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MINIREVIEWS

- 1892 Go-Ichi-Ni-San 2: A potential biomarker and therapeutic target in human cancers
Shan DD, Zheng QX, Chen Z
- 1903 Neoadjuvant therapy in resectable pancreatic cancer: A promising curative method to improve prognosis
Zhang HQ, Li J, Tan CL, Chen YH, Zheng ZJ, Liu XB

ORIGINAL ARTICLE**Basic Study**

- 1918 Transcriptional factor III A promotes colorectal cancer progression by upregulating cystatin A
Wang J, Tan Y, Jia QY, Tang FQ
- 1933 VCAN, expressed highly in hepatitis B virus-induced hepatocellular carcinoma, is a potential biomarker for immune checkpoint inhibitors
Wang MQ, Li YP, Xu M, Tian Y, Wu Y, Zhang X, Shi JJ, Dang SS, Jia XL
- 1949 Overexpression of ELL-associated factor 2 suppresses invasion, migration, and angiogenesis in colorectal cancer
Feng ML, Wu C, Zhang HJ, Zhou H, Jiao TW, Liu MY, Sun MJ
- 1968 Interleukin-34 promotes the proliferation and epithelial-mesenchymal transition of gastric cancer cells
Li CH, Chen ZM, Chen PF, Meng L, Sui WN, Ying SC, Xu AM, Han WX
- 1981 Cuproptosis-related long non-coding RNAs model that effectively predicts prognosis in hepatocellular carcinoma
Huang EM, Ma N, Ma T, Zhou JY, Yang WS, Liu CX, Hou ZH, Chen S, Zong Z, Zeng B, Li YR, Zhou TC

Retrospective Study

- 2004 Multi-slice spiral computed tomography in differential diagnosis of gastric stromal tumors and benign gastric polyps, and gastric stromal tumor risk stratification assessment
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- 2014 Predictive value of a serum tumor biomarkers scoring system for clinical stage II/III rectal cancer with neoadjuvant chemoradiotherapy
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- 2025 Role of sex on psychological distress, quality of life, and coping of patients with advanced colorectal and non-colorectal cancer
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- 2085** Correction to "Genome-wide CRISPR-Cas9 screening identifies that hypoxia-inducible factor-1 α -induced CBX8 transcription promotes pancreatic cancer progression *via* IRS1/AKT axis"

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AIMS AND SCOPE

The primary aim of *World Journal of Gastrointestinal Oncology* (*WJGO*, *World J Gastrointest Oncol*) is to provide scholars and readers from various fields of gastrointestinal oncology with a platform to publish high-quality basic and clinical research articles and communicate their research findings online.

WJGO mainly publishes articles reporting research results and findings obtained in the field of gastrointestinal oncology and covering a wide range of topics including liver cell adenoma, gastric neoplasms, appendiceal neoplasms, biliary tract neoplasms, hepatocellular carcinoma, pancreatic carcinoma, cecal neoplasms, colonic neoplasms, colorectal neoplasms, duodenal neoplasms, esophageal neoplasms, gallbladder neoplasms, *etc.*

INDEXING/ABSTRACTING

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Retrospective Study

Predictive value of a serum tumor biomarkers scoring system for clinical stage II/III rectal cancer with neoadjuvant chemoradiotherapy

Jie-Yi Zhao, Qing-Qing Tang, Yu-Ting Luo, Shu-Min Wang, Xiao-Rui Zhu, Xiao-Yu Wang

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Abstract

BACKGROUND

Multiple classes of molecular biomarkers have been studied as potential predictors for rectal cancer (RC) response. Carcinoembryonic antigen (CEA) is the most widely used blood-based marker of RC and has proven to be an effective predictive marker. Cancer antigen 19-9 (CA19-9) is another tumor biomarker used for RC diagnosis and postoperative monitoring, as well as monitoring of the therapeutic effect. Using a panel of tumor markers for RC outcome prediction is a practical approach.

AIM

To assess the predictive effect of pre-neoadjuvant chemoradiotherapy (NCRT) CEA and CA19-9 levels on the prognosis of stage II/III RC patients.

METHODS

CEA and CA19-9 levels were evaluated 1 wk before NCRT. According to the receiver operating characteristic curve analysis, the optimal cut-off point of CEA and CA19-9 levels for the prognosis were 3.55 and 19.01, respectively. The novel serum tumor biomarker (NSTB) scores were as follows: score 0: Pre-NCRT CEA < 3.55 and CA19-9 < 19.01; score 2: Pre-NCRT CEA > 3.55 and CA19-9 > 19.01; score 1: Other situations. Pathological information was recorded according to histopathological reports after the operation.

RESULTS

In the univariate analysis, pre-NCRT CEA < 3.55 [$P = 0.025$ for overall survival

(OS), $P = 0.019$ for disease-free survival (DFS)], pre-NCRT CA19-9 < 19.01 ($P = 0.014$ for OS, $P = 0.009$ for DFS), a lower NSTB score (0-1 *vs* 2, $P = 0.009$ for OS, $P = 0.005$ for DFS) could predict a better prognosis. However, in the multivariate analysis, only a lower NSTB score (0-1 *vs* 2; for OS, HR = 0.485, 95%CI: 0.251-0.940, $P = 0.032$; for DFS, HR = 0.453, 95%CI: 0.234-0.877, $P = 0.019$) and higher pathological grade, node and metastasis stage (0-I *vs* II-III; for OS, HR = 0.363, 95%CI: 0.158-0.837, $P = 0.017$; for DFS, HR = 0.342, 95%CI: 0.149-0.786, $P = 0.012$) were independent predictive factors.

CONCLUSION

The combination of post-NCRT CEA and CA19-9 was a predictive factor for clinical stage II/III RC patients receiving NCRT, and the combined index had a stronger predictive effect.

Key Words: Rectal cancer; Neoadjuvant chemoradiotherapy; Scoring system; Carcinoembryonic antigen; Carbohydrate antigen 19-9; Predictive

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Core Tip: Tumor microenvironment (TME) combined with neoadjuvant chemotherapy (NCRT) is the standard treatment for resectable stage II/III rectal cancer (RC). Multiple classes of molecular biomarkers have been studied as potential predictors for RC response but there is no sufficient evidence for any of them to be introduced into clinical practice. By retrospectively evaluating clinical stage II/III RC patients undergoing NCRT followed by standard TME, we found that the combination of NCRT carcinoembryonic antigen and carbohydrate antigen 19-9 levels could be a prognostic predictor for clinical stage II/III RC patients receiving NCRT, and the combined indexes had a stronger predictive effect than the index alone.

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INTRODUCTION

In the United States, tumor microenvironment (TME) combined with neoadjuvant chemotherapy (NCRT) is the standard treatment for resectable stage II/III rectal cancer (RC)[1-3]. Although numerous studies have shown that NCRT can reduce the rate of local recurrence, it is difficult to improve overall survival (OS)[4-6]. Multiple classes of molecular biomarkers have been studied as potential predictors for RC response but there is no sufficient evidence for any of them to be introduced into clinical practice [7]. Moreover, additional systematic chemotherapy could increase the toxicity [8,9]. Therefore, it is critical to identify predictive factors for clinical stage II/III patients and give additional chemotherapy or more aggressive treatment strategies.

Pathological indicators are generally considered to be the most effective predictive factors[10,11]. Unfortunately, pathological characteristics are difficult to obtain and quantitate and are usually affected by the operation and specimen-processing quality[8,9]. Moreover, the pathological indicators, which can only be obtained after surgery, do not assist in judging whether patients need additional chemotherapy before undergoing NCRT or surgery.

A glycoprotein, carcinoembryonic antigen (CEA), is the most widely used blood-based marker of RC and has proven to be an effective predictive marker[12-14]. According to You *et al*[15], the increment in postoperative serum CEA levels (CEA < 5 *vs* > 5) was an independent predictor of a poor prognosis. However, the major problem with the use of CEA as a marker of RC is its association with other types of cancer and benign diseases (inflammatory bowel disease)[16-18]. Cancer antigen 19-9 (CA19-9) is another tumor biomarker used for RC diagnosis and postoperative monitoring, as well as monitoring of the therapeutic effect[19,20]. Due to the highly heterogeneous nature of RC, a single tumor marker is unlikely to become a stand-alone predictive factor. Using a panel of tumor markers for RC outcome prediction is a practical approach.

In this study, we analyzed the predictive value of the combination of pre-NCRT serum tumor markers (CEA and CA19-9) in clinical stage II/III RC patients.

MATERIALS AND METHODS

Patients screening

We retrospectively evaluated clinical stage II/III RC patients undergoing NCRT followed by standard TME in our hospital from February 2011 to August 2020. We included the following categories of patients: (1) Patients receiving preoperative NCRT; (2) patients with colorectal adenocarcinoma confirmed by pathological biopsy; (3) patients whose serum CEA and CA19-9 levels were measured within one week before NCRT; and (4) patients undergoing NCRT followed by standard TME. We excluded the following categories of patients: (1) Patients with distal metastasis; (2) patients with other concomitant tumors; (3) patients with insufficient blood, clinicopathological, or follow-up data; and (4) patients with unresectable RC. The patient-screening flowchart is shown in [Figure 1](#).

This retrospective study was approved by the ethics committee of our hospital. The requirement for patients' informed consent was waived due to the retrospective nature of the study.

Treatment and follow-up of patients

All patients in this study received NCRT followed by standard TME. Their CEA and CA19-9 levels were evaluated within 1 wk pre-NCRT. Pathological, node and metastasis (TNM) stages and histological grades were noted according to histopathological reports. The receiver operating characteristic (ROC) curve was adopted to determine the best cut-off values of pre-NCRT CEA and CA19-9 levels for predicting OS. The novel serum tumor biomarker (NSTB) scores were as follows: score 0: Pre-NCRT CEA < 3.55 and CA19-9 < 19.01; score 2: Pre-NCRT CEA > 3.55 and CA19-9 > 19.01; score 1: Pre-NCRT CEA < 3.55 and CA19-9 > 19.01 or pre-NCRT CEA > 3.55 and CA19-9 < 19.01.

Postoperative follow-up was performed according to the National Comprehensive Cancer Network guidelines[13]. Generally, patients were followed up clinically and radiographically at three-month intervals in the first 2 years after surgery, then every 6 mo for 3 years, and annually thereafter[13]. Follow-up data were obtained from medical records, telephone follow-ups, out-patient clinics, or visits.

OS was defined as the survival time until death by any reason[21]. DFS was defined as the time-lapse between surgery and either RC recurrence or death[22]. Patients lost to follow-up or still alive at the final follow-up were included in the analysis as censored data[21].

Statistical analysis

Data were analyzed using SPSS for Windows (IBM Corp. Released 2013. IBM SPSS Statistics for Windows, Version 22.0. Armonk, NY: IBM Corp.). Continuous data were described in terms of the median and interquartile range (IQR) whereas categorical variables were described in terms of frequencies and percentages. Significant parameters identified in the univariate analysis ($P < 0.05$) were entered into the multivariate Cox regression analysis to determine independent predictive factors[23, 24]. All statistical tests were two-sided, and a P value of < 0.05 was considered statistically significant [25].

In general, pathological characteristics have the strongest predictive value for patient outcomes[21]. To compare the predictive effect of the NSTB score, several pathological indicators were included. To prevent the effects of pre-NCRT CEA and CA19-9 levels on the NSTB score, two models, one including and the other excluding the NSTB score in the multivariate analysis, were established.

RESULTS

Patient characteristics

Eighty-seven (36.7%) patients were female and 150 (63.3%) were male. The distribution of the patients according to pathological evaluation was as follows: vascular invasion was detected in 13 (5.5%) patients, lymphatic invasion in 13 (5.5%) patients, perineural invasion in 41 (17.3%) patients, and circumferential resection margin (CRM) positivity in 8 (3.8%) patients. Regarding the pathological TNM classification, 45 (19.0%) patients were in stage 0, 57 (24.1%) were in stage I, 72 (30.4%) were in stage II, and 63 (26.6%) were in stage III ([Table 1](#)). A total of 118 (49.8%) patients were in pT stage 0-2 while 119 (50.2%) were in pT stage 3-4. Sixty (25.3%) patients had pN metastases while 177 (74.7%) did not have pN metastases. The median (IQR) level of pre-NCRT CEA was 4.15 (2.18-10.07) and that of pre-NCRT CA19-9 was 13.56 (7.80-25.40).

During follow-up, 9 (3.8%) patients were lost to follow-up and 36 (15.2%) developed cancer recurrence and died.

Kaplan-Meier curves stratified by pre-NCRT CEA, CA19-9, and the NSTB score

ROC curves identified the optimal cut-off for survival prediction by pre-NCRT CEA and CA19-9 were 3.55 and 19.01, respectively. They divided patients into different groups. [Figures 2-4A](#) show the OS of included patients stratified by pre-NCRT CEA, CA19-9, and the NSTB score, respectively, and [Figures 2-4B](#) show their DFS stratified by the same parameters. According to the Kaplan-Meier curves, increased

Table 1 Clinicopathological characteristics

Features	Median (IQR)
Pre-NCRT CEA	4.15 (2.18-10.07)
Pre-NCRT CA19-9	13.56 (7.80-25.40)
Age in yr	57.0 (50.0-66.5)
Sex	
Male	150
Female	87
Pathological T stage	
T0-2	118
T3-4	119
Pathological N stage	
N0	177
N+	60
Pathological TNM stage	
0	45
I	57
II	72
III	63
Pathological vascular invasion	
Yes	13
No	224
Pathological lymphatic invasion	
Yes	13
No	224
Pathological perineural invasion	
Yes	41
No	196
Pathological CRM	
Positive	8
Negative	229

NCRT: Neoadjuvant chemoradiotherapy; CEA: Carcinoembryonic antigen; CA19-9: Cancer antigen 19-9; TNM: Tumor, node and metastasis; CRM: Circumferential resection margin; IQR: Interquartile range.

pre-NCRT CEA and CA19-9 levels and higher NSTB scores were all associated with decreased OS and DFS.

Cox regression analysis of factors affecting the prognosis

Possible clinicopathological parameters that may predict patient outcome were reviewed. In the univariate analysis, pre-NCRT CEA > 3.55, pre-CA19-9 > 19.01, a higher pathological TNM stage, and a higher NSTB score were significantly associated with decreased OS (Table 2) and DFS (Table 3).

In the multivariate analysis of OS (Table 4), only a lower pathological TNM stage (stage 0-I vs II-III, HR = 0.363, 95%CI: 0.158-0.837, $P = 0.017$) and the NSTB score (score 0-1 vs 2, HR = 0.485, 95%CI: 0.251-0.940, $P = 0.032$) were significant predictors of a better outcome while pre-NCRT CEA < 3.55 (HR = 0.529, 95%CI: 0.23-1.205, $P = 0.130$) and CA19-9 < 19.01 (HR = 0.604, 95%CI: 0.300-1.215, $P = 0.158$) were not. In the multivariate analysis of DFS (Table 5), a lower pathological TNM stage (stage 0-I vs II-III, HR = 0.342, 95%CI: 0.149-0.786, $P = 0.012$) and the NSTB score (score 0-1 vs 2, HR = 0.453, 95%CI: 0.234-0.877, $P = 0.019$) could also predict a better outcome while pre-NCRT CEA < 3.55 (HR = 0.521, 95%CI:

Table 2 Univariate analysis of factors affecting the overall survival

Characteristics	HR (95%CI)	P value
Pre-NCRT CEA (< 3.55/> 3.55)	0.407 (0.185-0.893)	0.025
Pre-NCRT CA19-9 (< 19.01/> 19.01)	0.437 (0.225-0.849)	0.014
Sex (male/female)	0.478 (0.218-1.049)	0.066
Pathological TNM stage (0-I/II-III)	0.321 (0.141-0.732)	0.007
Pathological vascular invasion (absent/present)	0.556 (0.170-1.821)	0.332
Pathological lymphatic invasion (absent/present)	0.400 (0.141-1.136)	0.085
Pathological perineural invasion (absent/present)	0.534 (0.250-1.141)	0.105
Pathological CRM (negative/positive)	0.826 (0.198-3.449)	0.793
NSTB score (0-1/2)	0.416 (0.217-0.800)	0.009

NCRT: Neoadjuvant chemoradiotherapy; CEA: Carcinoembryonic antigen; CA19-9: Cancer antigen 19-9; TNM: Tumor, node and metastasis; CRM: Circumferential resection margin; NSTB: Novel serum tumor biomarker score.

Table 3 Univariate analysis of factors affecting disease-free survival

Characteristic	HR (95%CI)	P value
Pre-NCRT CEA (< 3.55/> 3.55)	0.391 (0.178-0.859)	0.019
Pre-NCRT CA19-9 (< 19.01/> 19.01)	0.413 (0.213-0.802)	0.009
Sex (male/female)	0.466 (0.213-1.023)	0.057
Pathological TNM stage (0-I/II-III)	0.302 (0.132-0.690)	0.005
Pathological vascular invasion (absent/present)	0.571 (0.175-1.863)	0.353
Pathological lymphatic invasion (absent/present)	0.435 (0.154-1.231)	0.117
Pathological perineural invasion (absent/present)	0.595 (0.279-1.265)	0.177
Pathological CRM (negative/positive)	0.657 (0.158-2.738)	0.564
NSTB score (0-1/2)	0.391 (0.203-0.751)	0.005

NCRT: Neoadjuvant chemoradiotherapy; CEA: Carcinoembryonic antigen; CA19-9: Cancer antigen 19-9; TNM: Tumor, node and metastasis; CRM: Circumferential resection margin; NSTB: Novel serum tumor biomarker score.

0.226-1.162, $P = 0.109$) and CA19-9 < 19.01 (HR = 0.570, 95%CI: 0.284-1.141, $P = 0.112$) could not. The nomogram of OS (Figure 5) and DFS (Figure 6) shows the precise prognosis.

DISCUSSION

Our data showed that the combination of pre-NCRT tumor markers had better predictive value than a single marker. Although univariate analyses demonstrated that lower pre-NCRT CEA and CA19-9 levels were potential indicators of a better prognosis, the multivariate analysis proved that only the NSTB score and pathological TNM stage could independently determine the prognosis. In general, pathological indicators had a more robust predictive value than other indicators in determining the prognosis[8]; however, the multivariate analysis revealed that the NSTB score could predict outcomes better than pathological characteristics of lymphatic invasion, vascular invasion, nerve infiltration, and CRM invasion. Thus, we propose that the NSTB score should be used to guide the treatment and determine the prognosis of patients with RC of clinical stage II/III.

Pathological findings were generally recognized as the most effective indicators to predict the prognosis[8]. A previous study revealed that lymphatic invasion, perineural invasion, vascular invasion, CRM invasion, LN metastasis, and a higher tumor invasion stage can predict a worse outcome[4]. However, pathological characteristics were difficult to identify as they are usually affected by the quality of surgery and specimen-processing, and their analysis is significantly subjective and difficult to quantitate[15]. Moreover, pathological indicators could only be obtained after surgery, which means

Table 4 Multivariate analysis of factors affecting the overall survival

Characteristic	Multivariate analysis			
	Model 1		Model 2	
	HR (95%CI)	P value	HR (95%CI)	P value
Pre-NCRT CEA (< 3.55/> 3.55)	0.529 (0.232-1.205)	0.130		
Pre-NCRT CA19-9 (< 19.01/> 19.01)	0.604 (0.300-1.215)	0.158		
Pathological TNM stage (0-I/II-III)	0.373 (0.162-0.859)	0.020	0.363 (0.158-0.837)	0.017
NSTB score (0-1/2)			0.485 (0.251-0.940)	0.032

Model 1: Including pre-neoadjuvant chemoradiotherapy (NCRT) carcinoembryonic antigen (CEA) and cancer antigen 19-9 (CA19-9) into multivariate analysis, not including novel serum tumor biomarker (NSTB) score; Model 2: Including NSTB score into multivariate analysis, not including pre-NCRT CEA and CA19-9. NCRT: Neoadjuvant chemoradiotherapy; CEA: Carcinoembryonic antigen; CA19-9: Cancer antigen 19-9; NSTB: Novel serum tumor biomarker score; TNM: Tumor, node and metastasis.

Table 5 Multivariate analysis of factors affecting the disease-free survival

Characteristic	Multivariate analysis			
	Model 1		Model 2	
	HR (95%CI)	P value	HR (95%CI)	P value
Pre-NCRT CEA (< 3.55/> 3.55)	0.512 (0.226-1.162)	0.109		
Pre-NCRT CA19-9 (< 19.01/> 19.01)	0.570 (0.284-1.141)	0.112		
Pathological TNM stage (0-I/II-III)	0.350 (0.152-0.806)	0.014	0.342 (0.149-0.786)	0.012
NSTB score (0-1/2)			0.453 (0.234-0.877)	0.019

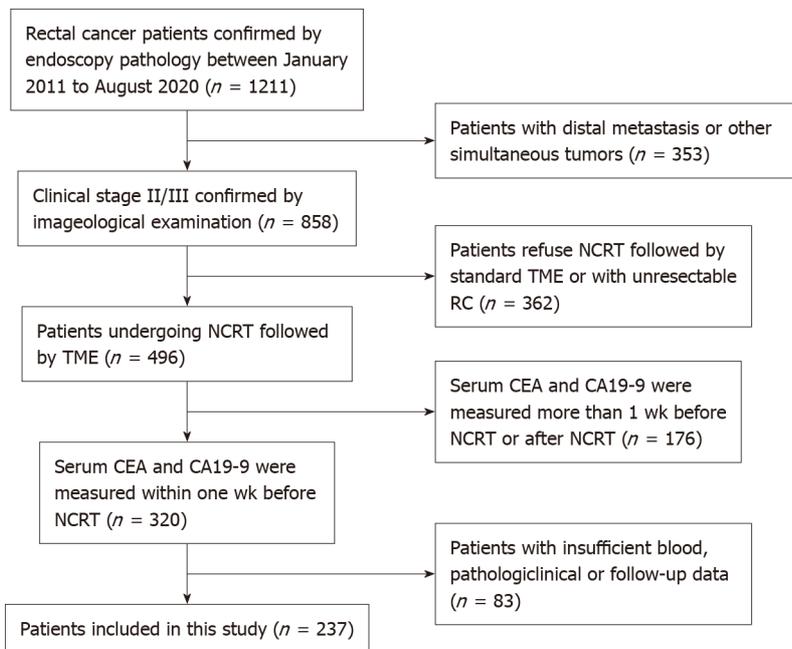
Model 1: Including pre-neoadjuvant chemoradiotherapy (NCRT) carcinoembryonic antigen (CEA) and cancer antigen 19-9 (CA19-9) into multivariate analysis, not including novel serum tumor biomarker (NSTB) score; Model 2: Including NSTB score into multivariate analysis, not including pre-NCRT CEA and CA19-9. NCRT: Neoadjuvant chemoradiotherapy; CEA: Carcinoembryonic antigen; CA19-9: Cancer antigen 19-9; NSTB: Novel serum tumor biomarker score; TNM: Tumor, node and metastasis.

that it was impossible to judge whether patients needed additional chemotherapy before undergoing NCRT or surgery. Moreover, the NSTB score could be obtained before NCRT and surgery.

Some molecules or proteins can determine a patient's prognosis. Lin *et al*[26] reported that the high expression of EphA4 served as an independent adverse predictor for DFS. Rödel *et al*[27] found that an increase in survivin levels was a significant risk factor for local recurrence and decreased DFS. Hiyoshi *et al*[28] demonstrated that serum miR-143 was a non-invasive and novel predictive marker for locally advanced rectal cancer (LARC) patients. Unfortunately, all of these molecular or protein markers had the following disadvantages: First, the detection cost of these markers was high, which increased the economic burden for patients; second, these novel markers could only be detected in large medical centers, which made them difficult to be used widely in clinical practice; finally, these new indicators lack uniform standards, and the test results may vary a lot in different medical centers. CEA and CA19-9 levels are widely used clinically because they are cheap, convenient to detect, and have uniform standards in different hospitals.

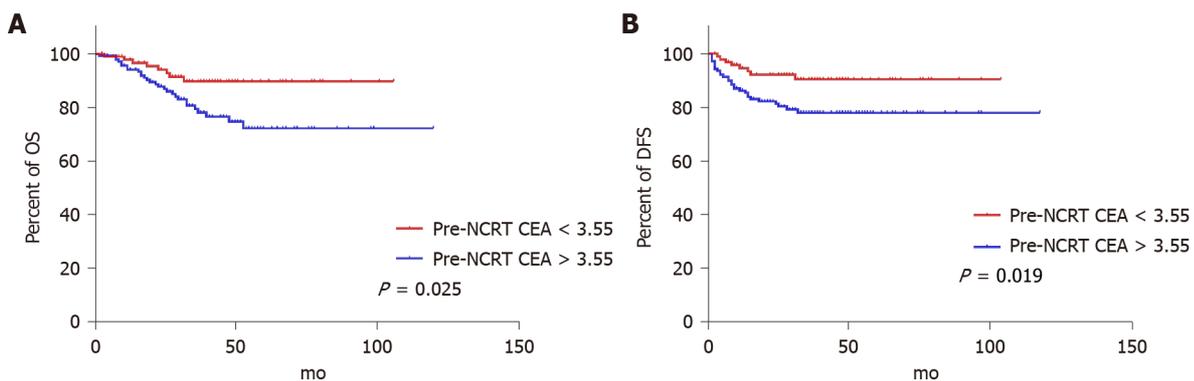
CEA is currently one of the primary markers for the diagnosis and follow-up of RC[2,18,19]. We found that lower CEA levels could predict a better prognosis in a univariate analysis. CA19-9 has shown great value for the differential diagnosis of malignant tumors and disease monitoring and evaluation [19]. Compared to either CA19-9 or CEA alone, an essential advantage of the combination was that it could reduce the interference of other factors and increase the predictive effectiveness. Although some studies also focused on the influence of CEA and CA19-9 levels on the prognosis, the two markers were studied separately[2,18-20,29]. Consequently, they failed to identify CEA and CA19-9 as predictive factors, which was similar to our findings. However, the predictive value increased significantly and was even stronger than that of several pathological factors when they were combined.

Our study had a few strengths: Firstly, to the best of our knowledge, this is the first study that combined CEA and CA19-9 to evaluate the prognosis of clinical stage II/III patients undergoing NCRT. Secondly, we adopted an ROC curve to determine the cut-off point of CEA and CA19-9 instead of just evaluating whether they were higher than the normal values, which optimized the efficiency of the OS prediction. Ultimately, the NSTB score was cheap and easily accessible before treatment.



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Figure 1 Patient-screening flowchart. NCRT: Neoadjuvant chemoradiotherapy; TME: Tumor microenvironment; RC: Rectal cancer; CEA: Carcinoembryonic antigen; CA19-9: Cancer antigen 19-9.



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Figure 2 Overall survival curves and disease-free survival curves stratified by pre-neoadjuvant chemoradiotherapy carcinoembryonic antigen levels. A: Overall survival curves stratified by pre-neoadjuvant chemoradiotherapy carcinoembryonic antigen (CEA) levels; B: Disease-free survival curves stratified by pre-neoadjuvant chemoradiotherapy CEA levels. NCRT: Neoadjuvant chemoradiotherapy; CEA: Carcinoembryonic antigen; OS: Overall survival; DFS: Disease-free survival.

Our study also had some shortcomings. First, this was a retrospective study conducted in a single medical center. Second, the cut-off points of pre-NCRT CEA and CA19-9 levels in our center may not always be reproducible in other centers.

CONCLUSION

This study established a NSTB score by combining pre-NCRT CEA and CA19-9 levels. The NSTB score can independently predict the prognosis of patients with LARC of clinical stage II/III who underwent NCRT. Its predictive value was stronger than that of either marker alone, and even some pathologic characteristics.

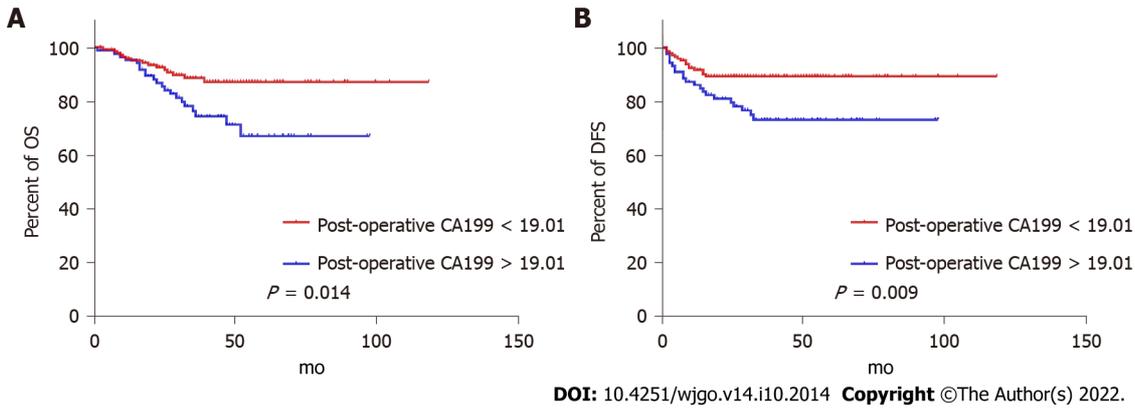


Figure 3 Overall survival curves and disease-free survival curves stratified by pre-neoadjuvant chemoradiotherapy cancer antigen 19-9 levels. A: Overall survival curves stratified by pre-neoadjuvant chemoradiotherapy cancer antigen 19-9 (CA19-9) levels; B: Disease-free survival curves stratified by pre-neoadjuvant chemoradiotherapy CA19-9 levels. CA19-9: Cancer antigen 19-9; OS: Overall survival; DFS: Disease-free survival.

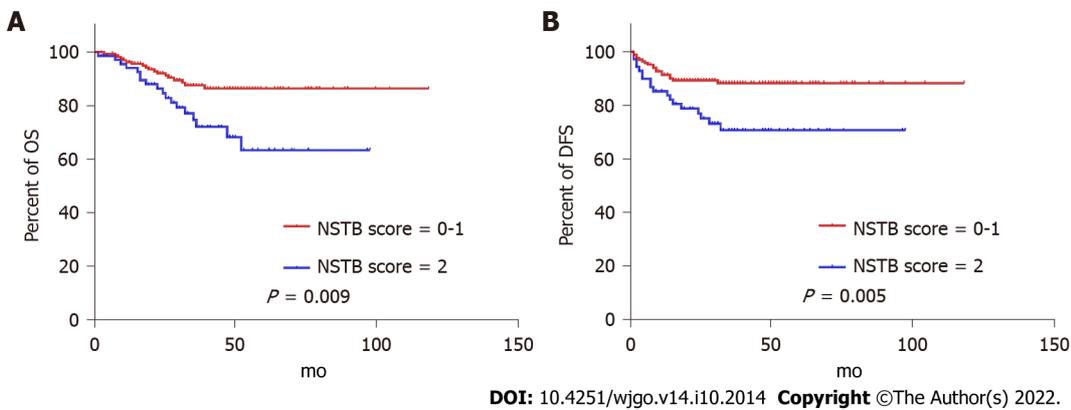


Figure 4 Overall survival curves and disease-free survival curves stratified by novel serum tumor biomarker scores. A: Overall survival curves stratified by novel serum tumor biomarker scores; B: Disease-free survival curves stratified by novel serum tumor biomarker scores. NSTB: Novel serum tumor biomarker; OS: Overall survival; DFS: Disease-free survival.

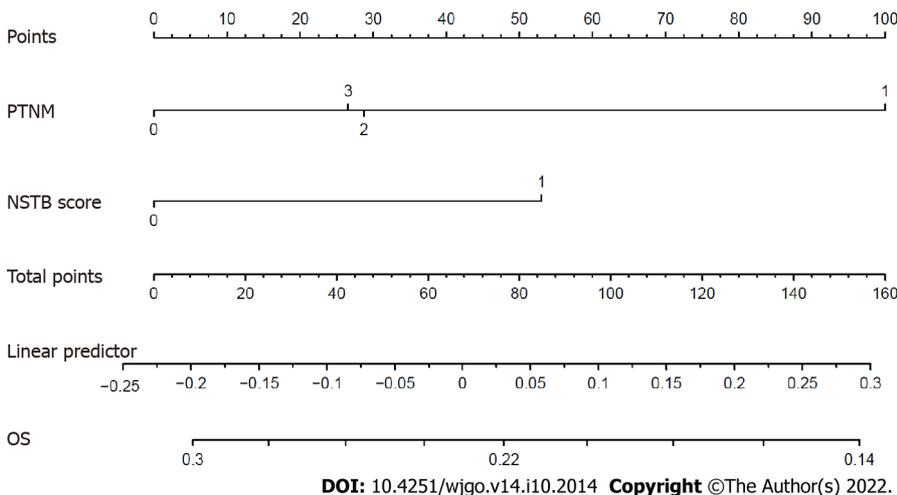


Figure 5 Predictive nomogram predicting overall survival in clinical stage II/III RC patients undergoing NCRT. PTNM: Pathological tumor-node-metastasis; NSTB: Novel serum tumor biomarker; OS: Overall survival.

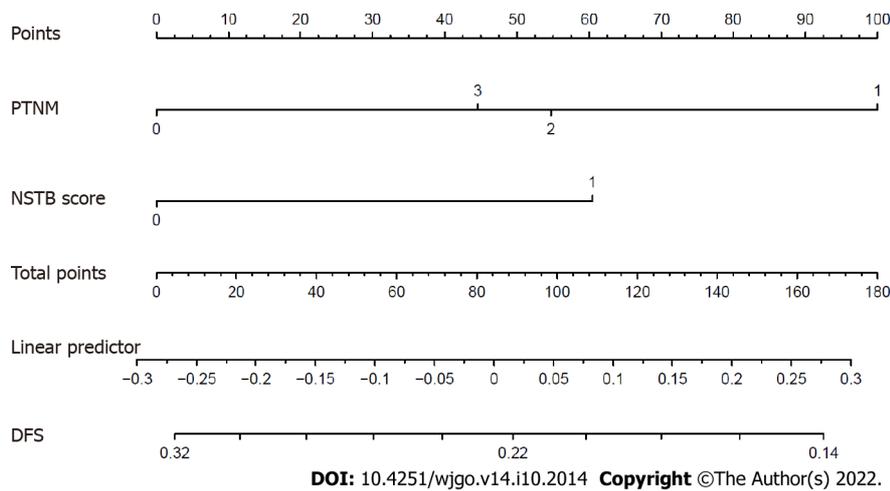


Figure 6 Predictive nomogram predicting disease-free survival in clinical stage II/III RC patients undergoing NCRT. PTNM: Pathological tumor-node-metastasis; NSTB: Novel serum tumor biomarker; DFS: Disease-free survival.

ARTICLE HIGHLIGHTS

Research background

Multiple classes of molecular biomarkers were studied as potential predictors for rectal cancer (RC) response but there was no sufficient evidence for any of them to be introduced into clinical practice.

Research motivation

To assess the predictive effect of pre-neoadjuvant chemoradiotherapy (NCRT) carcinoembryonic antigen (CEA) and cancer antigen 19-9 (CA19-9) levels on the prognosis of stage II/III RC patients.

Research objectives

The objective of this study is to establish a novel serum tumor biomarker score by combining pre-NCRT CEA and CA19-9 levels. The novel serum tumor biomarker (NSTB) score may predict the prognosis of patients with locally advanced rectal cancer (LARC) of clinical stage II/III who underwent NCRT independently.

Research methods

A total of 237 patients were included in this study. CEA and CA 19-9 levels were evaluated 1 wk before NCRT. The NSTB score was as follows: score 0: pre-NCRT CEA < 3.55 and CA19-9 < 19.01; score 2: pre-NCRT CEA > 3.55 and CA19-9 > 19.01; score 1: other situations. Pathological information was recorded according to histopathological reports after the operation.

Research results

In the univariate analysis, pre-NCRT CEA < 3.55, pre-NCRT CA19-9 < 19.01, and a lower NSTB score could predict a better prognosis. However, in the multivariate analysis, only a lower NSTB score and higher pathological tumor-node-metastasis (TNM) stage were independent predictive factors.

Research conclusions

We established a novel serum tumor biomarker score by combining pre-NCRT CEA and CA19-9 levels. The NSTB score can independently predict the prognosis of patients with LARC of clinical stage II/III who underwent NCRT.

Research perspectives

More accurate prediction models need to be established by studies with a larger number of patients.

FOOTNOTES

Author contributions: Zhao JY and Wang XY designed the research and wrote the manuscript; Tang QQ and Luo YT analyzed the data; Wang SM and Zhu XR performed data extraction; all authors have read and approved the final version.

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