

Establishment and validation of a prediction model for gastric cancer with perineural invasion based on preoperative inflammatory markers

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Background: Gastric cancer (GC) is a prevalent malignant tumor of the digestive system, characterized by a poor prognosis and high recurrence rate. Perineural invasion (PNI), the neoplastic infiltration of nerves, is a significant predictor of survival outcome in GC. Accurate preoperative identification of PNI could facilitate patient stratification and optimal preoperative treatment. We therefore established and validated a preoperative risk assessment model for GC patients with PNI.

Methods: We collected data from 1,195 patients who underwent surgical resection at our hospital between October 2020 and December 2023, with PNI confirmed by pathological examination. We gathered laboratory data, including blood cell count, blood type, coagulation index, biochemical indexes, and tumor markers. Eligible patients were randomly divided into a training set and a testing set at a ratio of 7:3. The important risk factors of PNI were evaluated by random forest package in RStudio. Receiver operating characteristic-area under the curve (ROC-AUC) analysis was used to evaluate the discriminatory ability of the factors for PNI. Univariate and multivariate logistic regression analyses were utilized to verity independent risk factors for patients with PNI, and the logistic regression model and nomogram were constructed based on the results. Calibration curve and decision curve analysis (DCA) were conducted to assess the predictive model. Finally, we verified the prediction equation model using the testing set.

Results: In the training set, 416 GC patients were pathologically diagnosed with PNI. The top 5 important risk factors for PNI were identified as carcinoembryonic antigen (CEA), fibrinogen-to-lymphocyte ratio (FLR), D-dimer, platelet-to-lymphocyte ratio (PLR), and carbohydrate antigen 19-9 (CA19-9), with optimal cut-off values of 3.89 ng/mL, 2.08, 0.24 mg/L, 122.37, and 14.85 U/mL, respectively. Multivariate logistic regression analysis confirmed that CEA, FLR, D-dimer, PLR, CA19-9, and CA72-4 as independent risk factors for PNI (P<0.05). We formulated the following predictive equation: Logit(P) = $-1.211 + 0.695 \times CEA + 0.546 \times FLR + 0.686 \times D$ -dimer + 0.653 × PLR + 0.515 × CA19-9 + 0.518 × CA72-4 (χ^2 =105.675, P<0.001). The model demonstrated an ROC-AUC value of 0.719 [95% confidence interval (CI): 0.681–0.757] in the training set, with a sensitivity of 68.51% and a specificity of 67.60%. The ROC-AUC value was 0.791 (95% CI: 0.750–0.831) in the testing set (sensitivity: 69.57%, specificity: 56.41%). Calibration curve and DCA confirmed that the model has good discrimination and accuracy.

Conclusions: We successfully established and validated a prediction model for GC patients with PNI

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based on hematological indicators, hoping that this model can provide an adjunctive tool for predicting PNI in clinical work.

Keywords: Gastric cancer (GC); perineural invasion (PNI); inflammatory markers; risk factors; prediction model

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Introduction

Gastric cancer (GC) is the fifth most common cancer and the third leading cause of cancer-related death globally (1). In China, it is also a prevalent digestive tract tumor and a major cause of cancer mortality (2). Due to the lack of early symptoms in GC patients, 80–90% of patients are diagnosed with advanced-stage disease, with a 5-year overall survival of no more than 30% (3). Previous studies have confirmed that tumor size, histologic type, age, lymphovascular invasion, and perineural invasion (PNI), distant metastasis are important prognostic factors for GC (4-7).

PNI, first described in the 1800s by Russian and French pathologists, is characterized by cancer cell infiltration into the perineurium or neural fascicles, indicating an early step in local tumor spread (8,9). More and more studies have affirmed that PNI indicates a high invasive ability of cancer cells, and its prognostic value has been shown

Highlight box

Key findings

• We successfully established and validated a prediction model for gastric cancer (GC) patients with perineural invasion (PNI) based on hematological indicators.

What is known and what is new?

- Previous prediction models have focused on the predictive value of common tumor marker in the occurrence of PNI in GC, and have been tested on limited sample sizes.
- In this study, we established a prediction model for PNI in GC based on peripheral blood inflammatory markers by multivariate logistic regression analysis.

What is the implication, and what should change now?

 When constructing a prognostic model, it is crucial to consider the patients' treatment mode, as this will enhance the accuracy, reliability, specificity, and sensitivity of the model. Our model can provide valuable insights for evaluating the risk of PNI in GC patients. This approach may serve as an adjunctive tool in clinical decision-making processes regarding PNI. in various tumors, including colon cancer, squamous cell carcinoma of the tongue, pancreatic cancer, and prostate cancer (10-13). In GC, PNI is a common and significant independent prognostic factor affecting tumor recurrence and patient survival post-resection (14,15). The presence of PNI correlates significantly with tumor size, tumor, node, metastasis (TNM) stage, histological type, blood vessel invasion, and lymph node metastasis (14,16).

At present, paraffin-embedded tissue sections and histopathology of surgical specimens are used to evaluate PNI. The invasiveness of biopsy and the efficiency and timeliness PNI detection status may limit clinical decisionmaking (17). Preoperative identification of PNI may facilitate patient stratification and formulation of optimal treatment options. Several studies have documented the predictive value of medical images as an adjunctive diagnostic instrument in identifying PNI in GC (17-19). Consequently, considering that laboratory indicators also have the advantages of high cost-effectiveness and easy operation as ancillary assessments, their potential application in the predicting PNI in GC should not be disregarded. In recent years, some laboratory indicators have been confirmed to be significantly related to GC patients with PNI. For instance, carcinoembryonic antigen (CEA) $\geq 5 \mu g/L$ [odds ratio (OR) =5.870, 95% confidence interval (CI): 3.281-10.502] was shown to be a significant independent risk factor of PNI in advanced GC, with an area under the curve (AUC) value of 0.731 in a nomogram model (20). Immunoglobulin A (IgA) level (OR=1.61, 95% CI: 1.03-2.61) was found in the multivariable regression model to be one of the independent predictors of PNI in 261 GC patients (21). In addition, carbohydrate antigen 125 (CA125) was also shown to effectively indicate PNI positivity in gastric adenocarcinoma samples of 135 GC patients, with an AUC value of 0.714 (95% CI: 0.625-0.804) (22). Nevertheless, the above studies almost all reported the predictive value of common tumor markers in the occurrence of PNI in GC, and the limited sample size. Nowadays, accumulating

studies have confirmed that systemic inflammatory response is associated with the tumor microenvironment (TME), serving a pivotal role in tumor progression (23). In the TME, various inflammatory cells such as lymphocytes, macrophages, monocytes, and neutrophils are present (24). During the onset of inflammation, conventional blood indicators such as neutrophils, lymphocytes, and platelets count often become abnormal (25,26). Studies have demonstrated that cellular inflammatory markers, including the neutrophil-to-lymphocyte ratio (NLR), platelet-tolymphocyte ratio (PLR), and lymphocyte-to-monocyte ratio (LMR) are dysregulated in a variety of tumors and are often considered predictors of poor prognosis or malignant progression (27-29). Besides, fibrinogen may be involved in tumor-specific inflammatory response and is significantly elevated during an infection or inflammatory disease. Interestingly, malignant tumor cells can partially express fibrinogen (30,31). A recent study revealed that the systemic inflammation response index (SIRI = neutrophil count × monocyte count/lymphocyte count) is an independent risk factors for PNI in advanced GC, with an AUC value of 0.707 (95% CI: 0.653-0.761), sensitivity of 53.60%, and specificity of 87.9% (32). Despite this, the correlation between inflammatory markers in peripheral blood and PNI in GC is still unclear.

Therefore, this study was undertaken to establish a prediction model for PNI in GC based on peripheral blood inflammatory markers by multivariate logistic regression analysis. In addition, we assessed the prediction model using receiver operating characteristic-AUC (ROC-AUC), calibration curve analysis, and decision curve analysis (DCA). We present this article in accordance with the TRIPOD reporting checklist (available at https://tcr.amegroups.com/article/view/10.21037/tcr-24-481/rc).

Methods

Study patients and processing

The study was conducted in accordance with the Declaration of Helsinki (as revised in 2013) and approved by the Clinical Research Ethics Committee of Jiangsu Cancer Hospital (approval number: #KY-2024-012). The requirement for informed consent was waived in this retrospective study. A total of 1,195 patients with GC from the hospital were enrolled between October 2020 and December 2023. The diagnosis was based on the tissue sample obtained during gastroscopy and confirmed by

postoperative pathology. Patients were staged according to the American Joint Committee on Cancer (AJCC)/Union for International Cancer Control (UICC) 8th edition staging system. The exclusion criteria for patients were as follows: (I) receipt of prior treatment before surgery; (II) history of other malignancies or hematological disorders; (III) distant metastasis; (IV) recurrent GC; (V) history of blood transfusion or related conditions; (VI) disorders or dysfunction of the heart, liver, or kidneys; (VII) incomplete collection of laboratory data. After applying these criteria, 1,004 GC patients were included in the study. All cases were randomly divided into a training set (modeling group) and a testing set (validation group) at a ratio of 7:3. Among them, 703 patients were assigned to the modeling group for building predictive models, and 301 patients were assigned to the validation group for internal validation. There was no statistically significant difference in clinical data between the two groups mentioned above (Table S1).

The research procedures of this study were as follows: firstly, we obtained laboratory data and pathological characteristics in GC patients treated with curative gastrectomy. Secondly, we analyzed the differences in laboratory indicators between PNI and PNI-negative groups to identify significant risk factors for PNI. Then, the optimal cut-off value for PNI was determined using ROC-AUC analysis. Finally, we formulated the prediction equation model and nomogram based on multivariate logistic regression results and assessed the model using ROC-AUC, calibration curve analysis, and DCA. Additionally, we verified the prediction model using the testing set. A flow chart of the study is provided in *Figure 1*.

Laboratory methods

The fasting venous blood sample was collected from all patients in the morning. All laboratory indicators were checked before the cases received surgery treatment. Sysmex XE-2100 hematology analyzer and CS-5100 Hemagglutination analyzer (Sysmex, Kobe, Hyogo, Japan) were used to detect blood cell count and hemagglutination index. An Erytra automatic blood analyzer (Grifols, Barcelona, Spain) was used to detect patient blood type. The cobas[®] 8000 modular analyzer (Roche Diagnostics, Indianapolis, IN, USA) was used to detect the concentration of albumin and lactate dehydrogenase (LDH). The calculation formula was as follows: NLR = neutrophil-tolymphocyte ratio, PLR = platelet-to-lymphocyte ratio, FLR = fibrinogen-to-lymphocyte ratio, LMR = lymphocyte



Figure 1 The flowchart of the study design and analysis. GC, gastric cancer; PNI, perineural invasion; ROC-AUC, receiver operating characteristic-area under the curve; DCA, decision curve analysis.

to-monocyte ratio, ALR = albumin-to-lymphocyte ratio. Then, the concentration of serum tumor markers [CEA, CA125, carbohydrate antigen 72-4 (CA72-4), carbohydrate antigen 19-9 (CA19-9), alpha-fetoprotein (AFP)] was measured with electrochemiluminescence assays in a Roche E601 Immunoassay Analyzer according to the manufacturer's protocols.

PNI detection

The criteria of PNI were cancer cells inside the perineurium

involving at least 33% of the circumference (33). Nerve structure was confirmed by hematoxylin and eosin staining. For each specimen, three independent representative tissue sections were prepared, and when PNI was observed in any of the three sections, the specimens were classified as PNI.

Statistical analysis

The software SPSS 20.0 (IBM Corp., Armonk, NY, USA) and GraphPad Prism v9.4.1 (GraphPad Software, San Diego, CA, USA) were employed for statistical analysis.



Figure 2 Representative micrographs of PNI by tumors at 200× magnification. Representative micrographs of surrounding the nerve sheath (A) and invading through the nerve sheath (B) with HE staining. The yellow arrow indicates PNI. PNI, perineural invasion; HE, hematoxylin and eosin.

The continuous variables in this study were judged by Kolmogorov-Smirnov test to be non-normally distributed. They were expressed as medians with 25th and 75th percentiles, and results were compared using Mann-Whitney U test. Chi-square test was used to compare categorical variables. The important risk factors of PNI were evaluated by random forest package in RStudio (RStudio, Boston, MA, USA). The ROC-AUCs were used to evaluate the discriminatory ability of the factors for PNI. The sensitivity-specificity relationship was determined using Youden's index. Univariate and multivariate logistic regression analysis identified independent risk factors for PNI, with OR and 95% CI calculated. The prediction model was constructed from the multivariate logistic regression results, and its discriminatory ability was determined by ROC-AUC. The consistency between predicted and actual results was assessed through calibration curve analysis by the rms [6.4.0] software package (https:// cran.r-project.org/web/packages/rms/). DCA was performed to assess the clinical utility of the model by the rmda [1.6] software package of R (4.2.1; https://cran. r-project.org/web/packages/rmda/). Subsequently, we verified the prediction model using the testing set. A twosided P value of less than 0.05 was considered statistically significant.

Results

Patient characteristics in modeling group

In the modeling group, 416 GC patients (59.14%) were pathologically diagnosed with PNI; a typical micrograph of PNI is displayed in *Figure 2*. As shown in the *Table 1*, the medians of age and preoperative D-dimer, NLR, PLR, FLR, LMR, ALR, CEA, CA72-4, CA19-9, and CA125,

and the difference between the two groups were statistically significant (all P<0.01). In contrast, gender (P=0.49), blood type (P=0.66), LDH (P=0.70), and AFP (P=0.67) showed no statistically significant differences between the two groups.

Random forest algorithm to extract important ranking for PNI in GC patients

Random forest analysis showed that variables were the most important ranking influencing the occurrence of PNI. As depicted in the *Figure 3*, variables with statistically significant differences between PNI and PNI-negative were analyzed using a random forest package in RStudio. The variable with the highest relative importance was CEA, followed by FLR, D-dimer, PLR, CA19-9, CA72-4, CA125, LMR, ALR, NLR, and age.

ROC curve of important risk factors for PNI in GC patients

In order to evaluate the discriminative power of CEA, FLR, D-dimer, PLR, CA19-9, CA72-4, CA125, LMR, ALR, NLR, and age between PNI and PNI-negative patients, the ROC-AUC was performed. As shown in *Table 2*, the best cut-off values of CEA, FLR, D-dimer, PLR, CA19-9, and CA72-4 for PNI prediction using the Youden's index were 3.89 ng/mL, 2.08, 0.24 mg/L, 122.37, 14.85 U/mL, and 2.31 U/mL, respectively. The ROC-AUC of CEA diagnosis for PNI was 0.581 (95% CI: 0.539–0.623), corresponding to a sensitivity of 30.84% and a specificity of 86.07%. The ROC-AUC of FLR was 0.640 (95% CI: 0.599–0.681), with a sensitivity of 52.16% and a specificity of 71.43%. The ROC-AUC of D-dimer was 0.614 (95% CI: 0.572–0.656) with a sensitivity of 80.24% and specificity of 39.03%.

Table 1 Patient characteristics according to status of perineural invasion in modeling group

Characteristics	PNI-negative (n=287)	PNI-positive (n=416)	χ²/Z	P value
Age (years)	63 [55–69]	66 [58–71]	-3.003	0.003
Gender			0.569	0.49
Women	74 (25.78)	118 (28.37)		
Men	213 (74.22)	298 (71.63)		
Blood type			0.195	0.66
А	92 (32.06)	143 (34.38)		
AB	36 (12.54)	33 (7.93)		
В	78 (27.18)	105 (25.24)		
0	81 (28.22)	135 (32.45)		
D-dimer (mg/L)	0.31 [0.19–0.53]	0.40 [0.26–0.77]	-5.182	<0.001
NLR	1.85 [1.47–2.50]	2.14 [1.62–2.92]	-4.032	<0.001
PLR	117.33 [89.94–160.57]	141.32 [110.19–190.77]	-5.779	<0.001
FLR	1.68 [1.28–2.19]	2.12 [1.59–2.85]	-6.349	<0.001
LMR	3.96 [3.01–5.16]	3.60 [2.74–4.60]	-3.376	<0.001
ALR	25.14 [20.98–31.93]	27.91 [22.07–34.37]	-2.825	0.005
LDH (mmol/L)	176.00 [156.00–199.00]	175.00 [154.00–199.00]	-0.393	0.70
CEA (ng/mL)	1.97 [1.34–3.16]	2.46 [1.41–4.53]	-3.654	<0.001
CA72-4 (U/mL)	1.76 [1.50–3.43]	2.34 [1.50–5.79]	-4.177	<0.001
CA19-9 (U/mL)	8.08 [5.42–14.20]	10.55 [6.03–20.53]	-3.690	<0.001
AFP (ng/mL)	2.45 [1.77–3.40]	2.39 [1.76–3.33]	-0.421	0.67
CA125 (U/mL)	8.79 [6.26–12.70]	9.97 [7.23–15.18]	-3.471	<0.001

The measurement data were expressed as the median and quartile [25–75%], and the enumeration data were expressed as frequency and rate (%). PNI, perineural invasion; NLR, neutrophil-to-lymphocyte ratio; PLR, platelet-to-lymphocyte ratio; FLR, fibrinogen-to-lymphocyte ratio; LMR, lymphocyte-to-monocyte ratio; ALR, albumin-to-lymphocyte ratio; LDH, lactic dehydrogenase; CEA, carcinoembryonic antigen; CA72-4, carbohydrate antigen 72-4; CA19-9, carbohydrate antigen 19-9; AFP, alpha-fetoprotein; CA125, carbohydrate antigen 125.

The ROC-AUC of PLR was 0.627 (95% CI: 0.586–0.669, sensitivity: 66.11%, specificity: 55.41%). The ROC-AUC of CA19-9 was 0.581 (95% CI: 0.539–0.624, sensitivity: 37.11%, specificity: 77.36%).

Univariate logistic regression analysis for PNI in GC patients

Using the occurrence of PNI as the dependent variable (negative =0, positive =1), CEA (≤3.89 ng/mL =0, >3.89 ng/mL =1), FLR (≤2.08 =0, >2.08 =1), D-dimer (≤0.24 mg/L =0, >0.24 mg/L =1), PLR (≤122.37 =0, >122.37=1), CA19-9 (≤14.85 U/mL =0, >14.85 U/mL =1),

CA72-4 (≤ 2.31 U/mL =0, >2.31 U/mL =1), CA125 (≤ 14.45 ng/mL =0, >14.45 ng/mL =1), LMR (≤ 3.65 =0, >3.65 =1), ALR (≤ 25.72 =0, >25.72 =1), and NLR (≤ 2.05 =0, >2.05 =1), age (≤ 63 years =0, >63 years =1), as independent variables. The results showed that the above indicators were related to occurrence of PNI (*Table 3*).

Establishment a predictive risk prediction model for PNI in GC

To establish an accurate method for evaluating the possibility of PNI by using the above hematological indicators, we obtained a classification discriminant

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equation using the multivariate logistic regression analysis to ascertain whether GC patients have PNI as follows: Logit(P)= $-1.211 + 0.695 \times X1 + 0.546 \times X2 + 0.686 \times X3 +$ $0.653 \times X4 + 0.515 \times X5 + 0.518 \times X6 (\chi^2=105.675, P<0.001;$ where X1 = CEA, X2 = FLR, X3 = D-dimer, X4 = PLR,



Figure 3 Random forest algorithm to extract important ranking for PNI. CEA, carcinoembryonic antigen; FLR, fibrinogen-tolymphocyte ratio; PLR, platelet-to-lymphocyte ratio; CA19-9, carbohydrate antigen 19-9; CA72-4, carbohydrate antigen 72-4; CA125, carbohydrate antigen 125; LMR, lymphocyte-to-monocyte ratio; ALR, albumin-to-lymphocyte ratio; NLR, neutrophil-tolymphocyte ratio; PNI, perineural invasion.

X5 = CA19-9, X6 = CA72-4, *Table 4*), for which the critical value is 0.50, thus, if the Logit (P) of a case is larger than 0.50, it belongs to the PNI group. On the contrary, if the Logit (P) \leq 0.50, it belongs to the PNI-negative groups. Based on the logistic regression model, we established a nomogram prediction model for PNI in GC patients (*Figure 4A*).

Immediately after, based on the predicted probability of PNI in GC patients using this model, the ROC-AUC was 0.719 (95% CI: 0.681–0.757, P<0.01), with a sensitivity of 68.51% and a specificity of 67.60% (*Figure 4B*). The calibration ability of the prediction model was evaluated through the Hosmer-Lemeshow goodness of fit test. The results indicated that there were no statistically significant differences between the predicted values of the model and the actual observed values (χ^2 =13.632, P=0.09) (*Figure 4C*). Besides, the DCA suggested that the clinical net benefit of intervention based on the predicted probability of the model is highest when the threshold probability is between 0.36 and 0.83 (*Figure 4D*).

Validation of the prediction model in testing set

Subsequently, we proceeded to validate the discriminant equation model using the testing set data, comprising 184 patients with PNI and 117 PNI-negative patients. There were 51 patients with GC mistaken for PNI and 56

Table 2 Relevant results of important risk factors for PNI patients with GC in modeling group

Factors	AUC (95% CI)	SE	P value	Sensitivity (%)	Specificity (%)	Cut-off value	Youden's index
CEA (ng/mL)	0.581 (0.539–0.623)	0.021	<0.001	30.84	86.07	3.89	0.169
FLR	0.640 (0.599–0.681)	0.021	<0.001	52.16	71.43	2.08	0.236
D-dimer (mg/L)	0.614 (0.572–0.656)	0.022	<0.001	80.24	39.03	0.24	0.193
PLR	0.627 (0.586–0.669)	0.021	<0.001	66.11	55.41	122.37	0.215
CA19-9 (U/mL)	0.581 (0.539–0.624)	0.021	<0.001	37.11	77.36	14.85	0.145
CA72-4 (U/mL)	0.590 (0.548–0.633)	0.022	<0.001	50.84	64.46	2.31	0.153
CA125 (U/mL)	0.577 (0.535–0.620)	0.022	<0.001	28.92	82.93	14.45	0.118
LMR	0.575 (0.532–0.618)	0.022	0.001	60.63	51.45	3.65	0.121
ALR	0.562 (0.519–0.605)	0.022	0.005	59.62	56.45	25.72	0.161
NLR	0.590 (0.547–0.632)	0.022	<0.001	56.01	58.89	2.05	0.149
Age (years)	0.566 (0.523–0.609)	0.022	0.003	61.93	49.48	63.00	0.114

PNI, perineural invasion; GC, gastric cancer; CEA, carcinoembryonic antigen; FLR, fibrinogen-to-lymphocyte ratio; PLR, platelet-to-lymphocyte ratio; CA19-9, carbohydrate antigen 19-9; CA72-4, carbohydrate antigen 72-4; CA125, carbohydrate antigen 125; LMR, lymphocyte-to-monocyte ratio; ALR, albumin-to-lymphocyte ratio; NLR, neutrophil-to-lymphocyte ratio; AUC, area under the curve; SE, standard error; CI, confidence interval.

Factors	В	SE	Wald	P value	OR	95% CI
CEA >3.89 ng/mL	1.010	0.201	25.271	<0.001	2.744	1.851-4.068
FLR >2.08	0.992	0.163	36.946	<0.001	2.700	1.960–3.719
D-dimer >0.24 mg/L	0.899	0.170	28.044	<0.001	2.457	1.761–3.426
PLR >122.37	0.885	0.158	31.535	<0.001	2.423	1.779–3.299
CA19-9 >14.85 U/mL	0.697	0.174	16.082	<0.001	2.008	1.428-2.822
CA72-4 >2.31 U/mL	0.624	0.158	15.695	<0.001	1.867	1.371–2.542
CA125 >14.45 ng/mL	0.655	0.192	11.607	0.001	1.925	1.321-2.806
LMR >3.65	-0.480	0.156	9.504	0.002	0.619	0.456-0.840
ALR >25.72	0.649	0.155	17.422	<0.001	1.913	1.411–2.594
NLR >2.05	0.567	0.155	13.299	<0.001	1.762	1.300–2.389
Age >63 years	0.458	0.155	8.761	0.003	1.581	1.167–2.142

Table 3 Univariate logistic regression analysis of PNI in patients with GC in modeling group

PNI, perineural invasion; GC, gastric cancer; CEA, carcinoembryonic antigen; FLR, fibrinogen-to-lymphocyte ratio; PLR, platelet-to-lymphocyte ratio; CA19-9, carbohydrate antigen 19-9; CA72-4, carbohydrate antigen 72-4; CA125, carbohydrate antigen 125; LMR, lymphocyte-to-monocyte ratio; ALR, albumin-to-lymphocyte ratio; NLR, neutrophil-to-lymphocyte ratio; B, Beta coefficient; SE, standard error; OR, odds ratio; CI, confidence interval.

 Table 4 Multivariate logistic regression analysis of PNI in patients with GC in modeling group

Factors	В	SE	Wald	P value	OR	95% CI
CEA (X1)	0.695	0.216	10.405	0.001	2.004	1.314–3.507
FLR (X2)	0.546	0.186	8.620	0.003	1.726	1.199–2.484
D-dimer (X3)	0.686	0.182	14.259	<0.001	1.986	1.391–2.835
PLR (X4)	0.653	0.179	13.337	<0.001	1.922	1.353–2.729
CA19-9 (X5)	0.515	0.189	7.452	0.006	1.674	1.156-2.424
CA72-4 (X6)	0.518	0.168	9.462	0.002	1.678	1.207–2.333
Constant	-1.211	0.197	37.895	<0.001	0.298	-

Method: forward LR. PNI, perineural invasion; GC, gastric cancer; CEA, carcinoembryonic antigen; FLR, fibrinogen-to-lymphocyte ratio; PLR, platelet-to-lymphocyte ratio; CA19-9, carbohydrate antigen 19-9; CA72-4, carbohydrate antigen 72-4; B, Beta coefficient; SE, standard error; OR, odds ratio; CI, confidence interval; LR, likelihood ratio.

PNI patients whose diagnoses were missed by the equation results (*Table 5*). The diagnostic efficiency for PNI in testing set was calculated by Clinical Calculator 1 (http:// vassarstats.net/clin1.html). The sensitivity, specificity, positive predictive value (PPV), and negative predictive value (NPV) were 69.57%, 56.41%, 71.51%, and 54.10%, respectively (*Table 6*). As shown in *Figure 5A*, the ROC-AUC of the model was 0.791 (95% CI: 0.750–0.831, P<0.01). Further, the calibration curve showed that the model had good discrimination and accuracy (*Figure 5B*).

The DCA indicated that when the threshold probability is between 0.27 and 0.87, the clinical net benefit is highest (*Figure 5C*).

Discussion

PNI is characterized by an encounter between the cancer cells and neuronal fibers and holds an extremely poor prognosis in malignant tumors (34). A meta-analysis reported that the median of PNI in GC is 40.9% (range,



Figure 4 Establishment of a predictive risk prediction model for PNI in GC. (A) Nomogram prediction model for PNI. (B) ROC curve of model in the modeling group. (C) Calibration curve of model in the modeling group. (D) DCA of model in the modeling group. CEA, carcinoembryonic antigen; FLR, fibrinogen-to-lymphocyte ratio; PLR, platelet-to-lymphocyte ratio; CA19-9, carbohydrate antigen 19-9; CA72-4, carbohydrate antigen 72-4; AUC, area under curve; CI, confidence interval; PNI, perineural invasion; GC, gastric cancer; ROC, receiver operating characteristic; DCA, decision curve analysis.

Table 5 The discriminant value of validation group						
Results of criterion	Pathologic	Total				
equation	PNI-positive PNI-negative		iotai			
PNI-positive	128	51	179			
PNI-negative	56	66	122			
Total	184	117	301			

PNI, perineural invasion.

6.8–75.6%) (35). In our modeling group, PNI was pathologically diagnosed in 416 patients, representing 59.14% of the group. Paraffin-embedded tissue section of specimens and histopathology were used to evaluate PNI.

Nevertheless, the positivity of PNI might be affected by the number of tissues obtained, biopsy techniques, tissue sections, and inter-observer differences. Additionally, some patients with PNI may not be surgical candidates. Therefore, identifying non-invasive, highly sensitive, and easily accessible indicators is crucial. Hematological indicators, known for their simplicity and clinical utility in tumor screening, have become a common method for cancer evaluation. This raises the question of whether blood-based biomarkers offer clinical value for PNI assessment in GC. To our knowledge, there are few studies on PNI and hematological indicators in GC.

In our study, the cases were randomly divided into a training set (modeling group) and a testing set (validation

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Disgnastia officiancy	Estimated value	95% Cl		
Diagnostic enciency	Estimated value	Lower limit	Upper limit	
Sensitivity (%)	69.57	62.29	76.01	
Specificity (%)	56.41	46.94	65.45	
Positive predictive value (%)	71.51	64.21	77.87	
Negative predictive value (%)	54.10	44.86	63.07	
Positive likelihood ratio	1.59	1.27	2.00	
Negative likelihood ratio	0.54	0.43	0.68	

Table 6 Diagnostic efficiency for PNI in validation group

PNI, perineural invasion; CI, confidence interval.



Figure 5 Validation of the prediction model. (A) ROC curve of model in the validation group. (B) Calibration curve of model in the validation group. (C) DCA of model in the validation group. AUC, area under the curve; CI, confidence interval; ROC, receiver operating characteristic; DCA, decision curve analysis.

group) at a ratio of 7:3. In the modeling group, the medians of age, D-dimer, NLR, PLR, FLR, LMR, ALR, CEA, CA72-4, CA19-9, and CA125 were remarkably higher in GC patients with PNI than in those who were PNInegative. Moreover, through the random forest algorithm, the top 5 important risk factors for PNI patients were CEA, FLR, D-dimer, PLR, and CA19-9. These results suggest that the occurrence of PNI in GC patients has a certain relationship with hematological indicators. Multivariate logistic regression analysis indicated that CEA >3.89 ng/mL, FLR >2.08, D-dimer >0.24 mg/L, PLR >122.37, CA19-9 >14.85 U/mL, and CA72-4 >2.31 U/mL are independently associated factors with PNI in GC patients. Meanwhile, we developed the equation: $Logit(P) = -1.211 + 0.695 \times CEA +$ 0.546 × FLR + 0.686 × D-dimer + 0.653 × PLR + 0.515 × CA19-9 + 0.518 × CA72-4 (χ^2 =105.675, P<0.001) to predict the presence of PNI. The model demonstrated that an ROC-AUC value of 0.719 (95% CI: 0.681–0.757) in the training set, with a sensitivity of 68.51% and a specificity of 67.60%. In addition, the calibration curve and DCA showed that the model has good discrimination and accuracy.

Growing evidence has indicated that there is an inflammatory link between tumor, microenvironment, and the systemic response. Different inflammatory markers have been analyzed in many cancers, including FLR and C-reactive protein in head and neck (36,37), PLR in thyroid carcinoma (28), and NLR in GC (38). Lymphocytes play a role in antitumor immune activity and tumor-related immune response (39). Lymphopenia has been observed in advanced cancer patients and its association with poor outcomes in patients with various types of cancer has been demonstrated (40). Fibrinogen plays a key role

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in coagulation, cell adhesion, systemic inflammation, and cancer progression (41,42). High levels of fibrin are indicators of coagulation and fibrinolysis activation, and almost all types of tumors exhibit abnormalities in coagulation and fibrinolysis, thereby promoting the body to be in a hypercoagulable state (43). D-dimer-containing species are soluble fibrin degradation products derived from plasmin-mediated degradation of cross-linked fibrin, which could be considered a biomarker of in vivo activation of both coagulation and fibrinolysis (44). Beyond that, platelets can secrete various growth factors to stimulate tumor cell differentiation, especially in advanced stage tumor patients who often have elevated platelets (45). Despite these insights, the relationship between blood-based biomarkers and PNI in GC remains inconclusive. PNI is posited as a novel pathway for tumor metastasis, with patients exhibiting increased levels of FLR, PLR, and D-dimer, suggesting heightened fibrinogen and platelet counts and reduced lymphocyte levels. These changes are associated with an elevated risk of PNI, indicating a poor prognosis (36,46,47). Common tumor markers such as CEA, CA19-9, and CA72-4 were identified as independent risk factors of GC with PNI in the present study, consistent with the findings of Liu et al. This study suggested that CEA $\geq 5 \mu g/L$ is a significant independent risk factor of PNI in advanced GC (20). Meanwhile, Huang et al. evaluated an optimal cutoff value of FLR of 2.555 (OR: 1.266, 95% CI: 1.031-1.555, P=0.02), where an FLR above 2.555 is a risk factor for GC with peritoneal dissemination (48).

Immediately after, we verified the discriminant equation model using the data of the testing set. The AUC-ROC was 0.791 (95% CI: 0.750-0.831, P<0.01) in the validation group. The sensitivity, specificity, PPV, and NPV were 69.57%, 56.41%, 71.51%, and 54.10%, respectively. These results suggested that the model has good predictive value for the risk of PNI in GC patients. The calibration curve also indicated that the predictive results were good in accordance with the actual results. PNI can serve as an important reference indicator for the biological behavior of GC, which is of great significance for improving tumor staging, selecting treatment plans rationally, and evaluating prognosis. A previous study suggested that PNI may be related to positive surgical margins (49). Therefore, preoperative prediction of PNI and appropriate expansion of tumor resection range during surgery maybe reduce tumor recurrence. In addition, PNI prediction is invaluable for patients ineligible for surgery, guiding the development of alternative treatment strategies.

The present study, a single-center retrospective analysis, has several inherent limitations. Firstly, potential selection bias and limitations in data extraction may have influenced the study's conclusions. Secondly, the cut-off value of CEA, FLR, D-dimer, PLR, CA19-9, and CA72-4 was calculated only by mathematical methods, and the sensitivity and specificity of these laboratory data in predicting PNI require further verification in multi-center and large cohorts. Thirdly, some other inflammatory indicators associated with cancer progression, such as procalcitonin, interleukin-6, and C-reactive protein were not included in our study. Nonetheless, this study represents a novel approach, being the first to suggest the potential of preoperative CEA, FLR, D-dimer, PLR, CA19-9, and CA72-4 as predictors of PNI in GC patients.

Conclusions

We have successfully established and validated a predictive model for GC patients with PNI based on hematological indicators. The preoperative assessment of hematological inflammatory markers offers valuable insights for evaluating the risk of PNI in GC patients. This approach may serve as an adjunctive tool in clinical decision-making processes regarding PNI.

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Footnote

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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) and approved by the Clinical Research Ethics Committee of Jiangsu Cancer hospital (approval number: #KY-2024-012). The requirement for informed consent was waived in this retrospective study.

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