Lymphovascular invasion quantification could improve risk prediction of lymph node metastases in patients with submucosal (T1b) esophageal adenocarcinoma


1Department of Gastroenterology and Hepatology, Erasmus MC Cancer Institute, University Medical Center, Rotterdam, The Netherlands
2Department of Pathology, Erasmus MC Cancer Institute, University Medical Center, Rotterdam, The Netherlands
3Department of Pathology, Isala Clinics, Zwolle, The Netherlands
4Department of Public Health, Erasmus MC Cancer Institute, University Medical Center, Rotterdam, The Netherlands
5Department of Gastroenterology and Hepatology, University Medical Center Utrecht, Utrecht, The Netherlands
6Department of Gastroenterology and Hepatology, St. Antonius Hospital, Nieuwegein, The Netherlands
7Department of Pathology, University Medical Center Utrecht, Utrecht, The Netherlands
8Department of Surgery, University Medical Center Utrecht, Utrecht, The Netherlands
9Department of Pathology, Pathology DNA, St. Antonius Hospital, Nieuwegein, The Netherlands
10Department of Gastroenterology and Hepatology, Isala Clinics, Zwolle, The Netherlands
11Department of Gastroenterology and Hepatology, Catharina Hospital, Eindhoven, The Netherlands
12Department of Pathology, PAMM, Eindhoven, The Netherlands
13Department of Gastroenterology and Hepatology, IJsselland Hospital, Capelle aan den IJssel, The Netherlands
14Department of Pathology, IJsselland Hospital, Capelle aan den IJssel, The Netherlands
15Department of Gastroenterology and Hepatology, University Medical Center Groningen, Groningen, The Netherlands
16Department of Pathology, University Medical Center Groningen, Groningen, The Netherlands
17Department of Surgery, University Medical Center Groningen, Groningen, The Netherlands
18Department of Gastroenterology and Hepatology, Haga Teaching Hospital, Den Haag, The Netherlands
19Department of Pathology, Haga Teaching Hospital, Den Haag, The Netherlands
20Department of Gastroenterology and Hepatology, Amsterdam University Medical Centers, Amsterdam, The Netherlands
21Department of Pathology, Amsterdam University Medical Centers, Amsterdam, The Netherlands
Abstract

Aim: To quantify lymphovascular invasion (LVI) and to assess the prognostic value in patients with pT1b esophageal adenocarcinoma.

Methods: In this nationwide, retrospective cohort study, patients were included if they were treated with surgery or endoscopic resection for pT1b esophageal adenocarcinoma. Primary endpoint was the presence of metastases, lymph node metastases, or distant metastases, in surgical resection specimens or during follow-up. A prediction model to identify risk factors for metastases was developed and internally validated.

Results: 248 patients were included. LVI was distributed as follows: no LVI (n = 196; 79.0%), 1 LVI focus (n = 16; 6.5%), 2–3 LVI foci (n = 21; 8.5%) and ≥4 LVI foci (n = 15; 6.0%). Seventy-eight patients had metastases. The risk of metastases was increased for tumors with 2–3 LVI foci [subdistribution hazard ratio (SHR) 3.39, 95% confidence interval (CI) 2.10–5.47] and ≥4 LVI foci (SHR 3.81, 95% CI 2.37–6.10). The prediction model demonstrated a good discriminative ability (c-statistic 0.81).

Conclusion: The risk of metastases is higher when more LVI foci are present. Quantification of LVI could be useful for a more precise risk estimation of metastases. This model needs to be externally validated before implementation into clinical practice.

KEYWORDS
endoscopic mucosal resection, esophagectomy Lymphovascular invasion, LVI, lymph node metastases, prediction, quantification, risk assessment, submucosal esophageal adenocarcinoma, T1b adenocarcinoma

INTRODUCTION

The presence of lymph node metastases (LNM) is an important prognostic factor in submucosal esophageal adenocarcinoma (pT1b EAC). Various risk factors for LNM have been identified including depth of infiltration, grade of differentiation, and lymphovascular invasion (LVI).

We recently developed a prediction model based on histopathological risk factors to estimate the risk for metastases in patients with pT1b EAC. We showed that LVI was the strongest predictor for metastases with an estimated 5-year risk of developing metastases ranging between 15.7% and 70.1% for pT1b EAC with LVI as compared to 5.9%–35.2% for those without LVI. Other studies reported a worse 5-year survival in patients with pT1b EAC with LVI compared to pT1b EAC without LVI (27% vs. 77%, p < 0.01).

LVI is often reported as a dichotomous parameter, either absent or present. Intuitively, there may be a differential metastatic risk in case of only a single LVI focus versus a case with extensive LVI. This is supported by a study on gastric cancer in which a higher number of lymphatic tumor emboli was an independent predictor for LNM. We hypothesized that the number of LVI foci in pT1b EAC is of influence on patient outcome. Quantification of LVI may possibly further classify these patients into low and high risk for the development of metastases.
The aim of this study was to quantify LVI and to assess the prognostic value in patients with pT1b EAC.

MATERIALS AND METHODS

This was a nationwide retrospective cohort study. Patients diagnosed with pT1b EAC between 1989 and 2016 were included. This study was approved by all Medical Ethical Review Committees in the participating centers (MEC-2016-050). A detailed description of inclusion and exclusion criteria, data collection, patient selection, reassessment of histopathological variables, and the statistical analyses can be found in our previous study, as the present study is an extension to our previous study. Histopathological reassessment was performed by three gastrointestinal pathologists for 84 patients to calculate the inter-observer agreement as explained in our previous study. For the other 164 patients, histopathological reassessment was performed by one pathologist for LVI ($\kappa = 0.88$) and differentiation grade ($\kappa = 0.77$) because of the good and excellent interobserver agreement.

A model was developed to predict the risk of metastases in patients with pT1b EAC. In the prediction model of the present study, quantification of LVI on a 4-tier assessment scale is incorporated instead of interpreting LVI as a dichotomous parameter. The same patient cohort is used, and the same statistical analyses are performed as in our previous study. The added value of LVI quantification to the model was assessed using the likelihood ratio test and quantified by the increase in Harrell’s concordance index ($c$-statistic).

Histopathological reassessment of LVI and LVI quantification

Whole case hematoxylin and eosin stained (H&E) slide sets were used for LVI reassessment. Immunohistochemical staining was not performed because formalin-fixed paraffin embedded tissue blocks were not available. Initially, LVI was defined as being present or absent as in standard clinical practice. All slides were assessed with high power up to and including the submucosa. When LVI was present, all LVI foci were counted in every H&E slide that was available. In general, resection specimens of 2-mm thickness were assessed when endoscopic resection was performed in contrast to 5-mm thickness when surgical resection was performed. Figure 1 shows an example of an H&E slide with three LVI foci. The LVI focus number was the sum of all counted LVI foci in all H&E slides from one patient. In patients in whom LVI was analyzed by three pathologists, the highest LVI focus number was included for analysis. Patients were categorized into four groups based on the number of LVI foci: no LVI, 1 LVI focus, 2–3 LVI foci, or ≥4 LVI foci. This 4-tier assessment scale was used to create different LVI foci groups with less LVI (1 LVI focus), moderate LVI (2–3 LVI foci), or extensive LVI (≥4 LVI foci).

![Figure 1](image.jpg)

**Figure 1** Quantification of LVI: three LVI foci, indicated by an arrow. This case nicely illustrated an example of three foci of LVI, which may very well all be located in the same lymph vessel.

**Table 1** Patient characteristics and number of LVI foci ($n = 248$)

<table>
<thead>
<tr>
<th>Parameter</th>
<th>Total cohort (n = 248)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Gender, n (%)</td>
<td></td>
</tr>
<tr>
<td>Male</td>
<td>217 (87.5%)</td>
</tr>
<tr>
<td>Female</td>
<td>31 (12.5%)</td>
</tr>
<tr>
<td>Median age, years (IQR)</td>
<td>65.6 (57.8–72.5)</td>
</tr>
<tr>
<td>Presence of LVI, n (%)</td>
<td>52 (21.0%)</td>
</tr>
</tbody>
</table>

<table>
<thead>
<tr>
<th>LVI foci</th>
<th>Total number</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>16</td>
</tr>
<tr>
<td>2</td>
<td>10</td>
</tr>
<tr>
<td>3</td>
<td>11</td>
</tr>
<tr>
<td>4</td>
<td>6</td>
</tr>
<tr>
<td>5–10</td>
<td>9</td>
</tr>
</tbody>
</table>

<table>
<thead>
<tr>
<th>Metastases, n (%)</th>
<th></th>
</tr>
</thead>
<tbody>
<tr>
<td>LNM</td>
<td>49 (19.8%)</td>
</tr>
<tr>
<td>Distant metastasis</td>
<td>6 (2.4%)</td>
</tr>
<tr>
<td>LNM + distant metastasis</td>
<td>23 (9.3%)</td>
</tr>
</tbody>
</table>

Abbreviations: IQR, interquartile range; LNM, lymph node metastases; LVI, lymphovascular invasion.

Endpoints

The primary endpoint was the presence metastases defined as LNM in surgical resection specimens or the development of metastases (LNM or distant metastases) during follow-up. In case surgical resection was performed, at least 12 lymph nodes were required for adequate assessment of LNM. The development of metastases during follow-up was used as a surrogate endpoint in case no surgery was performed after endoscopic resection (ER), or if surgical resection specimen contained <12 lymph nodes without LNM. In these cases, a minimum follow-up period of 2 years was required. If patients died within these 2 years, they were only included if cause of death was known.
RESULTS

In total, 248 patients with pT1b EAC were included. Patient characteristics are presented in Table 1. Primary endoscopic resection was performed in 82 of 248 (33.1%) patients and primary surgery was performed in 166 of 248 (66.9%) patients. Of patients who were treated with endoscopic resection, additional surgery was performed in 49/82 (59.8%) patients. Local recurrence was detected during follow-up in 12 of 248 (4.8%) patients, of whom 11 also developed metastasis.

Quantification of LVI

There were 52 patients with LVI positive tumors. The LVI focus number was distributed from 1 to 10 foci (Table 1). LVI quantification was determined by three gastrointestinal pathologist in 17/52 patients (F.T.K., M.D., K.B.). In the remaining 35 patients, LVI quantification was determined by one pathologist (F.T.K.) because the inter-observer variability was excellent for LVI ($\kappa = 0.88$). The median LVI focus number was 2.5 [interquartile range (IQR) 1–4]. LVI quantification was determined by three gastrointestinal pathologist in 17/52 patients (F.T.K., M.D., K.B.). In the remaining 35 patients, LVI quantification was determined by one pathologist (F.T.K.) because the inter-observer variability was excellent for LVI ($\kappa = 0.88$). The median LVI focus number was 2.5 [interquartile range (IQR) 1–4]. LVI was categorized as follows: no LVI ($n = 196; 79.0\%$), 1 LVI focus ($n = 16; 6.5\%$), 2–3 LVI foci ($n = 21; 8.5\%$), and $\geq 4$ LVI foci ($n = 15, 6.0\%$).

Metastases

Some 78 of 248 patients had metastases (Table 1), with a 5-year cumulative incidence of 30.9% (95% CI 25.1–36.8%) (Figure 2). In patients treated with primary surgery ($\geq 12$ lymph node dissections), the median follow-up time was 3.3 years (IQR 1.8–5.3). In patients treated with surgery ($< 12$ lymph node dissections) or endoscopic resection only, the median follow-up period was 5.5 years (IQR 4.9–7.7) The majority of patients developed LNM only (49/78). These
Patients were all treated with surgery; LNM were detected during surgery in 47 patients and during follow-up in 2 patients. Six patients developed distant metastases only, and all were treated with surgery for EAC (without LNM). Adequate lymph nodes (>12) were resected in only 2 of these 6 patients. Twenty-three patients developed both LNM and distant metastases. Surgery was performed in 22/23 patients, LNM was found in surgical resection specimen in 19 patients and during follow-up in 3 patients. One patient was treated with endoscopic resection only due to comorbidity, metastases were detected during follow-up. Of all patients who developed metastases during follow-up, the median interval between EAC diagnosis and detection of metastases was 1.9 years (IQR: 1.2–4.9).

**Risk factor analysis**

**Univariable analysis**

The number of LVI foci was associated with the presence of metastases. Tumors with 1 LVI focus had metastases in 37.5%, tumors with 2–3 LVI foci had metastases in 71.4%, and tumors with ≥4 LVI foci had metastases in 93.3% (Table 2). In contrast, 22% of patients without LVI developed metastases.

**Multivariable analysis**

For every increase of tumor invasion with 500 μm, the subdistribution hazard of developing metastases increased with 1.08 (95% CI 1.02–1.14) (Table 2). The presence of 2–3 LVI foci (SHR 3.39, 95% CI 2.10–5.47) and the presence of ≥4 LVI foci (SHR 3.81, 95% CI 2.37–6.10) were independent predictors for metastases compared to patients without LVI. In contrast, the presence of only one LVI focus was not significantly correlated with metastases after multivariable analysis. For every increase of tumor size with 10 mm, the subdistribution hazard of developing metastases increased with 1.23 (95% CI 1.09–1.38). Poor differentiation grade was not an

### Table 2

Univariable and multivariable subdistributional hazard regression analyses of risk factors associated with metastases (no metastases; n = 170, metastasis; n = 78)

<table>
<thead>
<tr>
<th>Variable</th>
<th>No metastases (n = 170)</th>
<th>Metastases (n = 78)</th>
<th>Univariable subdistributional hazard regression</th>
<th>Multivariable subdistributional hazard regression</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td></td>
<td></td>
<td>SHR (95% CI) p-value</td>
<td>SHR (95% CI) p-value</td>
</tr>
<tr>
<td>Sex, n (%)</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Female</td>
<td>24 (77%)</td>
<td>7 (23%)</td>
<td>Reference</td>
<td>-</td>
</tr>
<tr>
<td>Male</td>
<td>146 (67%)</td>
<td>71 (33%)</td>
<td>1.51 (0.74–3.10) 0.26</td>
<td>-</td>
</tr>
<tr>
<td>Differentiation grade, n (%)</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>G1/G2 (good/moderate)</td>
<td>121 (75%)</td>
<td>41 (25%)</td>
<td>Reference</td>
<td>Reference</td>
</tr>
<tr>
<td>G3/4 (poor/undifferentiated)</td>
<td>49 (57%)</td>
<td>37 (43%)</td>
<td>1.78 (1.20–2.65) &lt;0.01</td>
<td>0.98 (0.65–1.50) 0.94</td>
</tr>
<tr>
<td>Median tumor length (mm) (IQR)</td>
<td>20 (13–29)</td>
<td>30 (25–44)</td>
<td>1.39 (1.25–1.53) *&lt;0.01</td>
<td>1.23 (1.09–1.38) *&lt;0.01</td>
</tr>
<tr>
<td>Median submucosal invasion (μm) (IQR)</td>
<td>958 (409–2100)</td>
<td>2165 (1173–3500)</td>
<td>1.13 (1.08–1.18)** &lt;0.01</td>
<td>1.08 (1.02–1.14)** &lt;0.01</td>
</tr>
<tr>
<td>LVI, n (%)</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>No LVI</td>
<td>153 (78%)</td>
<td>43 (22%)</td>
<td>Reference</td>
<td>-</td>
</tr>
<tr>
<td>LVI 1</td>
<td>10 (62.5%)</td>
<td>6 (37.5%)</td>
<td>1.81 (0.84–3.90) 0.13</td>
<td>1.72 (0.89–3.32) 0.11</td>
</tr>
<tr>
<td>LVI 2–3</td>
<td>6 (28.6%)</td>
<td>15 (71.4%)</td>
<td>3.90 (2.47–6.15) *&lt;0.01</td>
<td>3.39 (2.10–5.47) *&lt;0.01</td>
</tr>
<tr>
<td>LVI ≥4</td>
<td>1 (6.7%)</td>
<td>14 (93.3%)</td>
<td>5.54 (3.82–8.04) *&lt;0.01</td>
<td>3.81 (2.37–6.10) *&lt;0.01</td>
</tr>
</tbody>
</table>

Abbreviations: CI, confidence interval; IQR, interquartile range; LVI, lymphovascular invasion; SHR, subdistribution hazard ratio.

*For every increase of 10 mm.

**For every 500 μm increase of sm invasion.

### Table 3

Score chart; 5-year risk (%) of developing metastases for different combinations of histopathological characteristics in pT1b esophageal adenocarcinoma; LVI incorporated as present or absent

<table>
<thead>
<tr>
<th>Tumor size</th>
<th>Submucosal invasion</th>
<th>LVI–% (95% CI)</th>
<th>LVI+% (95% CI)</th>
</tr>
</thead>
<tbody>
<tr>
<td>&lt;20 mm</td>
<td>sm1</td>
<td>5.9 (2.3–11.2)</td>
<td>15.7 (6.0–29.3)</td>
</tr>
<tr>
<td></td>
<td>sm2</td>
<td>7.3 (2.6–13.8)</td>
<td>19.3 (6.3–36.8)</td>
</tr>
<tr>
<td></td>
<td>sm3</td>
<td>14.1 (7.9–21.9)</td>
<td>34.7 (19.7–50.8)</td>
</tr>
<tr>
<td>≥20 mm</td>
<td>sm1</td>
<td>16.1 (6.2–29.2)</td>
<td>38.8 (17.0–61.4)</td>
</tr>
<tr>
<td></td>
<td>sm2</td>
<td>19.4 (8.6–32.2)</td>
<td>45.6 (20.8–67.9)</td>
</tr>
<tr>
<td></td>
<td>sm3</td>
<td>35.2 (25.8–44.7)</td>
<td>70.1 (60.5–78.7)</td>
</tr>
</tbody>
</table>

Abbreviations: CI, confidence interval; LVI, lymphovascular invasion; sm, submucosal.
independent risk factor for metastases (SHR 0.98, 95% CI 0.65–1.50),
and this variable was therefore not incorporated into the prediction
model.

Prediction model development and validation

The 5-year risk of developing metastases for different combinations
of histopathological tumor characteristics is illustrated in Table 4.
The prediction model with LVI incorporated as a dichotomous vari-
able is presented in Table 3.6 In Table 4, the predicted score was
16.1% (95% CI 6.2–29.2) for patients with pT1b EAC ≥20 mm, sm1
invasion depth without LVI, compared to 22.2% (95% CI 6.2–45.3)
when one LVI focus is present, 37.0% (95% CI 16.0–62.3) when 2–3
LVI foci are present, and 47.1% (95% CI 21.1–72.9) when ≥4 LVI foci
are present. Internal validation of the prediction model showed a
good discriminative ability with a c-statistic of 0.81 (95% CI 0.74–
0.87), which did not increase compared to the c-statistic of the pre-
vious model (p = 0.71).

DISCUSSION

LVI has so far only been evaluated as a dichotomous parameter:
either absent or present.6–9 In this study, we aimed to determine
whether quantification of LVI provides additional prognostic infor-
mation in patients with pT1b EAC. The results of our study show that
the presence of more LVI foci was correlated with a higher risk for
metastases. We believe that quantification of LVI may further classify
pT1b EAC for a more accurate risk estimation of metastases.

Accurate risk estimation is important for shared decision-making,
whether or not to undergo adjuvant therapy after ER of pT1b EAC.
Compared to the risk in LVI+ patients in the previous model, the
metastases risk in the current model is lower in case of 1 LVI focus
and higher in case of 2–3 LVI foci and ≥4 LVI foci.6 As a consequence,
a lower metastases risk can result in the decision to perform endo-
scopic surveillance instead of esophagectomy and a high metastases
risk may result in an advice to offer neo-adjuvant therapy before
esophagectomy. The more accurate this prediction is, the better a
patient can be informed and the better the decision about (neo-) adjuvant therapy can be made.

Remarkable is that the presence of only 1 LVI focus was not an
independent predictor for metastases. The predicted 5-year metas-
tases risk in patients with only 1 LVI focus is more in line with the risk
in patients without LVI. One could argue that the presence of 1 LVI
focus is negligible.

To our knowledge, this is the first study that evaluates the
quantification of LVI in pT1b EAC. We could therefore not compare
our results with previous studies. LVI has also been identified as a
strong predictor for LNM in other cancer types.15,18 The number of
lymphatic tumor emboli was an independent predictor for LNM in
early gastric cancer with LVI.15 The more lymphatic tumor emboli
were present, the higher the LNM risk.

Despite the promising results of our study, it is unlikely that the
application of the current model will change clinical practice soon.
The difference in predicted metastasis risk between the LVI foci
groups is relatively small and still high in all scenarios. Although the
metastases risk in patients with only 1 LVI focus is much lower than
the risk in patients with 2–3 or ≥4 LVI foci, the advice for adjuvant
treatment or active surveillance after ER will probably not change
based on the number of LVI foci. We are looking for a better risk
assessment for metastasis to make active surveillance possible on the
one hand. On the other hand, the presence of extensive LVI could be
a reason to combine surgery with neo-adjuvant therapy. The better
this risk assessment, the better we can apply tailored therapy for
patients with pT1b EAC. Therefore, further research on LVI quanti-
fication is necessary to assess whether it is possible to identify a
subgroup of patients with LVI-positive tumors with a low or high risk
of metastases.

Our study is subject to certain limitations. First, the retro-
pective design of our study could result in information and selec-
tion bias. There was no standard follow-up regime, which could
have resulted in a different number of reported and actual numbers
of metastases. Second, although the inter-observer agreement for
LVI was excellent as presented in our previous study, differences in
LVI foci between pathologists were not discussed in a consensus

---

TABLE 4 Score chart; 5-year risk (%) of developing metastases for different combinations of histopathological characteristics in pT1b esophageal adenocarcinoma; LVI incorporated as absent, 1 LVI focus, 2–3 LVI foci, or ≥4 LVI foci

<table>
<thead>
<tr>
<th>Tumor size</th>
<th>Submucosal invasion</th>
<th>LVI–</th>
<th>LVI 1x</th>
<th>LVI 2–3</th>
<th>LVI ≥4</th>
</tr>
</thead>
<tbody>
<tr>
<td>&lt;20 mm</td>
<td>sm1</td>
<td>5.9</td>
<td>10.9</td>
<td>19.5</td>
<td>25.7</td>
</tr>
<tr>
<td></td>
<td>sm2</td>
<td>7.3</td>
<td>13.4</td>
<td>24.1</td>
<td>31.6</td>
</tr>
<tr>
<td></td>
<td>sm3</td>
<td>14.1</td>
<td>26.3</td>
<td>43.5</td>
<td>54.4</td>
</tr>
<tr>
<td>≥20 mm</td>
<td>sm1</td>
<td>16.1</td>
<td>22.2</td>
<td>37.0</td>
<td>47.1</td>
</tr>
<tr>
<td></td>
<td>sm2</td>
<td>19.4</td>
<td>26.3</td>
<td>44.4</td>
<td>55.2</td>
</tr>
<tr>
<td></td>
<td>sm3</td>
<td>35.2</td>
<td>48.4</td>
<td>70.2</td>
<td>80.6</td>
</tr>
</tbody>
</table>

Abbreviations: CI, confidence interval; LVI, lymphovascular invasion; sm, submucosal.
Moreover, when multiple LVI foci were detected, it was uncertain whether these foci represented LVI in different vascular structures. It is very well possible that multiple foci in the H&E section actually represented tumor localization in the same vascular structure. Third, additional immunohistochemical staining was not performed in the assessment of LVI and this may improve LVI detection. Moreover, retraction artifacts related to tissue specimen preparation are difficult to distinguish from LVI in H&E slides. Fourth, endoscopic and surgical resection specimens were combined in the results of our study, which introduces heterogeneity. Sampling and processing of endoscopic and surgical resection specimens were different (2 vs. 5 mm) and this might introduce bias in histological interpretation. In addition, the number of LVI foci was counted in every available H&E slide. It might be that the number of LVI foci is higher when more slides are available in one patient. However, studies about this subject are lacking. The last limitation is that external validation of our prediction model was not performed. Although our sample size was large, it was still not large enough to perform external validation. As a consequence, we do not know whether our model has adequate predictive value to be used in clinical practice. External validation is desirable when implementing it into daily clinical practice.

A major strength of our study is that all histopathological slides and tumor characteristics were reassessed by gastrointestinal pathologists. The incorporated tumor characteristics were valid without missing data. Another strength of the study is the cooperation with the Netherlands Cancer Registration, which enabled us to incorporate all eligible patients.

To conclude, this prediction model suggests that quantification of LVI may further refine risk estimation in pT1b EAC. External validation of the model and more research regarding LVI quantification are required to confirm our findings.

CONFLICT OF INTEREST
The authors have no conflicts of interest to declare.

DATA AVAILABILITY STATEMENT
The data that support the findings of this study are available on request from the corresponding author. The data are not publicly available due to privacy or ethical restrictions.

ORCID
Steffi E. M. van de Ven https://orcid.org/0000-0001-9793-0957
Annieke W. Gotink https://orcid.org/0000-0003-0509-3472
Roos E. Pouw https://orcid.org/0000-0002-4707-1186

REFERENCES


How to cite this article: van de Ven, SEM, Suzuki, L, Gotink, AW, ten Kate FJC, Nieboer D, Weusten BLAM, et al. Lymphovascular invasion quantification could improve risk prediction of lymph node metastases in patients with submucosal (T1b) esophageal adenocarcinoma. United European Gastroenterol J. 2021;1–8. https://doi.org/10.1002/ueg2.12151