

# Predicting Peritoneal Metastasis of Gastric Cancer Patients Based on Machine Learning

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## Abstract

**Objective:** The aim is to explore the prediction effect of 5 machine learning algorithms on peritoneal metastasis of gastric cancer.

**Methods:** 1080 patients with postoperative gastric cancer were divided into a training group and test group according to the ratio of 7:3. The model of peritoneal metastasis was established by using 5 machine learning (gbm(Light Gradient Boosting Machine), GradientBoosting, forest, Logistic and DecisionTree). Python pair was used to analyze the machine learning algorithm. Gbm algorithm is used to show the weight proportion of each variable to the result.

**Result:** Correlation analysis showed that tumor size and depth of invasion were positively correlated with the recurrence of patients after gastric cancer surgery. The results of the gbm algorithm showed that the top 5 important factors were albumin, platelet count, depth of infiltration, preoperative hemoglobin and weight, respectively. In training group: Among the 5 algorithm models, the accuracy of GradientBoosting and gbm was the highest (0.909); the AUC values of the 5 algorithms are gbm (0.938), GradientBoosting (0.861), forest (0.796), Logistic(0.741) and DecisionTree(0.712) from high to low. In the test group: among the 5 algorithm models, the accuracy of forest, DecisionTree and gbm was the highest (0.907); AUC values ranged from high to low to gbm (0.745), GradientBoosting (0.725), forest (0.696), Logistic (0.680) and DecisionTree (0.657).

**Conclusion:** Machine learning can predict the peritoneal metastasis in patients with gastric cancer.

## Keywords

machine learning, peritoneal metastasis, gastric cancer, predictive modeling

## Introduction

Gastric cancer (GC) is a highly malignant and heterogeneous tumor. Among the malignant tumors worldwide, the incidence rate is the fourth and the mortality rate is the second.<sup>1</sup> Peritoneal membrane is the common metastatic site of gastric cancer. The studies<sup>2,3</sup> showed that 8.0% ~ 13.5% of patients with newly diagnosed gastric cancer were complicated with malignant ascites, while the incidence of peritoneal metastasis was higher than 39.0% ~ 43.0% in patients with advanced gastric cancer. Patients with advanced gastric cancer, especially those with malignant ascites due to peritoneal metastases, have a poorer prognosis. The median survival of patients with gastric cancer and peritoneal metastasis is only<sup>4-6</sup> months.<sup>2</sup> At the same time, malignant ascites can lead to complications such as intestinal obstruction, infection, malnutrition and renal insufficiency, which seriously affect the patients' quality of life. The early

stage of peritoneal metastasis of gastric cancer is mainly micro-metastasis, the small size and low density of peritoneal tumor nodules, so how to correctly diagnose peritoneal metastasis in early-stage has been the subject of clinical researchers'

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attention. Laparoscopic or laparotomy pathology is the “gold standard” for the diagnosis of peritoneal metastases, but both are invasive and are not suitable for routine performance. CT and PET/CT have commonly used imaging examinations for the diagnosis of recurrent metastasis after radical gastrectomy for gastric cancer, but there is some radiation and high cost.

At present, there are some predictive models for peritoneal metastasis of advanced gastric cancer. Studies<sup>4</sup> have shown that the clinical features combined with CT can predict the peritoneal metastasis of advanced gastric cancer. Similarly, some studies<sup>5</sup> have shown that texture features obtained from preoperative CT images of patients with advanced gastric cancer can be used to predict peritoneal metastasis. Other studies<sup>6</sup> have shown that venous CT radiological analysis based on primary tumors provides valuable information for predicting peritoneal metastasis of advanced gastric cancer. However, there is no research on the peritoneal metastasis of gastric cancer related to artificial intelligence.

Currently, based on large data sets and in-depth learning, researchers can use medical data and machine learning to predict disease risk better. Machine learning can translate measurements into relevant prediction models. Through machine learning, the diagnostic and drug genetics experts can find out the complexity of the disease, perform treatments, and customize medical options for individual patients. This study has reported that the combination of anti-cancer antigen 125 level and machine learning can predict the recurrence of abdominopelvic cancer.<sup>7</sup> Machine learning combined with MRI can predict the prognosis of breast tumor patients early<sup>8</sup>; Machine-based learning and radiomics can help improve the diagnostic performance of prostate cancer.<sup>9</sup> Convolutional neural network classifier can effectively distinguish bone metastasis of prostate cancer patients.<sup>10</sup> Other studies<sup>11</sup> have shown that machine learning can predict the effect of immune tumor-related gene expression on immune checkpoint inhibition in gastrointestinal cancer. Moreover, machine learning analysis can help clinicians determine the scope of lymph node dissection in gastric cancer before surgical resection.<sup>12</sup>

However, studies on peritoneal metastasis of gastric cancer related to machine learning have not been reported. Therefore, this study intends to investigate the effect of 5 machine learning algorithms on predicting peritoneal recurrence in gastric cancer.

## Materials and Methods

### Patients

In this retrospective analysis, we reviewed the data of 1199 GC patients who underwent GC surgery. Data is available at BioStudies database (<https://www.ebi.ac.uk/biostudies/studies?query=S-EPMC5383064>), accession numbers: S-EPMC5383064. All patients underwent a preoperative CT scan and were CT negative for peritoneal metastasis. The following information was collected and recorded: the patient's personal information (i.e., age, sex, body mass index, family history), tumor characteristics (i.e., location, size, type of pathology, histopathological differentiation, lymphatic invasion), and blood routine indices

**Table 1.** Baseline Data.

Peritoneal metastasis	NO	YES	P-value
N	979	101	
Age (years)	63.8 ± 11.2	63.6 ± 12.1	0.958
Height(cm)	165.2 ± 8.0	164.4 ± 9.1	0.584
Weight(kg)	59.4 ± 10.5	57.0 ± 11.0	0.035
BMI (kg/m <sup>2</sup> )	21.7 ± 3.0	21.1 ± 3.2	0.037
Tumor size (cm)	3.9 ± 2.1	5.0 ± 2.2	<0.001
PLR	148.8 ± 73.7	170.3 ± 77.7	<0.001
NLR	2.6 ± 1.6	2.9 ± 1.6	0.011
PREOPERATIVE.	118.4 ± 24.0	112.4 ± 22.6	0.003
HEMOGLOBIN			
Platelet count	225.3 ± 69.6	241.9 ± 72.4	0.035
Albumin (g/L)	39.7 ± 5.2	38.6 ± 5.1	0.016
Neutrophil count	4.1 ± 3.9	4.0 ± 1.3	0.151
Lymphocyte count	1.8 ± 2.0	1.6 ± 0.5	0.034
Monocyte count	0.5 ± 0.4	0.4 ± 0.2	0.144
WBC count	6.1 ± 1.6	6.2 ± 1.5	0.735
Sex			0.893
Male	760 (77.6%)	79 (78.2%)	
Female	219 (22.4%)	22 (21.8%)	
ASA			0.905
I	61 (6.2%)	5 (5.0%)	
2	828 (84.6%)	86 (85.1%)	
3	90 (9.2%)	10 (9.9%)	
TNM			0.958
I	236 (24.1%)	23 (22.8%)	
II	67 (6.8%)	8 (7.9%)	
III	513 (52.4%)	52 (51.5%)	
IV	163 (16.6%)	18 (17.8%)	
Borrmann types			0.041
I	55 (5.6%)	8 (7.9%)	
2	859 (87.7%)	80 (79.2%)	
3	65 (6.6%)	13 (12.9%)	
Pathological type [n, (%)]			0.015
Ulcerative	120 (12.3%)	21 (20.8%)	
Nonulcerative	859 (87.7%)	80 (79.2%)	
Depth of invasion [n, (%)]			<0.001
T1/T2	341 (34.8%)	6 (5.9%)	
T3/T4	638 (65.2%)	95 (94.1%)	

Note: NLR, neutrophil-to-lymphocyte ratio; PLR, platelet-to-lymphocyte ratio; WBC, white blood cell.

(i.e., neutrophils, lymphocytes, platelets, monocytes, NLR, and PLR). Pathological types were divided into ulcerative group and non-ulcerative group. The diagnosis was confirmed by histological examination in all patients. Exclusion criteria included (1) history of gastrectomy (8 patients), (2) liver disease such as cirrhosis (14 patients), (3) history of other malignancies (15 patients), (4) severe bleeding and autoimmune disease (22 patients), (5) preoperative chemoradiotherapy (2 patients), (6) severe inflammatory or hematological disorders (34 patients), and (7) distant metastasis (except abdominal metastases) (6 patients). Finally, the study enrolled in 1080 patients.

### Diagnosis of Peritoneal Metastases

According to the “Japanese Guidelines for the Treatment of Gastric Cancer” (fifth edition), the diagnostic criteria for

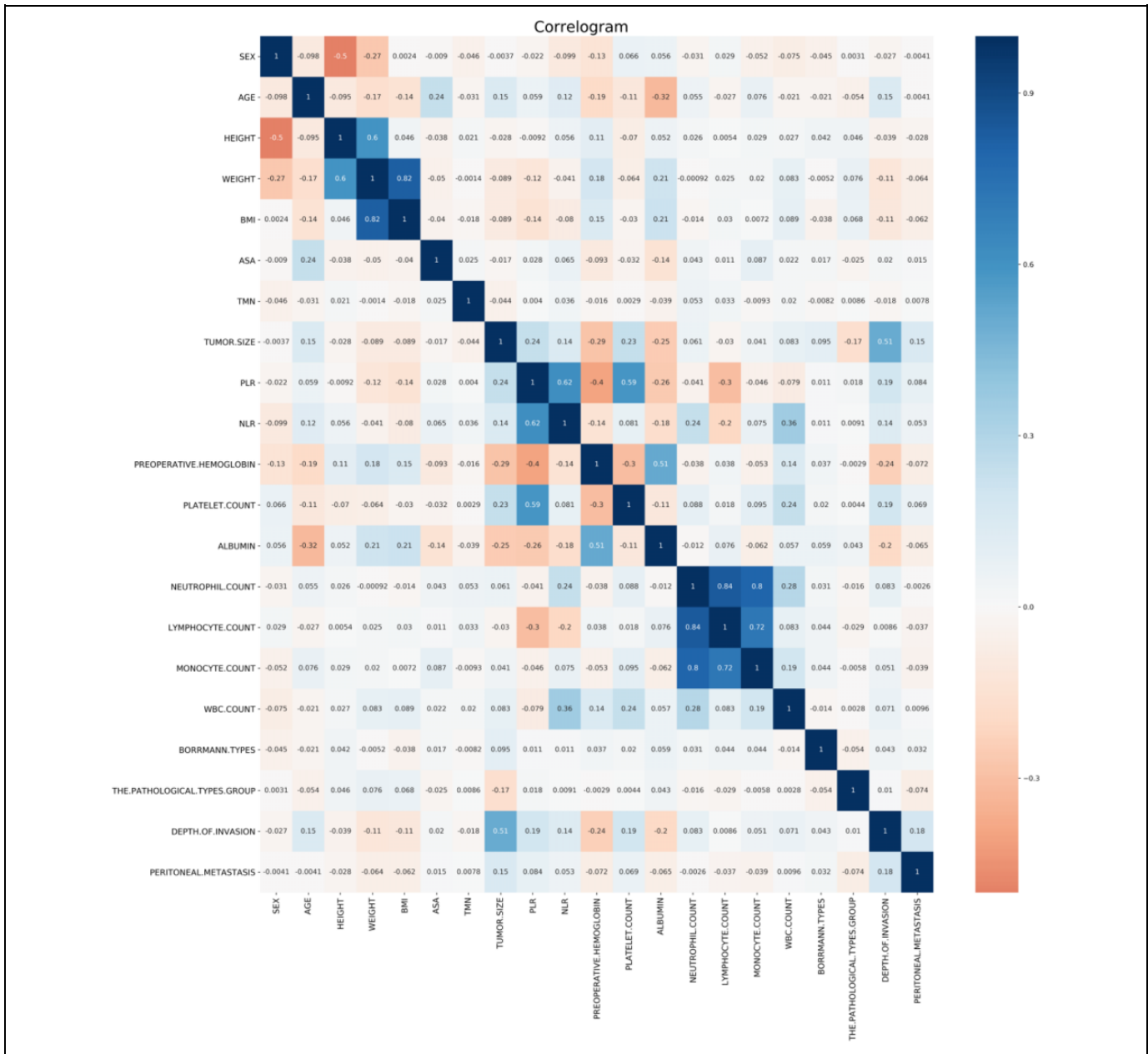


Figure 1. Correlation between variables.

peritoneal metastasis are as follows: metastasis is limited to the greater omentum, lesser omentum, anterior lobe of transverse colon, pancreatic capsule and spleen; Metastases and metastases to upper abdominal peritoneum (visceral peritoneum above transverse position and parietal peritoneum above umbilical cord). These patients with peritoneal metastases were diagnosed by intraoperative cryosections and postoperative pathological diagnosis.

### Machine Learning

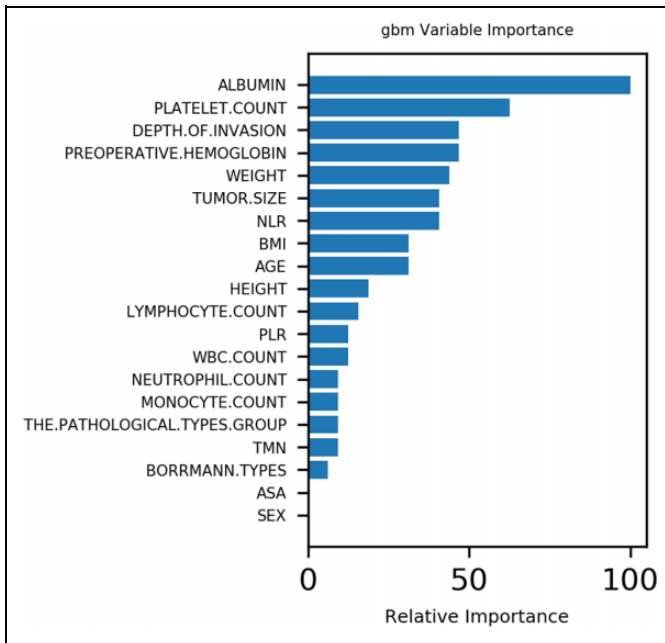
Logistic regression is essentially a surveillance classification algorithm. Logistic regression establishes a regression model to

predict the probability that a given data entry falls into a category numbered “1.”

Decision tree learning is a decision model established by tree structure based on the attributes of data, which can be used to solve the classification and regression problems.

A random forest is a proprietary noun that represents the overall decision tree. In order to classify it according to the attributes of a new object, each decision tree has a classification called this decision tree “vote” to the classification.

GBDT (Gradient Boosting Decision Tree) is a decision tree algorithm based on iteration. It is a Boosting method, and its main idea is that each establishment of the model is the gradient descent direction in which the model loss function was established before.



**Figure 2.** Variable importance of features included in machine learning algorithm for prediction of peritoneal metastasis.

LightGBM (Light Gradient Boosting Machine) adopts the Histogram algorithm. The idea is to divide the continuous floating-point characteristics into  $k$  discrete values and construct the Histogram with  $k$  width.

### Data Processing

The data were processed by R language, and the measurement data were expressed by the “mean, standard deviation” and tested by T-test. Counting data are expressed by the number of examples ( $n$ ) and percentage (%) and  $\chi^2$  test is adopted. The difference was statistically significant with  $p < 0.05$ . Multiple interpolations are used for missing variables. Python pair was used to analyze the machine learning algorithm. The total population was randomly divided into a training group and test group according to the ratio of 7:3. Meanwhile, the data were normalized and the prognostic weight was constructed.

Correlation analysis is used to observe the relationship between variables and show the correlation values. In particular, 5 different classification techniques have been evaluated: gbm, GradientBoosting, forest, Logistic and DecisionTree. Gbm algorithm is used to show the weight proportion of each variable to the result. Regularization is used to correct the over-fitting of the model. The prediction performance of the model is adjusted by manual and net style parameters. The following indicators are used to evaluate the prediction ability of the model: Accuracy, MSE and AUC.

Accuracy = (number of samples correctly classified)/(number of all samples classified). MSE (Mean Squared Error) is called mean squared error, the smaller, the better. ROC curve: receiver operating characteristic curve is a comprehensive index reflecting sensitivity and continuous specificity

variables. AUC value is between 0 -1, the greater, the better. The parameters and data packets used to build the machine learning model are shown in Appendix Table 1. Codes related to this research can be downloaded from GitHub website (<https://github.com/qazq124/-Peritoneal-Metastasis-of-Gastric-Cancer/blob/master/ML%20code12.pdf>).

### Result

Of the 1080 patients, 839 were men and 241 were women. Patients with GC and peritoneal metastases had a significantly higher PLR than patients without peritoneal metastases ( $P < 0.001$ ). Similarly, GC patients with peritoneal metastases had higher NLR ( $P = 0.011$ ). Age and height were not statistically different between the 2 groups. (See in Table 1) The basic information features of the training and test group are shown in Appendix Table 2.

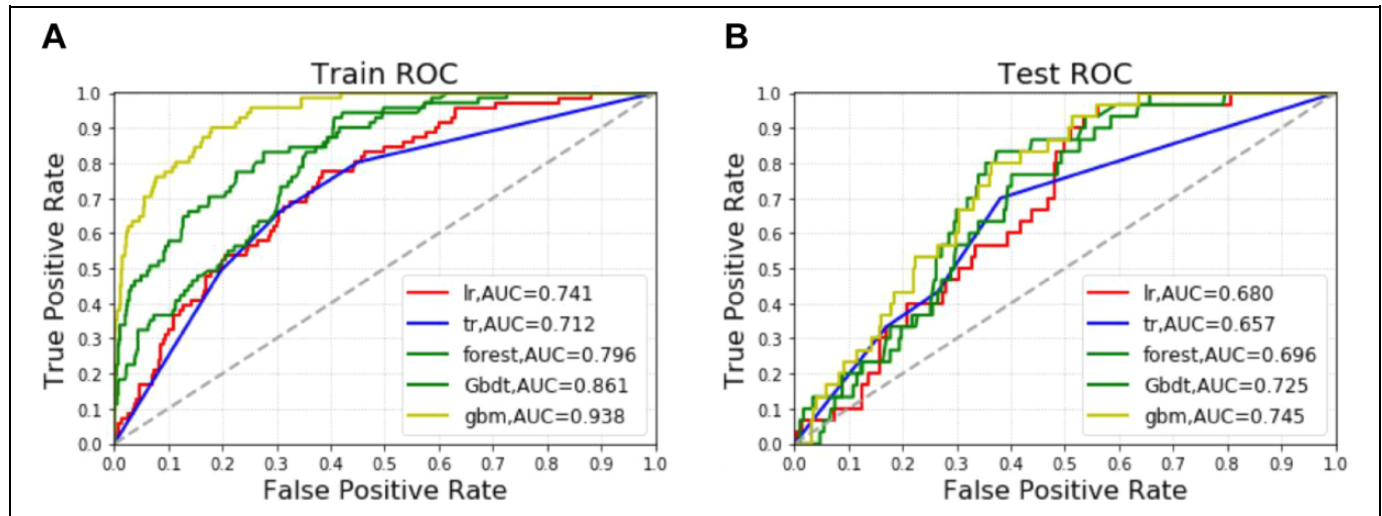
Correlation analysis showed that tumor size and depth of invasion were positively correlated with the recurrence of patients after gastric cancer surgery. Among them, tumor size and invasion depth were positively correlated with peritoneal metastasis, while body weight and BMI index were weakly correlated with peritoneal metastasis. (Figure 1) In addition, the results of the gbm (Light Gradient Boosting Machine) algorithm showed that the top 5 important factors were albumin, platelet count, depth of infiltration, preoperative hemoglobin and weight, respectively. (Figure 2)

Effect of postoperative recurrence model of gastric cancer patients in a training group: Among the 5 algorithm models, the accuracy of GradientBoosting and gbm was the highest (0.909), and the accuracy of the other 3 algorithm models was (0.906). The AUC values of the 5 algorithms are gbm(0.938), GradientBoosting(0.861), forest(0.796), Logistic(0.741) and DecisionTree(0.712) from high to low. Of the 5 algorithms, both GradientBoosting and gbm have the lowest MSE value of 0.091. (Figure 3A and Table 2)

The effect of postoperative recurrence model of gastric cancer patients in the test group: among the 5 algorithm models, the accuracy of the forest, DecisionTree and gbm was the highest 0.907, and the accuracy of the other 2 algorithm models was 0.904; AUC values ranged from high to low to gbm(0.745), GradientBoosting(0.725), forest(0.696), Logistic(0.680) and DecisionTree(0.657). Among the 5 algorithms, DecisionTree, gbm and forest had the lowest MSE value of 0.093. (Figure 3B and Table 2)

### Discussion

The peritoneum is a membrane-like structure covering the inner surface of the abdominopelvic wall and the surface of the viscera, which is mainly composed of mesothelial cells and a small amount of underlying connective tissue.<sup>13</sup> It is a predilection site for malignant tumor metastasis of the abdominopelvic cavity. The incidence of peritoneal metastasis in patients after radical gastrectomy for gastric cancer is 40% ~ 50%, which is the main type of recurrent metastasis after radical



**Figure 3.** Different machine learning algorithms predict the peritoneal metastasis in the training group(A) and test group(B). Note: gbm: Light Gradient Boosting Machine.

**Table 2.** Forecast Results for Training and Testing Group.

	Training			Testing		
	Accuracy	AUC	MSE	Accuracy	AUC	MSE
Logistic	0.906	0.741	0.094	0.904	0.680	0.096
DecisionTree	0.906	0.712	0.094	0.907	0.657	0.093
forest	0.906	0.796	0.094	0.907	0.696	0.093
GradientBoosting	0.909	0.861	0.091	0.904	0.725	0.096
gbm	0.909	0.938	0.091	0.907	0.745	0.093

Note: gbm: Light Gradient Boosting Machine.

gastrectomy for gastric cancer, and also the main cause of death. There was no obvious symptom in the early stage of peritoneal metastasis, but further development could lead to ascites, gastrointestinal and ureteral obstruction, hydronephrosis, etc. which seriously affected the quality of life and prognosis of patients.<sup>14</sup> Peritoneal metastasis of gastric cancer is often associated with a poor prognosis.<sup>15</sup> The results of this study show that the machine learning algorithm can better predict peritoneal metastasis in patients with gastric cancer, and the accuracy of the 5 algorithms is as high as 90%. In addition, the results of the gbm algorithm showed that the top 5 important factors were albumin, platelet count, depth of infiltration, preoperative hemoglobin and weight, respectively.

The high-risk factors for peritoneal metastasis of gastric cancer include TNM staging, extranodal infiltration, Borrmann type III ~ IV, Lauren type diffused and abdominal free cancer cells positive. The incidence of peritoneal metastasis was 25% in T3, T4, and N positive patients, and postoperative pathology showed that the incidence of peritoneal metastasis was 3.84 times higher in lymph node-positive patients than in negative patients, and the risk of peritoneal metastasis was higher in patients with extranodal metastases. Albumin improves prognosis in patients with peritoneal metastases from gastric

cancer.<sup>16</sup> Low pretreatment hemoglobin levels may reflect poor prognosis in patients with endometrial cancer.<sup>17</sup> This is similar to our findings, and albumin and preoperative hemoglobin are one of the top 5 important factors in peritoneal metastases in patients with gastric cancer.

In the report by Pawlik et al,<sup>18</sup> regarding BMI and gastric cancer, patients with underweight BMI <18.5 kg/m<sup>2</sup> had worse overall survival after gastric cancer resection than patients with BMI above 18.5. Low body mass index has been reported to be associated with more serious postoperative complications and poor prognosis compared with patients with normal body mass index.<sup>19</sup> Body mass index (BMI) may be a prognostic factor for diffuse gastric cancer in the peritoneum.<sup>20</sup> This is supported by our findings.

Recently, NLR has been reported as an important independent predictor of peritoneal metastasis in patients with advanced GC.<sup>21</sup> The predictive value of NLR for peritoneal metastasis during SL has been studied in early gastric cancer or lower esophageal cancer but has not been reported in advanced GC.<sup>22</sup> Studies<sup>23</sup> have also shown that high NLR is an important independent predictor of P/ CY-positive outcomes during SL in patients with advanced GC. Similarly, a meta-analysis<sup>24</sup> of 26 studies, including 13964 patients, showed that PLR was a poor prognostic factor for OS in patients with gastric cancer and colorectal cancer, hepatocellular carcinoma, ovarian cancer, and non-small cell lung cancer, and was not a poor prognostic factor for pancreatic cancer. Gunaldi et al.<sup>25</sup> studied the relationship between PLR and prognosis of gastric cancer by taking 160 as the cut-off value of PLR; The results showed that PLR was related to depth of invasion and stage of disease, and was not related to OS. Lian et al.<sup>26</sup> followed up 162 patients with gastric cancer surgery, and selected PLR = 208 as demarcation value. The tumor invasion degree was deep in patients with high PLR, the lymph node metastasis was large, the clinical stage was late, and the OS and DFS of

patients with high PLR were short. Our results also suggest that PLR and NLR are one of the important factors in peritoneal metastasis of gastric cancer.

There are still some limitations of this study that cannot be overlooked. This was a retrospective study. Therefore, tumor characteristics such as depth of invasion, lymphangitic invasion, and type of pathology were obtained after surgery. These parameters can be obtained pre-operatively by endoscopy. However, the results may not be accurate. In addition, we did not add prognostic analysis to the study due to incomplete data and partial loss of follow-up data. In addition, all these patients were from the same hospital. And in this study, T1, T2, T3 and T4 stages are not subdivided, which fails to compare the prediction performance of different stages. As this is a second retrospective analysis, we can only re-study the specific

classification of T when we carry out relevant prospective research in the future. Therefore, our findings still need to be validated through a large prospective multicenter study. Besides, the clinical-pathological factors analyzed were few, and Ki-67, C-reactive protein, carcinoembryonic antigen and CA199 could be combined to select more significant independent prognostic indicators in the future.

## Conclusion

This is the first attempt to study the use of machine learning to predict peritoneal metastases in patients with gastric cancer. We found that machine learning was a good predictor of peritoneal metastases in gastric cancer patients with an accuracy of up to 90%.

## Appendix A

**Table 1.** Functions, Packages, and Tuning Parameters in the Anaconda Software Used for Each Machine Learning Algorithm.

Algorithm	Classifier	Package	Tuning Parameters
Logistic regression	LogisticRegression	from sklearn.linear_model import LogisticRegression	Penalty = "l2," tol = 0.0001, C = 1, intercept_scaling = 1, max_iter = 100
DecisionTree	DecisionTreeClassifier	from sklearn.tree import DecisionTreeClassifier	splitter = "best," max_depth = 2, min_samples_split = 20, min_samples_leaf = 5, min_weight_fraction_leaf = 0.1
forest	RandomForestClassifier	from sklearn.ensemble import RandomForestClassifier	n_estimators = 10, max_depth = 3, min_samples_split = 70, min_samples_leaf = 6, random_state = 41
GradientBoosting	GradientBoostingClassifier	from sklearn.ensemble import GradientBoostingClassifier	learning_rate = 0.06, n_estimators = 50, max_depth = 2, random_state = 41
gbm	lgb.LGBMClassifier	lightgbm 2.2.0	learning_rate = 0.1, n_estimators = 30, max_depth = 3

Note: gbm:(Light Gradient Boosting Machine).

**Table 2.** Basic Characteristics of Training Group and Test Group.

	Training	Test	P-value
Number	756	324	
Age (years)	63.9 ± 11.1	63.3 ± 11.6	0.409
Height(cm)	165.0 ± 8.2	165.5 ± 7.7	0.359
Weight(kg)	58.7 ± 10.4	60.1 ± 10.7	0.051
BMI (kg/m <sup>2</sup> )	21.6 ± 3.0	21.9 ± 3.1	0.061
Tumor size (cm)	4.1 ± 2.2	3.8 ± 2.0	0.048
PLR	151.0 ± 71.5	150.4 ± 80.6	0.890
NLR	2.6 ± 1.5	2.6 ± 1.7	0.936
PREOPERATIVE HEMOGLOBIN	117.5 ± 23.7	118.5 ± 24.4	0.539
Platelet count	229.0 ± 70.2	221.9 ± 69.4	0.128
Albumin (g/L)	39.5 ± 5.2	40.0 ± 5.2	0.136
Neutrophil count	4.0 ± 3.0	4.2 ± 5.1	0.627
Lymphocyte count	1.8 ± 1.9	1.8 ± 1.9	0.865
Monocyte count	0.4 ± 0.3	0.4 ± 0.5	0.838
WBC count	6.2 ± 1.6	6.0 ± 1.5	0.022
Sex			0.034
Male	574 (75.9%)	265 (81.8%)	
Female	182 (24.1%)	59 (18.2%)	

(continued)

**Table 2.** (continued)

	Training	Test	P-value
ASA			0.885
I	47 (6.2%)	19 (5.9%)	
2	641 (84.8%)	273 (84.3%)	
3	68 (9.0%)	32 (9.9%)	
TNM			0.314
I	183 (24.2%)	76 (23.5%)	
II	56 (7.4%)	19 (5.9%)	
III	383 (50.7%)	182 (56.2%)	
IV	134 (17.7%)	47 (14.5%)	
Borrmann types			0.382
I	44 (5.8%)	19 (5.9%)	
2	652 (86.2%)	287 (88.6%)	
3	60 (7.9%)	18 (5.6%)	
Pathological type [n, (%)]			0.650
Ulcerative	101 (13.4%)	40 (12.3%)	
Nonulcerative	655 (86.6%)	284 (87.7%)	
Depth of invasion [n, (%)]			0.401
T1/T2	237 (31.3%)	110 (34.0%)	
T3/T4	519 (68.7%)	214 (66.0%)	

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## Authors' Contributions

All authors contributed to data analysis, drafting or revising the article, gave final approval of the version to be published, and agree to be accountable for all aspects of the work.

## Availability of Data and Material

Data is available at BioStudies database (<https://www.ebi.ac.uk/biostudies/studies?query=S-EPMC5383064>), accession numbers: S-EPMC5383064.

## Consent for Publication

All consent of the personal data were obtained from corresponding person.

## Declaration of Conflicting Interests

The author(s) declared no potential conflicts of interest with respect to the research, authorship, and/or publication of this article.


## Ethics Approval and Consent to participate

Data is available at BioStudies database (<https://www.ebi.ac.uk/biostudies/studies?query=S-EPMC5383064>), accession numbers: S-EPMC5383064. Our study did not require the approval of an ethics committee as it is a secondary analysis of BioStudies database of a public domain and of free access [28] [29]. This article does not contain any studies with human participants or animals performed by any of the authors.

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