Preoperative assessment of lymph node metastasis in clinically node-negative rectal cancer patients based on a nomogram consisting of five clinical factors

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\textbf{Background:} Currently, reliable approaches for accurate assessment of lymph node metastases (LNM), which is an important indication of preoperative chemoradiotherapy (CRT), are not available for clinically node-negative rectal cancer patients. This study aims to identify clinical factors associated with LNM and to establish a nomogram for LNM prediction in clinically node-negative rectal cancer patients.

\textbf{Methods:} The least absolute shrinkage and selection operator (LASSO) aggression and multivariate logistic regression analyses were applied to identify clinical factors associated with LNM. A nomogram was established to predict the probability of LNM in clinically node-negative rectal cancer patients based on the multivariate logistic regression model.

\textbf{Results:} Six potential risk factors were selected on the basis of LASSO aggression analysis, and five of them were identified as independent risk factors for LNM based on multivariate analysis, including MRI-reported tumor location, clinical T classification, MRI-reported tumor diameter, white blood cell count (WBC), and preoperative elevated tumor markers. A nomogram consisting of the five clinical factors was established and showed good discrimination. Decision curve analysis demonstrated that the established nomogram was reliable and accurate for LNM prediction in clinically node-negative rectal cancer patients.

\textbf{Conclusions:} A nomogram based on five clinical factors, including MRI-reported tumor location, clinical T classification, MRI-reported tumor diameter, WBC, and preoperative elevated tumor markers, are useful for assessing LNM in clinically node-negative rectal cancer patients, which is important for preoperative CRT regimens.

\textbf{Keywords:} Lymph node metastasis (LNM); nomogram; rectal cancer; risk factor

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Introduction

Colorectal cancer (CRC) is the fourth leading cause of cancer-related death worldwide, and approximately 30% of CRC have been identified in the rectum (1-3). While early screenings can significantly reduce cancer morbidity (4), many cancers are diagnosed at middle or late stage, which led to a high mortality and recurrence rates. For locally advanced rectal cancer, chemoradiotherapy (CRT) prior to surgery has been established as the standard of care (5). A great amount of evidence showed that preoperative CRT significantly prevents the progress of cancers and improves overall survival of cancer patients, compared with surgery alone (6-13).

Currently, locally advanced rectal cancer is a widely accepted indication of preoperative CRT for rectal cancers (9,14). Therefore, accurate preoperative staging of rectal cancer is critically important for CRC treatment. Nowadays, imaging methods including computed tomography (CT) and magnetic resonance imaging (MRI) are the main approaches to evaluate preoperative staging of CRC. MRI, due to excellent soft tissue contrast, is powerful for the identification of mesorectal node involvement as well as assessment of the circumferential resection margin (15-20). Compared to CT, which has approximately 60% of accuracy of the assessment of lymph node involvement (21), MRI showed 80% of accuracy in determining lymph node involvement in CRC patients (15,22). Therefore, one fifth of RC patients with lymph node metastases (LNM) are still underdiagnosed as clinically node-negative CRC and may miss the opportunity of preoperative CRT even MRI is used for preoperative staging of CRC.

Therefore, preoperative imaging alone is insufficient for accurate evaluation of lymph node involvement in CRC patients. New tools and evaluation system are highlighted to improve the assessment of lymph node involvement in CRC patients. Numerous factors have been proposed for the assessment of LNM in previous study, such as depth of submucosal invasion, lymphovascular invasion, high-grade mucin production, signet ring features, T-stage, and poorly-differentiated adenocarcinoma (23-29). However, most of these risk factors were based on the results from resected specimen, which were obtained postoperatively and may be inappropriate for preoperative staging of CRC.

In this study, we analyzed a number of preoperatively clinical factors, which were selected to develop a nomogram for preoperative prediction of LNM in clinically lymph-node negative rectal cancer patients. Our results are useful for physicians to screen CRC patients for preoperative CRT.

Methods

Patients

This study was approved by the Institutional Review Board (IRB) of The Sixth Affiliated Hospital of Sun Yat-sen University. The training cohort of this study includes 434 clinically lymph node-negative rectal cancer patients who were hospitalized at the Sixth Affiliated Hospital of Sun Yat-sen University (Guangzhou, China) between January 2010 and December 2014. The independent validation cohort includes 165 clinically lymph node-negative rectal cancer patients hospitalized at the same hospital between January 2015 and December 2016. Both the training and validation cohorts use the same inclusion and exclusion criteria (see below). Demographics and clinic-pathological variables were prospectively maintained in the CRC Database. Both paper charts and electronic medical records were carefully reviewed.

Inclusion and exclusion criteria

The inclusion criteria were: (I) patients with rectal cancer and hospitalized at the Sixth Affiliated Hospital of Sun Yat-sen University; (II) patients with clinically node-negative RC; (III) patients with pathological examination of a minimum of 12 lymph nodes. The exclusion criteria included: (I) patients with colon cancer; (II) patients without pre-therapy MRI images; (III) patients received preoperative CRT; (IV) patients with familiar adenomatous polyposis (FAP) or inflammatory bowel disease (IBD); (V) patients had incomplete clinical data.

Variables

Clinically node negative is defined as the short-axis diameter of lymph node <5 mm based on pre-therapy MRI images. Demographic and clinic-pathological variables include general information, age at the time of surgery, race, body mass index (BMI), preoperative total protein (<60 g/L vs. ≥ 60 g/L), preoperative albumin (<35 g/L vs. ≥35 g/L), preoperative hemoglobin (<110 g/L vs. ≥110 g/L), elevated PLT (>300x10^9/L), elevated WBC (>10x10^9/L), elevated CEA (>5 ng/mL), elevated CA199 (>37 U/mL), MRI-based distance between tumor and anal verge, MRI-based tumor diameter, clinical T classification, clinical N classification,
pathological T classification, pathological N classification, pathological M classification, and pathological TNM stage.

Statistical analysis

Statistical analysis was conducted using the R software (version 3.0.1; http://www.Rproject.org). The packages in R that were used in this study are reported in the data supplement. Descriptive statistics were computed for all variables. The least absolute shrinkage and selection operator (LASSO) method was used to identify predictive factors from the primary data set, weighted by their respective coefficients. Multivariate analyses of selected suboptimal risk factors associated with LNM were conducted using the logistic regression analysis. Bidirectional elimination was done to fit regression models based on the lowest Akaike information criterion. A prediction nomogram was established based on multivariate logistic regression analysis. Decision curve analysis was conducted to determine the clinical usefulness of the nomogram by quantifying the net benefits at different threshold probabilities. P values less than 0.05 were considered statistically significant.

Results

Patient characteristics

A total of 599 clinically lymph node-negative rectal cancer patients were enrolled in this study, including 511 patients (85.3%) underwent laparoscopic surgery and 88 (14.7%) underwent open surgery. The training cohort included 434 patients while the independent validation cohort included 165 patients. LNM was found in 24.2% and 30.9% of the training and validation cohorts, respectively (P=0.094). Preoperative clinical factors including age, gender, BMI, CEA, CA199, total protein, albumin, hemoglobin, PLT, WBC, MRI-based tumor location, MRI-based tumor diameter, and clinical T classification were shown in Table 1.

Risk factors for LNM

Risk factors for LNM in clinically node-negative rectal cancer patients were identified using multivariate logistic regression models of LASSO method. Of 13 clinical factors, six potential predictors were selected from 434 patients in the training cohort, based on nonzero coefficients in the LASSO logistic regression model (Figure 1). In order to optimize the predictive model, the six potential predictors were further evaluated using multivariate logistic regression model. The multivariate logistic regression analysis revealed that MRI-based tumor location (P=0.001), clinical T classification (P=0.001), MRI-based tumor diameter (P<0.001), preoperative WBC (P=0.048) and preoperative elevated tumor markers (P=0.005) were independent LNM risk factors for clinically node-negative rectal patients (Figure 2).

Development and validation of an individualized prediction model

A nomogram based on the five independent risk factors was established (Figure 3). When using, a vertical line is drawn for each variable to see their respective score. Each score was added together to get the total score, which determines the probability of LNM. For example, a patient with T3/T4 classification (points=32), with tumor located at 6 cm above from anal verge (points=39), with a tumor diameter of 10mm (points=50), without elevated tumor marker (points=0), with elevated preoperative WBC (points=35) would have a total score of 156, and a predicted LNM risk of 75%.

The area under ROC curve (AUC) of the model was 0.743 (95% CI: 0.691–0.795), which was greater than MRI-based tumor location (0.673, 95% CI: 0.620–0.727), clinical T classification (0.626, 95% CI: 0.567–0.685), MRI-based tumor diameter (0.630, 95% CI: 0.568–0.692), preoperative WBC (0.524, 95% CI: 0.460–0.589) and preoperative elevated tumor markers (0.588, 95% CI: 0.522–0.654) in the training cohorts, suggesting a better predictive value (Figure 4). The AUC of the model was shown to be 0.777 (95% CI: 0.705–0.848) in the validation cohort, which further confirmed the predictive value (Figure 4) The calibration graph showed favorable agreement between prediction and observation for both the training and validation cohorts because the calibration curve was close to the 45-degree line, which suggests that the model can perfectly predict the real event as shown in Figure 4.

Clinical applications

The decision curve analysis of the nomogram showed that the nomogram adds more benefit to LNM prediction in clinically node-negative rectal cancer patients than either the treat-all-patients scheme or the treat-none scheme when
the threshold probability of a patient or doctor is between 5% and 70% (Figure 5). The true and false positive rates in each risk threshold were also shown in Figure 5.

**Discussion**
Preoperative CRT is a standard regimen for advanced rectal cancer patients due to its prominent advantages on local control as well as overall survival (6,7,13). The widely accepted clinical indications for preoperative CRT are rectal cancer patients with lymph node involvement (9,14). Therefore, assessment of accurate preoperative staging of rectal cancer patients is crucial for clinical decision-making.

MRI is recommended as the best imaging approach for clinical staging of rectal cancer (15). However, MRI, which is not capable of detecting metastatic lymph nodes <3 mm, exhibits an accuracy rate of 80% for predicting lymph node involvement (15,22). In this study, the false negative rate of MRI for predicting lymph node involvement ranged from 20% to 30%, which is consistent with previous reports. The limitations of existing imaging tools may exclude prospective patients from CRT because of missing diagnoses of LNM. Therefore, finding other methods to assess accurate status of regional lymph node involvement in rectal cancer patients is of great importance. In the present study, we aimed to establish a clinical scoring system based on preoperative parameters to accurately assess the status of lymph nodes in clinically node-negative rectal cancer patients.

Nomograms have been widely used to visualize risk factors and prognosis in CRC patients (30,31). In our study, six potential predictors from 13 risk factor candidates were used to establish a nomogram by shrinking the regression coefficients using the LASSO method, which has been

<table>
<thead>
<tr>
<th>Characteristics</th>
<th>The training cohort</th>
<th>The validation cohort</th>
</tr>
</thead>
<tbody>
<tr>
<td>Number of patients</td>
<td>434</td>
<td>165</td>
</tr>
<tr>
<td>LNM</td>
<td>105 (24.2)</td>
<td>51 (30.9)</td>
</tr>
<tr>
<td>Age (years)</td>
<td>59.1±12.7</td>
<td>59.8±11.7</td>
</tr>
<tr>
<td>Gender</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Female</td>
<td>183 (42.2)</td>
<td>65 (39.4)</td>
</tr>
<tr>
<td>Male</td>
<td>251 (57.8)</td>
<td>100 (60.6)</td>
</tr>
<tr>
<td>Body mass index (kg/m²)</td>
<td>22.7±3.4</td>
<td>22.7±3.2</td>
</tr>
<tr>
<td>CEA level (ng/mL)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>≤5</td>
<td>318 (73.3)</td>
<td>119 (72.1)</td>
</tr>
<tr>
<td>&gt;5</td>
<td>116 (26.7)</td>
<td>46 (27.9)</td>
</tr>
<tr>
<td>CA199 level (U/mL)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>≤37</td>
<td>395 (91.0)</td>
<td>153 (92.7)</td>
</tr>
<tr>
<td>&gt;37</td>
<td>39 (9.0)</td>
<td>12 (7.3)</td>
</tr>
<tr>
<td>Preoperative total protein (g/L)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>&lt;60</td>
<td>26 (6.0)</td>
<td>10 (6.1)</td>
</tr>
<tr>
<td>≥60</td>
<td>408 (94)</td>
<td>155 (93.9)</td>
</tr>
<tr>
<td>Preoperative albumin (g/L)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>&lt;35</td>
<td>13 (3.0)</td>
<td>7 (4.2)</td>
</tr>
<tr>
<td>≥35</td>
<td>421 (97.0)</td>
<td>158 (95.8)</td>
</tr>
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</table>

Table 1 (continued)

<table>
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<tr>
<th>Characteristics</th>
<th>The training cohort</th>
<th>The validation cohort</th>
</tr>
</thead>
<tbody>
<tr>
<td>Preoperative hemoglobin (g/L)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>&lt;110</td>
<td></td>
<td></td>
</tr>
<tr>
<td>≥110</td>
<td>381 (87.8)</td>
<td>139 (84.2)</td>
</tr>
<tr>
<td>Preoperative PLT (10⁹/L)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>≤300</td>
<td>386 (88.9)</td>
<td>140 (84.8)</td>
</tr>
<tr>
<td>&gt;300</td>
<td>48 (11.1)</td>
<td>25 (15.2)</td>
</tr>
<tr>
<td>Preoperative WBC (10⁹/L)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>≤10</td>
<td>417 (96.1)</td>
<td>155 (93.9)</td>
</tr>
<tr>
<td>&gt;10</td>
<td>17 (3.9)</td>
<td>10 (6.1)</td>
</tr>
<tr>
<td>Location (cm)</td>
<td>5.0±2.8</td>
<td>5.8±2.8</td>
</tr>
<tr>
<td>Tumor diameter (cm)</td>
<td>5.4±3.5</td>
<td>3.2±1.5</td>
</tr>
<tr>
<td>Clinical T classification</td>
<td>T1</td>
<td>37 (8.5)</td>
</tr>
<tr>
<td></td>
<td>T2</td>
<td>145 (33.4)</td>
</tr>
<tr>
<td></td>
<td>T3</td>
<td>230 (53.0)</td>
</tr>
<tr>
<td></td>
<td>T4</td>
<td>22 (5.1)</td>
</tr>
</tbody>
</table>

CA199, carbohydrate antigen 19-9; CEA, carcinoembryonic antigen; LNM, lymph node metastasis; location, MRI-based distance between tumor and anal verge; PLT, platelet; tumor diameter, MRI-based diameter of tumor; WBC, white blood cell.
Figure 1 Suboptimal factor selection using the least absolute shrinkage and selection operator (LASSO) binary logistic regression model. (A) Tuning parameter (Lambda) selection in the LASSO model used 10-fold cross-validation via minimum criteria. The Binomial Deviance was plotted versus log (Lambda). Dotted vertical lines were drawn at the optimal values using the minimum criteria and the 1 standard error of the minimum criteria (the 1-SE criteria). (B) LASSO coefficient profiles of the 13 texture features. A coefficient profile plot was generated against the log (Lambda) sequence. Vertical line represents the values selected using 10-fold cross-validation, where optimal lambda resulted in 6 nonzero coefficients.

Figure 2 Forest plot of multivariable logistic model associations with LNM in clinically node-negative rectal cancer patients. Location: MRI-based distance between tumor and anal verge; tumor diameter: MRI-based diameter of tumor; tumor marker: 0= normal CEA and CA199, 1= elevated CEA or CA199, 2= elevated CEA and CA199; LNM, lymph node metastases; MRI, magnetic resonance imaging; WBC, white blood cell.

Figure 3 The nomogram for predicting LNM in clinically node-negative rectal cancer patients. The nomogram was established in the training cohort using multivariable regression, consisting of the location, clinical T classification, tumor diameter, WBC and tumor marker. LNM, lymph node metastases; WBC, white blood cell.
recommended for variable selection (21,32). Then, the multivariate logistic regression model was used for the determination of the optimal predictors (33,34).

T classification, which is an index of the depth of tumor invasion, has been well recognized as one of the most important predictors of LNM for CRC (24,35-38). A study based on 804 cases reported that the percentage of lymph node involvement arises from 5.7% to 78.8% accordingly when the T classification increases from 1 to 4 (36). In our study, we found that the clinical T classification is a significant independent predictor of LNM in clinically node-negative rectal cancer patients, which is consistent with previous studies.

The diameter of tumor or the tumor size was also a LNM predictor in our nomogram, which was in accordance with the study conducted by Zhang et al. (39). While several studies failed to identify a direct association between tumor size and LNM, they demonstrated an association between tumor size and tumor progression. For example, Cai et al. (40) reported that the tumor size was significantly related to local advancement of tumors. Chok and colleagues (41) also reported that the tumor size was associated with pT classification. Furthermore, a study conducted by Kikuchi et al. (42) found a direct correlation between tumor diameter and the depth of tumor invasion in 182 CRC cases. Therefore, increased diameter of tumor, which is an independent prognostic factor of tumor progression, is also relatively associated with LNM.

**Figure 4** (A,B) Calibration curves of the developed nomogram in each cohort. (A) Calibration curve of the nomogram in the training cohort. (B) Calibration curve of the nomogram in the validation cohort. The x-axis represents the predicted probability from the nomogram, and the y-axis is the actual probability of LNM in clinically node-negative rectal cancer patients. The dashed line indicates that predicted outcome perfectly corresponds with actual outcome. The blue line indicates the bias-corrected accuracy of the nomogram while a closer fit to the diagonal dotted line represents a better prediction. (C,D) Receiver operating characteristic (ROC) curve. (C) ROC curve in the training cohort [area under the curve (AUC): 0.743]. (D) ROC curve in the validation curve (AUC: 0.777). Diagonal segments are produced by ties. LNM, lymph node metastases.
CEA and CA19-9, which are the most common tumor markers of CRC, have been currently utilized in clinical practice for cancer screening and follow-up (43,44). Previous studies have fully demonstrated that increased levels of CEA and CA19-9 were risk factors of tumor recurrence and anti-tumor agent resistance (45). A study based on 130 CRC patients revealed that elevated CEA and CA19-9 were significantly correlated with lymph nodes involvement (45). In the present study, our results also suggest that the elevation of these two preoperative tumor markers were predictors for LNM.

The location of tumor also served as a prognostic factor of LNM in our study. Longer distance between tumor and the anal verge is a risk factor of LNM. To the best of our knowledge, only a few studies focus on the association between tumor location and LNM. Two studies (46,47) even demonstrated the opposite results. Therefore, it is difficult to draw a conclusion on the relationship between tumor location and LNM risk based on currently available studies. Further studies are highlighted to carefully assess the exact relationship between the location of tumor and LNM.

The white blood cell count (WBC) was a widely used marker of infection and inflammation (48). Numerous studies have demonstrated that inflammation is involved in tumorigenesis and metastasis (49), mainly through changing the systemic immune status and local microenvironment (50), which is partially reflected by the abnormalities of WBC count (51). Several studies have found that more WBCs were associated with carcinogenesis, tumor progression and cancer mortality (48,51-53). Our study suggests that elevated WBC level was an ominous prognostic sign of LNM.

While several studies evaluated predictors for LNM, no study focused on the prediction of LNM in clinically node-negative rectal cancer patients. In this study, we established a nomogram consisting of 5 clinical factors. The decision curve showed that our nomogram is better than either the treat-all-patients scheme or the treat-none scheme in LNM prediction when the threshold probability of a patient or doctor ranges between 10% and 67%. (B) The true positive rates and false positive rates in each risk thresholds. The black light represents the true positive rates of the model, while the red light represents the false one. LNM, lymph node metastases.
Thus, our nomogram, which has up to 75% of accuracy of detecting LNM, provides supplemental information for individual clinical decision, which can be used to properly enroll CRC patients for CRT.

There are several limitations in our study. For instance, our study did not fully take advantages of the MRI examination. Recently, radiomics, which incorporate the features of imaging examination, has emerged in the field of cancer research (21). The nomogram including imaging features may be more convincing than using the result of MRI examination alone. Besides, 30% of rectal cancer patients were initially considered to be node-positive but were negative in the pathology report, and these patients might be overtreated (21). Therefore, it is of great importance to develop models for preoperative individualized assessment of LN metastasis in clinically node-positive patients, which requires our further investigation.

In conclusion, we established a nomogram consisting of five preoperative clinical factors for individualized prediction of LNM in clinically node-negative rectal cancer patients. Our nomogram is useful for enrolling CRC patients for CRT.

Acknowledgments

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Footnote

Conflicts of Interest: The authors have no conflicts of interest to declare.

Ethical Statement: The authors are accountable for all aspects of the work in ensuring that questions related to the accuracy or integrity of any part of the work are appropriately investigated and resolved. This study was approved by the Institutional Review Board (IRB) of The Sixth Affiliated Hospital of Sun Yat-sen University. All patients provided written informed consent to participate in this study.

References


41. Chok KS, Law WL. Prognostic factors affecting survival

# Related Computerized Programs for the Nomogram Using R in Current Study

## For the least absolute shrinkage and selection operator (LASSO) binary logistic regression mode

```r
library(glmnet)
fit<-glmnet(model.matrix(~Age+Gender..........,data=training_data),as.matrix(PositiveN,data=training_data),family="binomial",alpha=1)
cv.fit<-cv.glmnet(model.matrix(~Age+Gender..........,data=training_data),as.matrix(PositiveN,data=training_data),family="binomial",alpha=1)
predict(cv.fit,type='coefficients',s=cv.fit$lambda.min)
plot(cv.fit)
plot(cv.fit$glmnet.fit,xvar ="lambda")
```

## For Nomogram

```r
library(rms)
f<-lrm(PositiveLNM ~ Location+ClinicalTstage......,x=T,y=T)
nom <- nomogram(f, fun=plogis,fun.at=c(.05, seq(.1,.9, by=.1), .95, 1),lp=F,maxscale=100, funlabel="Risk of PositiveLNM")
plot(nom)
```

## For Resampling Validation of Nomogram

```r
validate(f,method="boot",B=1000,dxy=T)
```

## For Computing the AUC (C-Index) and 95% CI

```r
library(ROCR)
pred<-prediction(pre,PositiveLNM)
performance(pred,'auc')
SE<-.SD/2; 95%CI=AUC(C-Index)±1.96×SE
```

## For Calibration Curve

```r
plot(calibrate(f,method="boot",B=1000),scat1d.opts=list(nhistSpike=240,side=1,frac=0.08,lwd=1,nint=50))
lines(calibrate(f,method="boot",B=1000), lwd=2, lty=5, col=c(rbg(0,0,255,maxColorValue= 255)))
abline(0,1,lty =3,lwd=2,col=c(rbg(255,0,0,maxColorValue= 255)))
```

## For External Validation of Nomogram

```r
f1<-lrm(PositiveLNM~predict(f, data=validation_data),x=T,y=T)
validate(f1,method="boot",B=1000,dxy=T)
```

## For the Calibration Curve of the Validation Cohort

```r
plot(calibrate(f1,method="boot",B=1000),scat1d.opts=list(nhistSpike=240,side=1,frac=0.08,lwd=1,nint=50))
lines(calibrate(f1,method="boot",B=1000), lwd=2, lty=5, col=c(rbg(0,0,255,maxColorValue= 255)))
abline(0,1,lty =3,lwd=2,col=c(rbg(255,0,0,maxColorValue= 255)))
```

## For Computing the AUC (C-Index) and 95% CI of the Validation Cohort

```r
f1 <- glm(PositiveLNM ~ predict(f, data=validation_data),family=binomial(link='logit'),data=validation_data)
pred<-predict(f1,type='response')
pred<-prediction(pre,PositiveLNM)
performance(pred,'auc')
SE<-.SD/2; 95%CI=AUC(C-Index)±1.96×SE
```

## For the Decision Curve of the Validation Cohort

```r
library(MASS)
validation_data$PositiveLNM = predict(f1, type="response")
plot(dca(data=validation_data, outcome="PositiveN", predictors="PositiveLNM", smooth=TRUE,xstop=1)$net.benefit.threshold, km$net.benefit.none, type = "l", lwd=2,xlim=c(0,1), ylim=c(0,0.32), xlab = "Threshold Probability",ylab = "Net Benefit")
```