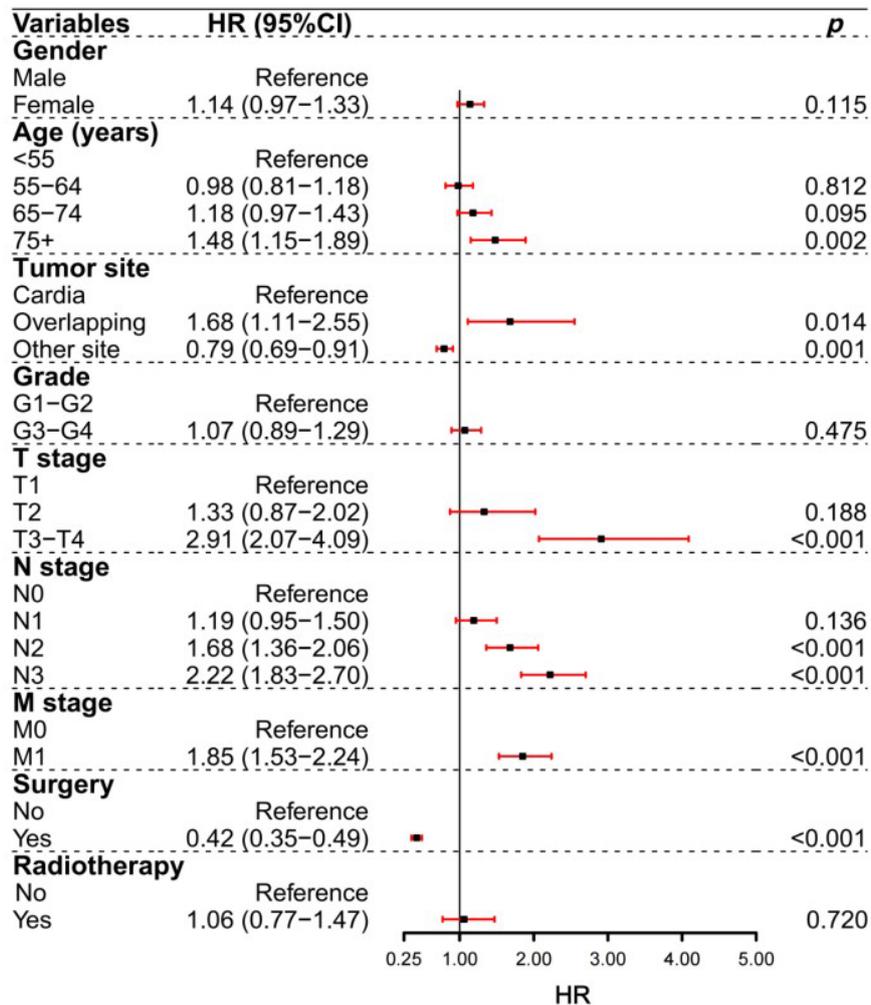


Figure S5 The outcome of multivariate Cox analysis of the specific survival of gastric cancer patients in the hospitals of the Hebei Cancer Registration Project. G1-G2: well-differentiated or moderately differentiated; G3-G4: poorly differentiated or undifferentiated. HR, hazard ratio; CI, confidence interval; T stage, tumor stage; N stage, node stage; M stage, metastasis stage.

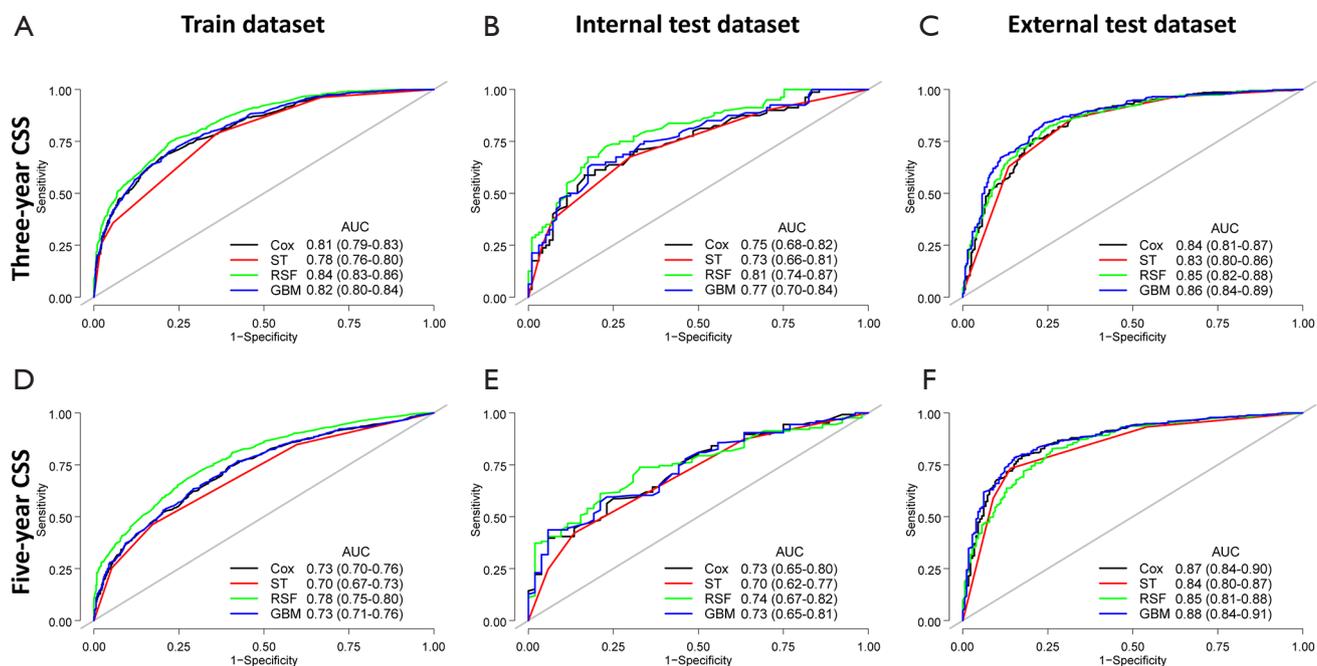


Figure S6 The ROC curve of different models of 3- and 5-year CSS in gastric cancer patients in the same variables group. (A) ROC curves in 3-year CSS in the same variables group of train dataset; (B) ROC curves in 3-year CSS in the same variables group of internal test dataset; (C) ROC curves in 3-year CSS in the same variables group of external test dataset; (D) ROC curves in 5-year CSS in the same variables group of train dataset; (E) ROC curves in 5-year CSS in the same variables group of internal test dataset; (F) ROC curves in 5-year CSS in the same variables group of external test dataset. AUC, area under the ROC curve; ROC, receiver operating characteristic; ST, survival trees; RSF, random survival forests; GBM, gradient boosting machines; TNM, Tumor Node Metastasis; CSS, cancer-specific survival.

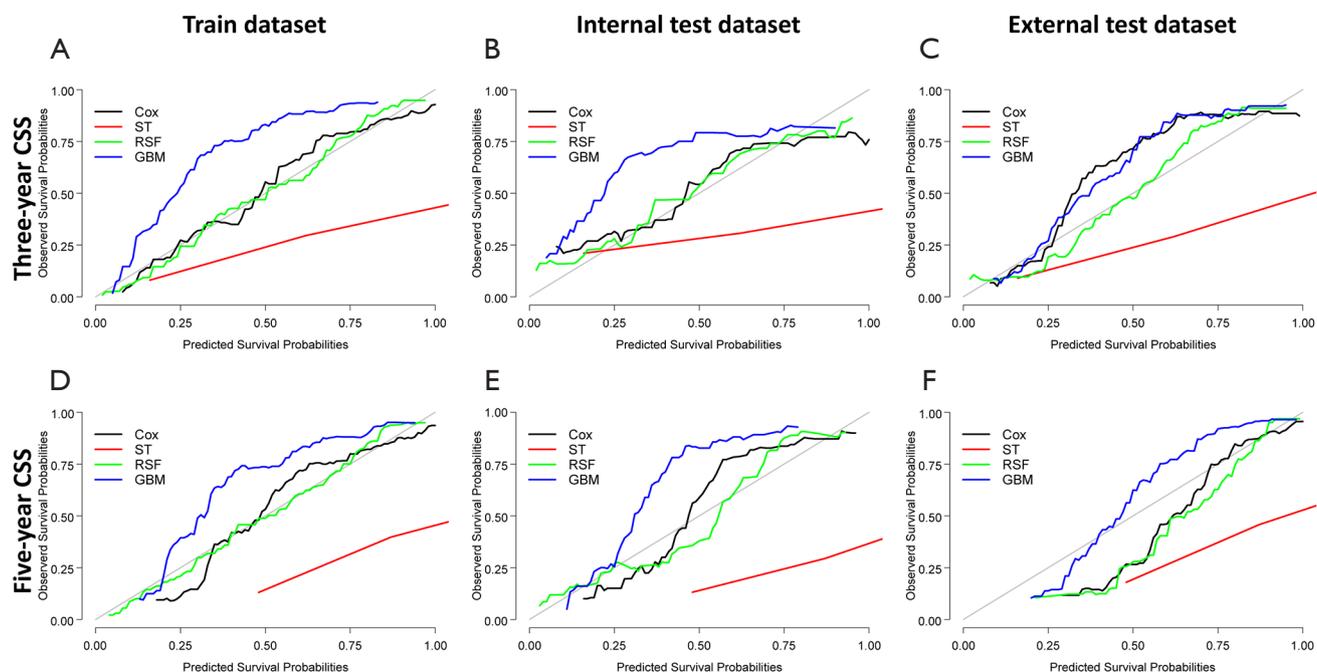


Figure S7 The calibration curve of different models of 3- and 5-year CSS in gastric cancer patients in the same variables group. (A) Calibration curves in 3-year CSS in the same variables group of train dataset; (B) calibration curves in 3-year CSS in the same variables group of internal test dataset; (C) calibration curves in 3-year CSS in the same variables group of external test dataset; (D) calibration curves in 5-year CSS in the same variables group of train dataset; (E) calibration curves in 5-year CSS in the same variables group of internal test dataset; (F) calibration curves in 5-year CSS in the same variables group of external test dataset. ST, survival trees; RSF, random survival forests; GBM, gradient boosting machines; TNM, Tumor Node Metastasis; CSS, cancer-specific survival.