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***Retrospective Study***

**Nomogram using F-18 fluorodeoxyglucose positron emission tomography/computed tomography for preoperative prediction of lymph node metastasis in gastric cancer**

Song BI. Preoperative prediction of lymph node metastasis in gastric cancer

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**Abstract**

BACKGROUND

Lymph node (LN) metastasis is an important prognostic factor in patients with gastric cancer (GC). However, the evaluation of LN metastasis status in the preoperative setting is not accurate. Therefore, precise preoperative prediction of LN metastasis status is crucial for optimal treatment in patients with GC.

AIM

To develop a preoperative nomogram for LN metastasis using F-18 fluorodeoxyglucose (F-18 FDG) positron emission tomography/computed tomography (PET/CT) and preoperative laboratory test findings in GC.

METHODS

In this study, the data of 566 GC patients who underwent preoperative F-18 FDG PET/CT and subsequent surgical resection were analyzed. The LN metastasis prediction model was developed in the training cohort and validated in the internal validation cohort. Routine preoperative laboratory tests, including albumin and carbohydrate antigen (CA) 19-9 were performed in all patients. Univariate and multivariable logistic regression was performed to validate the preoperative predictive indicators for LN metastasis.

RESULTS

Of the 566 patients, 232 (41%) had confirmed histopathologic LN metastasis. Univariate logistic regression revealed that the tumor location, blood hemoglobin, serum albumin levels, neutrophil to lymphocyte ratio, platelet to lymphocyte ratio, CA 19-9, maximum standardized uptake value (SUVmax) of the primary tumor (T\_SUVmax), and SUVmax of LN (N\_SUVmax) were significantly associated with LN metastasis. In multivariate analysis, T\_SUVmax (OR = 1.08; 95%CI: 1.02–1.15; *P* = 0.011) and N\_SUVmax (OR = 1.49; 95%CI: 1.19–1.97; *P* = 0.002) were found to be significant predictive factors for LN metastasis. The LN metastasis prediction model using T\_SUVmax, N\_SUVmax, serum albumin, and CA 19-9 yielded an area under the curve (AUC) of 0.733 (95%CI: 0.683–0.784, *P* = 0.025) in the training cohort and AUC of 0.756 (95%CI: 0.678–0.833, *P* < 0.001) in the test cohort.

CONCLUSION

T\_SUVmax and N\_SUVmax measured by preoperative F-18 FDG PET/CT are independent predictive factors for LN metastasis in GC.

**Key words** Gastric cancer; Lymph node metastasis; positron emission tomography/computed tomography; fluorodeoxyglucose; Prognostication; Standardized uptake value

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**Core tip:** The maximum standardized uptake values of the primary tumor and lymph node (LN), measured by preoperative F-18 fluorodeoxyglucose positron emission tomography/computed tomography, are independent predictive factors for LN metastasis in patients with gastric cancer. Moreover, a nomogram using a combination of these metabolic information and laboratory parameters, such as serum albumin and carbohydrate antigen 19-9, for risk estimation of LN metastasis in gastric cancer was successfully developed in the training cohort and validated in the internal validation cohort.

**INTRODUCTION**

Gastric cancer (GC) is one of the most commonly diagnosed malignancies and the second leading cause of cancer-related deaths worldwide[1]. The status of lymph node (LN) metastasis is an important prognostic factor in GC, and complete dissection of the metastatic LNs is the only curative treatment for GC[2]. Although contrast-enhanced computed tomography (CT) and endoscopic ultrasonography (EUS) are used for the diagnosis of LN metastasis in GC, the accuracy of diagnostic performance for LN metastasis is imperfect[3,4].

Positron emission tomography/computed tomography (PET/CT) with 18F-fluorodeoxyglucose (F-18 FDG) has become a useful diagnostic modality for staging, treatment response evaluation, and detection of recurrence in GC[5,6]. However, F-18 FDG PET/CT has shown relatively low sensitivity in the detection of LN metastasis in GC[7,8]. Recently, metabolic information of the primary tumor obtained using F-18 FDG PET/CT has been suggested as a promising predictive marker for LN metastasis[9–11]. Glucose metabolism in the primary tumor reflects not only the total tumor burden, but also the aggressiveness of cancer associated with LN metastasis. Therefore, a combination of the metabolic information of the primary tumor and metastatic LN could be useful in predicting LN metastasis in GC.

Recently, a few studies have been undertaken to develop a nomogram for the prediction of LN metastasis in GC[12,13]. However, these LN metastasis prediction models are based on postoperative parameters. Nevertheless, a preoperative LN metastasis prediction model, based on the tumor metabolic information as measured by F-18 FDG PET/CT, does not exist for GC. This model would be crucial for clinicians to determine the most effective treatment strategy.

The aim of this retrospective study was to determine whether the metabolic information of LN, as well as the primary tumor, could be prognostic factors for the prediction of LN metastasis in GC and to develop a preoperative nomogram for the prediction of LN metastasis in GC.

**MATERIALS AND METHODS**

***Patients***

Between January 2008 and December 2010, the medical records of 873 consecutive patients who underwent surgery for primary GC at Keimyung University Dongsan Medical Center (Daegu, South Korea) were retrospectively reviewed. Of these, 566 patients who underwent preoperative F-18 FDG PET/CT and subsequent surgical treatment without any preoperative intervention were enrolled in this study. The exclusion criteria were as follows: any other treatment prior to surgery such as gastric endoscopic submucosal dissection or chemotherapy, multiple primary malignancies, surgery for recurred GC, unavailable clinicopathological report, or an interval over 1 month between F-18 FDG PET/CT and surgery. A total of 566 GC patients were randomly divided into 377 of the training cohort and 189 of the internal validation cohort (2:1) (Figure 1). This retrospective study was approved by the Institutional Review Board of Keimyung University Dongsan Medical Center. The need for informed consent was waived, and all data were anonymized prior to the analysis.

All patients underwent subtotal or total gastrectomy along with D1+β or D2 lymphadenectomy in early GC and D2 lymphadenectomy in advanced GC. Clinicopathological data, including age at surgery, sex, location of the tumor, pathologic T (pT) stage, serum white blood cell (WBC) count, blood hemoglobin and serum albumin levels, platelet count, neutrophil count, lymphocyte count, platelet to lymphocyte ratio (PLR), neutrophil to lymphocyte ratio (NLR), preoperative carcinoembryonic antigen (CEA), and carbohydrate antigen (CA) 19-9 were retrieved from the patients’ medical records. The pT stage was classified according to the 8th edition of the American Joint Committee on Cancer (AJCC) tumor-node-metastasis (TNM) staging system.

***F-18 FDG PET/CT scan and image analysis***

All the patients fasted for at least 6 h before F-18 FDG injection, and the blood glucose level was managed to be lower than 150 mg/dL. The PET/CT scan was performed 60 min after F-18 FDG was administered. PET/CT scans were performed using a Discovery STE PET/CT scanner (GE Healthcare, Milwaukee, WI, United States). First, a low-dose CT scan (peak voltage of 120 kVp, automated tube current ranging from 60 to 150 mA, and slice thickness of 3.75 mm) for attenuation correction without contrast enhancement was acquired. After CT scan, PET scan was performed immediately with an acquisition time of 3 min per bed position in 3D mode. The PET images were reconstructed using an ordered-subset expectation maximum iterative reconstruction algorithm.

All the F-18 FDG PET/CT images were retrospectively interpreted on an Advantage Workstation 4.3 (GE Healthcare), blinded to the status of LN metastasis. First, all F-18 FDG PET/CT images were visually assessed and classified as positive or negative with respect to F-18 FDG uptake by the primary tumor or LN. A positive uptake was defined as abnormally increased F-18 FDG uptake that exceeded the physiologic uptake by the surrounding stomach wall and corresponding cancer lesions on esophagogastroduodenoscopy. Consequently, the maximum standardized uptake value (SUVmax) of the primary tumor (T\_SUVmax) was obtained only in positive F-18 FDG uptake lesions. In case of LNs, SUVmax of LN (N\_SUVmax) was acquired in the highest focal F-18 FDG avid LN on PET image regardless of size on CT. A spherical volume of interest was manually drawn over the maximum F-18 FDG uptake lesions on the attenuation-corrected transaxial F-18 FDG PET images for semi-quantitative analysis. The SUVmax was calculated using the following formula: SUVmax = maximum activity in the region of interest (MBq/g)/ [injected dose (MBq)/body weight (g)].

***Statistical analysis***

Numeric data were expressed as the mean ± sd. First, all the factors that were significantly associated (*P* < 0.05) with LN metastasis were identified in univariate analysis, and these significant factors were then evaluated to determine the variables independently associated with LN metastasis using multivariate logistic regression. Second, the LN metastasis prediction model was developed using the multivariate logistic analysis with a stepwise backward elimination method in the training cohort, and validated in the internal validation cohort. All variables with *P* < 0.05 in the univariate logistic analysis were selected for multivariate logistic analysis in the training cohort, and deleting the variable whose loss gives the most statistically insignificant deterioration of the prediction model fit. Lastly, we developed a nomogram as a graphical tool for calculating the risk of LN metastasis in individual patients. All statistical analyses were performed using R version 3.5.0 software (http://www.r-project.org, R Foundation for Statistical Computing, Vienna, Austria). A P < 0.05 was considered statistically significant.

**RESULTS**

***Patient characteristics***

The characteristics of the enrolled patients and the associations of these characteristics with LN metastasis in the training cohort (*n* = 377) and internal validation cohort (n = 189) are summarized in Table 1. Of the 566 patients enrolled in the present study, 232 (41.0%) had pathologically confirmed LN metastasis and 334 patients (59.0%) presented with no LN metastasis. The sensitivity, specificity, and accuracy of F-18 FDG PET/CT for the diagnosis of LN metastasis in GC patients were 28.9%, 97.3%, and 69.3%, respectively. Clinicopathological findings; tumor location, pT stage, blood hemoglobin levels, platelet count, lymphocyte count, PLR, NLR, CA 19-9, serum albumin, and metabolic parameters; T\_SUVmax, and N\_SUVmax were significantly different between the two groups (with or without LV metastasis); however, no significant differences were found with respect to age, sex, WBC count, neutrophil count, and serum CEA in the training cohort.

***Uni- and multivariate logistic regression analyses***

Univariate logistic regression analysis revealed that tumor location, blood hemoglobin levels, platelet count, lymphocyte count, PLR, NLR, CA 19-9, serum albumin, T\_SUVmax, and N\_SUVmax were significantly associated with LN metastasis in the training cohort. In multivariate analysis, T\_SUVmax (OR = 1.08; 95%CI: 1.02–1.15; *P* = 0.011) and N\_SUVmax (OR = 1.49; 95%CI: 1.19–1.97; *P* = 0.002) were found to be independent predictive factors for LN metastasis in the training set (Table 2). Also, T\_SUVmax (OR = 1.17; 95%CI: 1.07–1.29; *P* < 0.001) and N\_SUVmax (OR = 1.60; 95%CI: 1.09–2.69; *P* = 0.038) were independent predictive factors for LN metastasis in the test set (Table 3).

***LN metastasis prediction model and nomogram***

The result of the stepwise backward regression showed that a prediction model that combines T\_SUVmax, N\_SUVmax, serum albumin, and CA 19-9 was the best model to predict the risk of LN metastasis in the training cohort. The Hosmer and Lemeshow test generated a *P* value of 0.484, indicating that this predictive model fits well. A nomogram for predicting the probability of LN metastasis using pretreatment F-18 FDG PET/CT parameters and laboratory findings was successfully developed (Figure 2). The performance of this LN metastasis prediction model was good with the area under the receiver operating characteristic curve (AUC) of 0.733 (95%CI: 0.683–0.784, *P* = 0.025) in the training cohort and AUC of 0.756 (95%CI: 0.678–0.833, *P* < 0.001) in the test cohort (Figure 3).

**DISCUSSION**

In the present study, the incidence of LN metastasis in GC patients was 41% and the diagnostic performance of F-18 FDG PET/CT was highly specific for LN metastasis status; however, it had a limitation due to its relatively low sensitivity. The present study revealed that T\_SUVmax and N\_SUVmax measured by preoperative F-18 FDG PET/CT are independent predictive factors for LN metastasis in patients with GC. Moreover, the combination of these metabolic parameters with clinical laboratory findings (albumin and CA 19-9) significantly improved prediction of LN metastasis, compared with each parameter alone.

Several previous studies demonstrated that F-18 FDG PET/CT had relatively low sensitivity in detecting LN metastasis in GC patients[7,8,14]. In agreement with those studies, the results of the present study showed relatively low sensitivity. Despite the high specificity in the detection of LN metastasis, routine use of F-18 FDG PET/CT for GC stating is still controversial due to its low sensitivity[14–16]. A few studies have found that F-18 FDG uptake by the primary gastric tumor may predict LN metastasis status. Oh *et al*[17] reported that the peak-SUV of the primary gastric tumor is a useful indicator for LN metastasis. Kim *et al*[18] demonstrated that the T\_SUVmax was the only independent factor to be significantly related to sensitivity for LN metastasis. However, no study has evaluated the predictive value of the combination of T\_SUVmax and N\_SUVmax for LN metastasis in GC. Notably, the present study showed that T\_SUVmax and N\_SUVmax were independent predictive factors for LN metastasis in the validation cohort as well as the training cohort.

Several studies have found that F-18 FDG uptake by the primary tumor is positively correlated with the status of LN metastasis in various cancers[9,19,20]. The present study also suggested that T\_SUVmax was an independent prognostic factor for LN metastasis in patients with GC. This result could be explained by the fact that T\_SUVmax reﬂects not only the tumor’s metabolic information, but also the tumor aggressiveness[21,22]. In this regard, T\_SUVmax could have an additional value in predicting LN metastasis by reducing the high false-negative rate of F-18 FDG PET/CT for LN metastasis in patients with GC.

Some studies have found that pretreatment serum albumin[23,24] and CA 19-9[25–27] levels are correlated with LN metastasis. The chronic systematic inflammatory state increases the vascular permeability and loss of serum protein. Hypoalbuminemia, therefore, results from and reflects the systematic inflammatory condition. The inflammatory component contributes to tumor proliferation, angiogenesis, and metastasis[28]. For this reason, the serum albumin level is associated with LN metastasis. Meanwhile, CA 19-9 is a tumor-associated antigen and has recently been demonstrated to be a marker of digestive tract malignancies, especially pancreatic cancer[29]. Accordingly, the pretreatment serum albumin and CA 19-9 levels could be promising predictive factors for LN metastasis in patients with GC.

Meanwhile, the positive rate of LN differs by T stage, and the clinical significance of preoperative prediction of LN also depends on T stage. However, the endpoint of this study was the development of the preoperative LN metastasis prediction model. Therefore, despite the positive rate of LN differing by T stage, T stage could not be considered as a predictive parameter in this study. Although, there are several studies for the precise diagnosis of T stage using endoscopic ultrasonography (EUS), the accuracy of EUS for T stage ranged between < 50% and > 90%[30–33]. In GC, accurate preoperative prediction of LN status according to the specific T stage could provide more detailed pretreatment decision making. Since T stage is one of the most important factors for not only LN status prediction but also treatment decision making, establishment of reliable and objective method for the accurate T stage could be a useful co-consideration parameter for the prediction of LN metastasis and GC treatment.

Recently, validation of nomograms for calculating the risk of LN metastasis in GC has been reported[12,13]. However, no study has yet established a nomogram for prediction of LN metastasis using preoperative clinical parameters. The present study successfully developed an effective nomogram to predict LN metastasis in GC using T\_SUVmax, N\_SUVmax, serum albumin, and CA 19-9. Considering the feasibility of F-18 FDG PET/CT in the preoperative setting of GC, F-18 FDG PET/CT could be used as a non-invasive diagnostic tool for assessment of LN metastasis status in patients with GC and can be used to optimize current treatment strategy for patients with GC patients. The accurate preoperative prediction of LN can support clinicians in classifying patients who could receive minimal surgery or may derive greater clinical benefit from more extensive treatment.

This study had a few limitations. First, this study was a single-institution, retrospective study that might have been subject to selection bias. External validation is necessary to assess transferability of the LN prediction model. Second, the SUV of the small-sized primary tumor and LNs could be underestimated due to partial-volume effects. Lastly, since F-18 FDG uptake can be elevated by not only the malignant cell, but also the inflammatory lesion, SUVmax might be overestimated in some patients.

In conclusion, T\_SUVmax and N\_SUVmax were independent prognostic factors for the prediction LN metastasis in GC patients. Further, a prediction model using metabolic parameters (T\_SUVmax and N\_SUVmax) and laboratory findings (albumin and CA 19-9) could provide a more precise prediction of LN metastasis in the preoperative setting. The use of preoperative F-18 FDG PET/CT could be a useful tool for LN metastasis evaluation and treatment planning in patients with GC.

**ARTICLE HIGHLIGHTS**

***Research background***

Gastric cancer (GC) is one of the most commonly diagnosed malignancies and the second leading cause of cancer-related deaths worldwide. The status of lymph node (LN) metastasis is an important prognostic factor in patients with GC. However, the evaluation of LN metastasis status in the preoperative setting is not accurate.

***Research motivation***

A few studies have been conducted to develop a nomogram for the prediction of LN metastasis in GC. However, a preoperative LN metastasis prediction model, based on the tumor metabolic information as measured by F-18 fluorodeoxyglucose (F-18 FDG) positron emission tomography/computed tomography (PET/CT) and laboratory findings, does not exist for GC. The purpose of this study was to develop a preoperative nomogram for LN metastasis in patients with GC.

***Research objectives***

This study aims to identify predictive factors and to develop a preoperative nomogram for the prediction of LN metastasis using F-18 FDG PET/CT and preoperative laboratory findings in patients with GC.

***Research methods***

Between 2008 and 2010, a total of 566 GC patients who underwent preoperative F-18 FDG PET/CT and subsequent surgical treatment without any preoperative intervention were analyzed. The LN metastasis prediction model was developed in the training cohort (*n* = 377) and validated in the internal validation cohort (*n* = 189). Clinicopathological data were retrieved from the patients’ medical records and the F-18 FDG PET/CT images were retrospectively interpreted. Univariate and multivariable logistic regression was performed to validate the preoperative predictive factors for LN metastasis.

***Research results***

The multivariate logistic analysis showed that the combination of maximum standardized uptake value (SUVmax) of the primary tumor (T\_SUVmax) and SUVmax of LN (N\_SUVmax), serum albumin, and carbohydrate antigen (CA) 19-9 was the best model to predict the risk of LN metastasis. The preoperative nomogram for the prediction of LN metastasis using T\_SUVmax, N\_SUVmax, serum albumin, and CA 19-9 showed good performance in the validation cohort as well as the training cohort.

***Research conclusions***

The combination of preoperative F-18 FDG PET/CT metabolic parameters (T\_SUVmax and N\_SUVmax) and laboratory findings (albumin and CA 19-9) could be a useful tool for LN metastasis assessment and treatment planning in patients with GC.

***Research perspectives***

The preoperative nomogram for the prediction of LN should be verified on a larger and external validation cohort for widespread acceptance.

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**Footnotes**

**Institutional review board statement:** This study was reviewed and approved by the Institutional Review Board of Keimyung University Dongsan Medical Center (IRB No. 2018-06-028-003).

**Informed consent statement:** The patients were not required to give informed consent to the study because the analysis used anonymous clinical data that were obtained after each patient agreed to treatment by written consent.

**Conflict-of-interest statement:** Song BI declare no relevant conflicts of interests.

**Data sharing statement:** No additional data are available.

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Grade A (Excellent): 0

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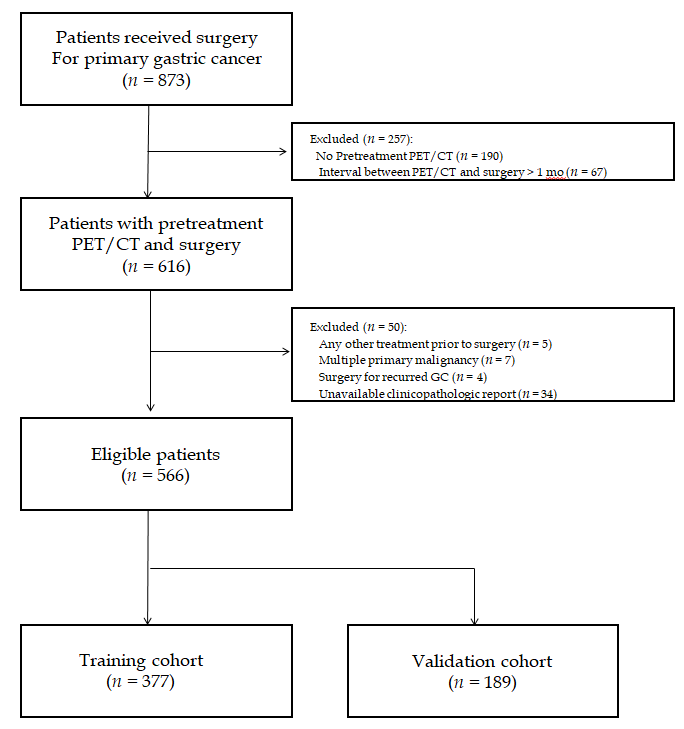
Grade C (Good): C, C

Grade D (Fair): D

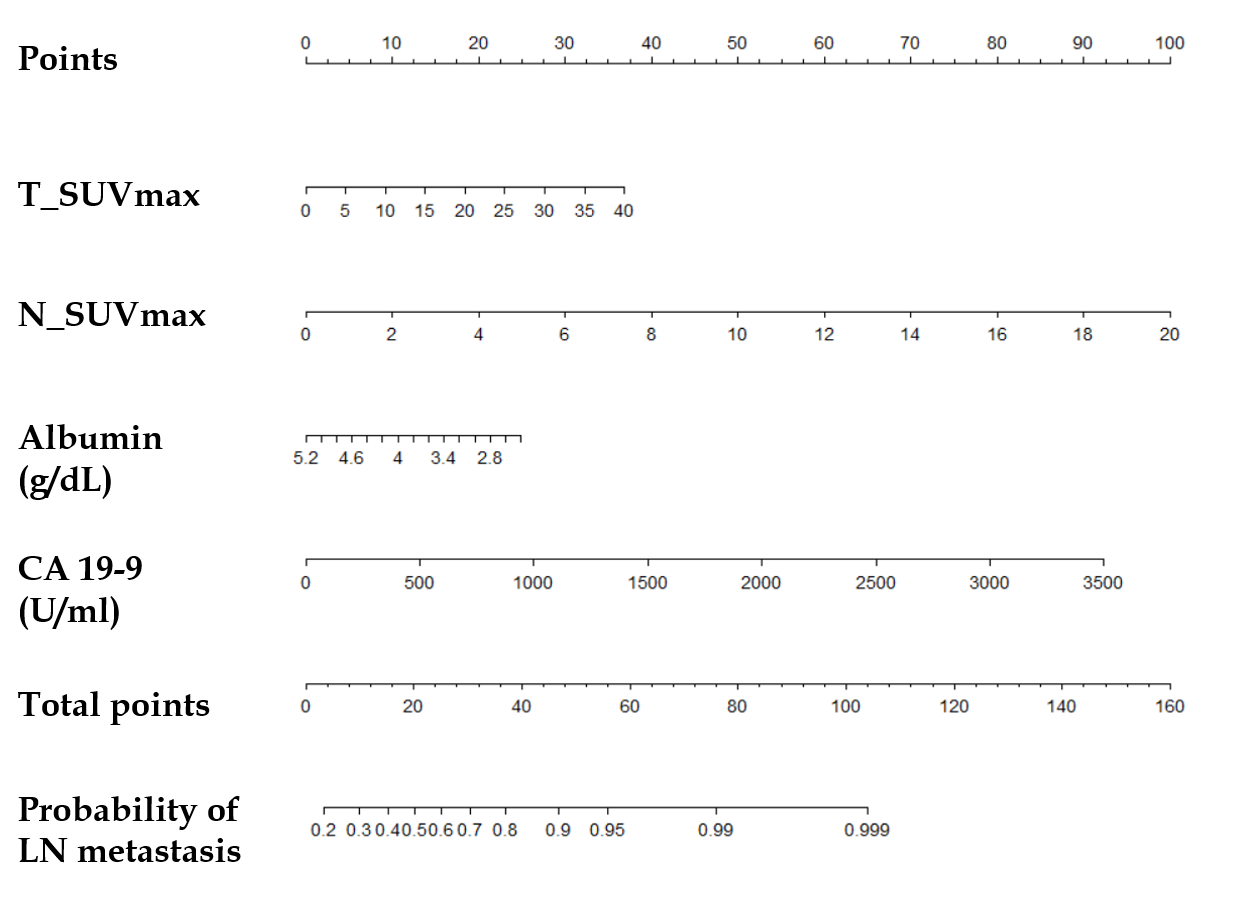
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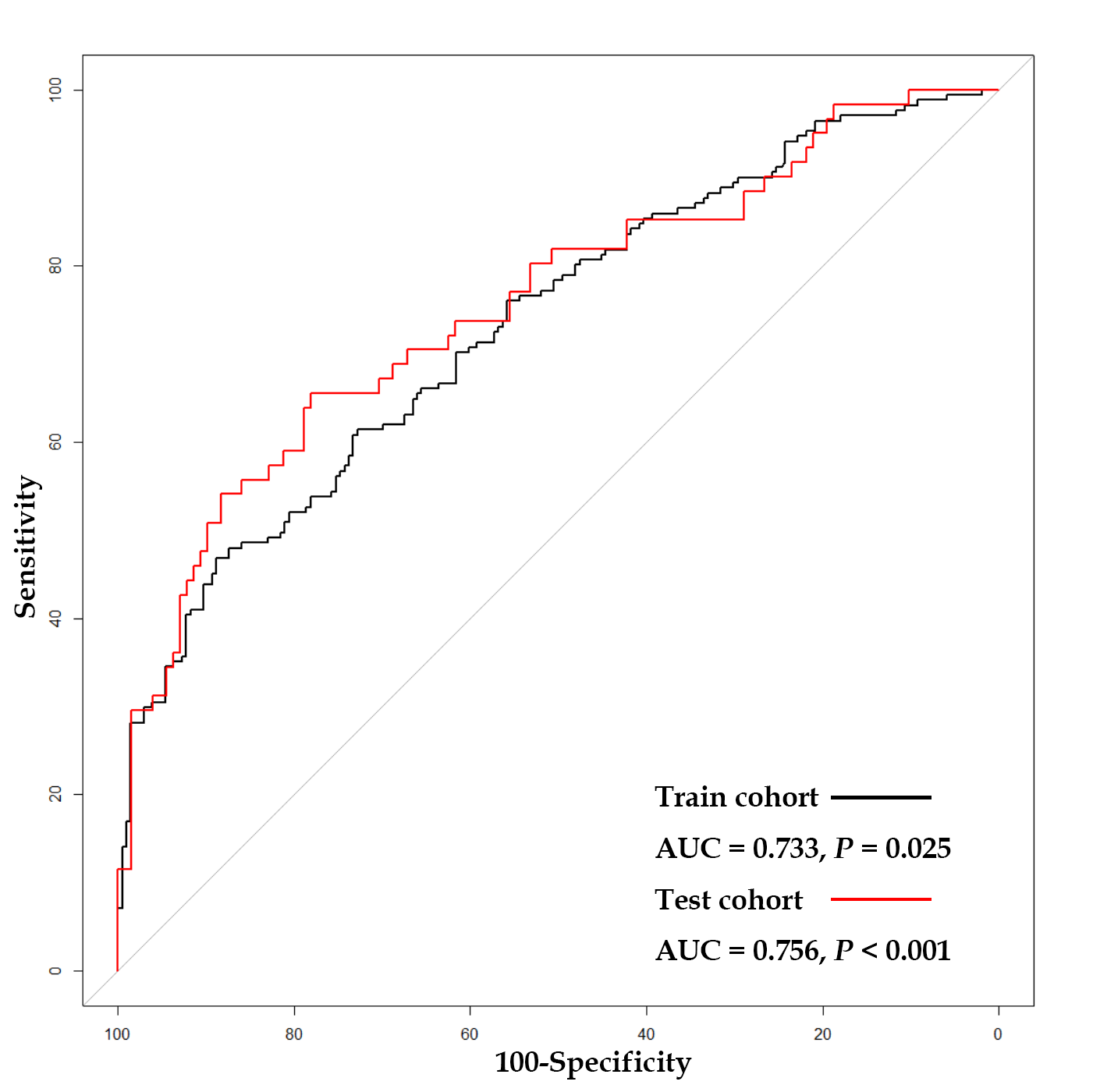
**Figure Legends**



**Figure 1 Flow diagram of patient selection.** PET/CT: Positron emission tomography/computed tomography; GC: gastric cancer.

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**Figure 2 Nomogram for predicting the risk of lymph node metastasis using preoperative F-18 Fluorodeoxyglucose positron emission tomography/computed tomography and laboratory parameters.** First, the number of points for each parameter –maximum standardized uptake value of primary tumor, maximum standardized uptake value of lymph node, albumin, and CA 19-9 – should be determined by drawing a vertical line from the exact value of variables to the points row. Subsequently, total points can be obtained by sum of four variables. The individual predictive risk of lymph node metastasis can be calculated by drawing a vertical line from the total points row to the probability of regional lymph node metastasis. T\_SUVmax: Maximum standardized uptake value of primary tumor; N\_SUVmax: Maximum standardized uptake value of lymph node; LN: Lymph node.



**Figure 3 C-statistic of the combination model using metabolic parameters (maximum standardized uptake value of primary tumor and maximum standardized uptake value of lymph node) and laboratory findings (albumin and CA 19-9).** C-statistic using receiver operating characteristic curve analysis, the area under the curve was 0.733 (95%CI: 0.683–0.784, *P* = 0.025) for lymph node metastasis prediction performance in the training cohort (black line), and area under the curve was of 0.756 (95%CI: 0.678–0.833, *P* < 0.001) in the test cohort (red line). AUC: Area under the curve.

T**able 1 Patient characteristics**

|  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- |
|  | **Training cohort** | | | **Validation cohort** | | | |
| **Variables** | **LN metastasis (-)** | **LN metastasis (+)** | ***P-*value** | | **LN metastasis (-)** | **LN metastasis (+)** | ***P-*value** |
| **(*n =* 206)** | **(*n =* 171)** | **(*n =* 128)** | **(*n =* 61)** |
| Age | 59.2 ± 11.5 | 59.6 ± 12.1 | 0.701 | | 59.5 ± 12.1 | 61.8 ± 12.8 | 0.218 |
| Sex |  |  | 0.577 | |  |  | 0.456 |
| Male | 122 (59.2%) | 107 (62.6%) |  | | 73 (57.0%) | 39 (63.9%) |  |
| Female | 84 (40.8%) | 64 (37.4%) |  | | 55 (43.0%) | 22 (36.1%) |  |
| Tumor Location |  |  | < 0.001 | |  |  | 0.034 |
| Upper | 46 (22.3%) | 27 (15.8%) |  | | 25 (19.5%) | 15 (24.6%) |  |
| Middle | 32 (15.5%) | 35 (20.5%) |  | | 24 (18.8%) | 11 (18.0%) |  |
| Low | 122 (59.2%) | 81 (47.4%) |  | | 78 (60.9%) | 30 (49.2%) |  |
| Mixed | 6 (2.9%) | 28 (16.4%) |  | | 1 ( 0.8%) | 5 (8.2%) |  |
| Pathologic T stage |  |  | < 0.001 | |  |  | < 0.001 |
| 1 | 158 (76.7%) | 28 (16.4%) |  | | 100 (78.1%) | 10 (16.4%) |  |
| 2 | 20 (9.7%) | 27 (15.8%) |  | | 11 (8.6%) | 12 (19.7%) |  |
| 3 | 18 (8.7%) | 27 (15.8%) |  | | 8 (6.2%) | 18 (29.5%) |  |
| 4 | 10 (4.9%) | 89 (52.0%) |  | | 9 (7.0%) | 21 (34.4%) |  |
| WBC counts (103 cells/μl) | 6.4 ± 1.8 | 6.2 ± 1.7 | 0.283 | | 6.2 ± 1.6 | 6.5 ± 1.8 | 0.295 |
| Blood hemoglobin levels (g/dl) | 12.5 ± 1.6 | 12.0 ± 1.8 | 0.004 | | 13.3 ± 9.7 | 12.2 ± 2.1 | 0.211 |
| Platelet counts (103 cells/μl) | 267.2 ± 72.6 | 287.9 ± 91.0 | 0.017 | | 267.3 ± 74.4 | 282.0 ± 67.3 | 0.192 |
| Neutrophil counts (cells/μl) | 3777.4 ± 1458.7 | 3810.9 ± 1333.2 | 0.818 | | 3686.8 ± 1367.7 | 3902.9 ± 1405.2 | 0.315 |
| Lymphocyte counts (cells/μl) | 1860.3 ± 609.4 | 1661.1 ± 589.8 | 0.001 | | 1835.0 ± 521.7 | 1855.1 ± 764.4 | 0.853 |
| PLR | 159.3 ± 72.8 | 196.4 ± 98.8 | < 0.001 | | 155.3 ± 60.3 | 184.8 ± 111.5 | 0.057 |
| NLR | 2.3 ± 1.4 | 2.5 ± 1.2 | 0.036 | | 2.2 ± 1.2 | 2.6 ± 2.0 | 0.136 |
| CEA (ng/ml) | 4.8 ± 25.9 | 7.9 ± 27.9 | 0.268 | | 2.6 ± 3.5 | 3.7 ± 6.1 | 0.195 |
| CA 19-9 (U/ml) | 18.1 ± 62.7 | 96.0 ± 364.9 | 0.006 | | 10.0 ± 7.6 | 12.7 ± 8.6 | 0.029 |
| Albumin (g/dl) | 4.1 ± 0.3 | 3.9 ± 0.4 | <0.001 | | 4.0 ± 0.3 | 3.9 ± 0.4 | 0.032 |
| T\_SUVmax | 2.9 ± 4.4 | 6.1 ± 5.5 | <0.001 | | 2.2 ± 3.7 | 6.9 ± 6.7 | < 0.001 |
| N\_SUVmax | 0.1 ± 0.9 | 1.8 ± 3.7 | <0.001 | | 0.1 ± 0.5 | 1.7 ± 3.9 | 0.002 |

LN: Lymph node; WBC: White blood cell; PLR: Platelet to lymphocyte ratio; NLR: Neutrophil to lymphocyte ratio; CEA: Carcinoembryonic Antigen; CA 19-9: Carbohydrate antigen 19-9; SUVmax: Maximum standardized uptake value; T\_SUVmax: SUVmax of primary tumor; N\_SUVmax: SUVmax of LN.

**Table 2 Uni- and multivariate logistic regression analyses for regional lymph node metastases in the training cohort**

|  | **Univariate logistic analysis** | | | | |  | **Multivariate logistic analysis** | |
| --- | --- | --- | --- | --- | --- | --- | --- | --- |
| **Variables** | **OR (95%CI)** | ***P* value** | | | |  | **OR (95%CI)** | ***P* value** |
| Age, yr | 1.00 (0.99–1.02) | | | 0.700 |  | |  |  |
| Sex (Male *vs* Female) | 0.87 (0.57–1.32) | | 0.507 | | |  |  |  |
| Tumor location | 1.31 (1.04–1.65) | | 0.022 | | |  | 1.19 (0.92–1.55) | 0.178 |
| WBC counts | 0.94 (0.83–1.05) | | 0.283 | | |  |  |  |
| Blood hemoglobin levels | 0.84 (0.74–0.94) | | 0.005 | | |  | 1.07 (0.91–1.26) | 0.433 |
| Platelet counts | 1.00 (1.00–1.01) | | 0.016 | | |  | 1.00 (1.00–1.01) | 0.625 |
| Neutrophil counts | 1.00 (1.00–1.00) | | 0.817 | | |  |  |  |
| Lymphocyte counts | 1.00 (1.00–1.00) | | 0.002 | | |  | 1.00 (1.00–1.00) | 0.342 |
| PLR | 1.01 (1.00–1.01) | | < 0.001 | | |  | 1.00 (0.99–1.01) | 0.816 |
| NLR | 1.19 (1.01–1.41) | | 0.041 | | |  | 0.85 (0.65–1.10) | 0.234 |
| CEA | 1.00 (1.00–1.01) | | 0.290 | | |  |  |  |
| CA 19-9 | 1.00 (1.00–1.01) | | 0.018 | | |  | 1.00 (1.00–1.01) | 0.133 |
| Albumin | 0.26 (0.13–0.49) | | < 0.001 | | |  | 0.52 (0.23–1.15) | 0.110 |
| T\_SUVmax | 1.17 (1.11–1.24) | | < 0.001 | | |  | 1.08 (1.02–1.15) | 0.011 |
| N\_SUVmax | 1.81 (1.45–2.40) | | < 0.001 | | |  | 1.49 (1.19–1.97) | 0.002 |

WBC: White blood cell; PLR: Platelet to lymphocyte ratio; NLR: Neutrophil to lymphocyte ratio; CEA: Carcinoembryonic Antigen; CA 19-9: Carbohydrate antigen 19-9; SUVmax: Maximum standardized uptake value; T\_SUVmax: SUVmax of primary tumor; N\_SUVmax: SUVmax of LN.

**Table 3 Uni- and multivariate logistic regression analyses for regional lymph node metastases in the test cohort**

|  | **Univariate Logistic Analysis** | |  | **Multivariate Logistic Analysis** | |
| --- | --- | --- | --- | --- | --- |
| **Variables** | **OR (95%CI)** | ***P* value** |  | **OR (95%CI)** | ***P* value** |
| Age, yr | 1.02 (0.99–1.04) | 0.218 |  |  |  |
| Sex (Male *vs* Female) | 0.75 (0.40–1.40) | 0.367 |  |  |  |
| Tumor location | 0.97 (0.68–1.40) | 0.881 |  |  |  |
| WBC counts | 1.10 (0.92–1.33) | 0.295 |  |  |  |
| Blood hemoglobin levels | 0.95 (0.82–1.02) | 0.451 |  |  |  |
| Platelet counts | 1.00 (1.00–1.01) | 0.194 |  |  |  |
| Neutrophil counts | 1.00 (1.00–1.00) | 0.315 |  |  |  |
| Lymphocyte counts | 1.00 (1.00–1.00) | 0.832 |  |  |  |
| PLR | 1.00 (1.00–1.01) | 0.027 |  | 1.00 (0.99–1.00) | 0.632 |
| NLR | 1.19 (0.98–1.46) | 0.087 |  |  |  |
| CEA | 1.05 (0.99–1.13) | 0.138 |  |  |  |
| CA 19-9 | 1.04 (1.00–1.08) | 0.032 |  | 1.03 (0.99–1.07) | 0.185 |
| Albumin | 0.31 (0.12–0.80) | 0.017 |  | 1.00 (0.30–3.32) | 0.994 |
| T\_SUVmax | 1.23 (1.14–1.34) | < 0.001 |  | 1.17 (1.07–1.29) | < 0.001 |
| N\_SUVmax | 2.02 (1.43–3.29) | < 0.001 |  | 1.60 (1.09–2.69) | 0.038 |

WBC: White blood cell; PLR: Platelet to lymphocyte ratio; NLR: Neutrophil to lymphocyte ratio; CEA: Carcinoembryonic Antigen; CA 19-9: Carbohydrate antigen 19-9; SUVmax: Maximum standardized uptake value; T\_SUVmax: SUVmax of primary tumor; N\_SUVmax: SUVmax of LN.