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Tumor Budding Is Associated with an Increased Risk of Lymph Node Metastasis and Poor Prognosis in Superficial Esophageal Adenocarcinoma

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Abstract

The treatment approach for superficial (stage T1) esophageal adenocarcinoma critically depends on the pre-operative assessment of metastatic risk. Part of that assessment involves evaluation of the primary tumor for pathologic characteristics known to predict nodal metastasis: depth of invasion (intramucosal versus submucosal), angiolymphatic invasion, tumor grade and tumor size. Tumor budding is a histologic pattern that is associated with poor prognosis in early stage colorectal adenocarcinoma and a predictor of nodal metastasis in T1 colorectal adenocarcinoma. In a retrospective study, we used a semi-quantitative histologic scoring system to categorize 210 surgically resected, superficial (stage T1) esophageal adenocarcinoma according to the extent of tumor budding (none, focal and extensive) and also evaluated other known risk factors for nodal metastasis, including depth of invasion, angiolymphatic invasion, tumor grade and tumor size. We assessed the risk of nodal metastasis associated with tumor budding in univariate analyses and controlled for other risk factors in a multivariate logistic regression model. Forty-one percent (24/59) of tumors with extensive tumor budding (tumor budding in ≥ 3 20X microscopic fields) were metastatic to regional lymph nodes, compared to 10% (12/117) of tumors with no tumor budding and 15% (5/34) of tumors with focal tumor budding ($p < 0.001$). When controlling for all pathologic risk factors in a multivariate analysis, extensive tumor budding remains an independent risk factor for lymph node metastasis in superficial esophageal adenocarcinoma associated with a 2.5-fold increase (95% CI, 1.1–6.3, $p = 0.039$) in the risk of nodal metastasis. Extensive tumor budding is also a poor prognostic factor with respect to overall survival and time to recurrence in univariate and multivariate analyses. As an independent risk factor for nodal metastasis and survival after esophagectomy, tumor budding should be evaluated in superficial (T1) esophageal adenocarcinoma as a part of a comprehensive pathologic risk assessment.

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DISCLOSURE

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INTRODUCTION

In the majority of patients, surgically resected, superficial (T1) adenocarcinoma of the esophagus or gastroesophageal junction has a favorable survival outcome relative to more deeply invasive cancers.¹ However, despite tumor that is confined to the mucosal or submucosal layers, up to 16% of patients with T1 esophageal adenocarcinoma will have nodal metastases identified at surgical resection.²⁻⁸ These patients have significantly worse prognosis.^{5, 9}

Based on a widespread consensus in the literature,^{2, 3, 5-8, 10-16} submucosal invasion is routinely evaluated by staging endoscopic resection of superficial esophageal adenocarcinoma and is regarded as the paramount risk factor for nodal metastasis.¹⁷ However, there are other established risk factors for nodal metastasis, including angiolymphatic invasion,^{2, 3, 11, 14, 15, 17-19} higher grade,^{2, 3, 8, 11, 15, 17} and larger tumor size^{3, 11, 14, 17} which are also associated with nodal metastasis.

In addition to these, tumor budding is another histologic feature that has been shown to be associated with lymph node metastasis or poor prognosis in other gastrointestinal neoplasms, including gastric,²⁰ colorectal,²¹ and ampullary adenocarcinomas²² and esophageal carcinomas^{23, 24}. A tumor bud is defined as a detached cluster of fewer than 5 cells at the invasive front of a tumor.²⁵ “Tumor budding” is present when the number and density of buds exceeds a threshold, with various scoring methods and thresholds proposed. At least some types of tumor budding are thought to be the morphologic manifestation of an epithelial-to-mesenchymal transition during which tumor cells lose their intercellular attachments and acquire an invasive, mesenchymal phenotype that facilitates metastasis.^{21, 26, 27}

Although tumor budding has been previously studied in esophageal carcinomas, little is known about its prognostic utility in superficial esophageal adenocarcinoma. An indication of its potential utility was suggested in a recent abstract in which tumor budding was found to be a risk factor for nodal metastasis and tumor recurrence in a cohort of 42 surgically resected superficial (stage T1) esophageal adenocarcinomas.²⁸ Because there are multiple known pathologic predictors of nodal metastasis in superficial esophageal adenocarcinoma, it is important to evaluate tumor budding relative to these other prognostic factors to see whether it adds additional, independent prognostic information. Previous studies have not been sufficiently powered to do so. Therefore, the aims of this study are to document the prevalence and extent of tumor budding in surgically resected superficial esophageal adenocarcinoma and determine whether tumor budding is predictive of lymph node metastasis and survival when controlling for the effects of other important prognostic variables.

METHODS

Case Selection

We identified 210 patients with stage pT1 esophageal or gastroesophageal junction adenocarcinoma who underwent esophagectomy without induction therapy at University of

Pittsburgh Medical Center from 1996 to 2013 and had representative tumor slides available for review. Patients diagnosed with high grade dysplasia only or staged as T2 or higher were not included, nor were patients with curative endoscopic resection of the tumor.

Evaluation of Pathologic and Clinical Features

The tumor slides from all 210 cases were reviewed and assessed for tumor size, tumor grade, submucosal invasion, angiolymphatic invasion, and tumor budding (MSL and JMD). An average of 3.9 blocks per tumor were evaluated (range 1–25), representing an average of 2.6 blocks per cm of tumor, not including deeper levels that were examined in some cases.

Tumor budding was semi-quantitatively scored for each tumor based on the maximum number of microscopic fields with tumor budding at the invasive front (illustrated in Figure 1). A tumor bud was defined as an isolated cluster of <5 tumor cells (including single tumor cells) completely surrounded by stroma and lacking gland lumen formation. A “tumor budding field” was defined as a 20X microscopic field (Olympus BX45, Olympus Plan N 20X objective lens; measuring 0.785 mm²) with 5 or more tumor buds based on the definition of Ueno et al.²⁵ We counted the number of budding fields in each tumor section and classified individual cases using the following cutoffs established *a priori*: no budding fields, 1 to 2 budding fields, or 3 or more budding fields based on the scoring methodology of Ohike et al.²² To compensate for differences in sampling density (number of tumor blocks per cm of tumor), the final budding score for each case was based on the tumor section with the greatest number of tumor budding fields.

Tumor size was determined based on the measurement reported in the original surgical pathology report and then confirmed by microscopic examination and review of gross photographs when necessary. For small tumors or tumors not grossly apparent, the size was determined by measuring the largest microscopic cross sectional size. For multifocal tumors, the size of the largest single focus was recorded. We used the median tumor size in our cohort (2.0 cm) as a cutoff for assessing risk of nodal metastasis. This cutoff has been previously reported as a risk factor for nodal metastasis.³

Tumor grade was determined based on current College of American Pathologists reporting guidelines for esophageal adenocarcinoma and the 2010 World Health Organization criteria for grading colorectal adenocarcinoma:²⁹ tumors composed of >95% isolated tubular glands were considered well-differentiated, those composed of 50% to 95% glandular structures were considered moderately differentiated, and those composed of <50% glands (i.e. > 50% solid growth or individual cells) were considered poorly-differentiated.

Angiolymphatic invasion was defined as tumor cells within endothelium-lined vascular or lymphatic spaces on H&E section within or adjacent to the primary tumor. Angiolymphatic invasion was not scored in the periesophageal or perigastric fat. Borderline cases were considered negative.

Submucosal invasion (stage T1b) was defined as definite extension of the tumor beyond the true muscularis mucosae and into the submucosa; tumors not satisfying this criterion were classified as intramucosal (pT1a). We further stratified depth of invasion according to Liu et

al.⁴ intramucosal adenocarcinomas were classified T1a-superficial if they were confined to the lamina propria, otherwise they were classified as T1a-deep. Submucosal adenocarcinomas were classified as T1b-superficial if they invaded less than half the thickness of the submucosa at the point of invasion and T1b-deep if they invaded more than half the submucosal thickness.

The tumor location, presence of lymph node metastases and the total lymph node count were determined based on the original surgical pathology report and confirmed on review. Tumors involving the esophagus or proximal stomach that invaded across the anatomic gastroesophageal junction were categorized as gastroesophageal junction adenocarcinomas. Tumors located in the esophagus that did not involve the anatomic gastroesophageal junction were categorized as esophageal. We did not include primary gastric carcinomas in the study. Barrett's esophagus was considered present if esophageal intestinal metaplasia was confirmed histologically in the esophagectomy specimen or in pre-operative biopsies. Age and sex, type of operation and presence of Barrett's esophagus were recorded from surgical pathology reports and the patient's electronic medical record. This study was approved by the University of Pittsburgh Institutional Review Board.

Statistical Analysis

Differences in categorical variables were analyzed by chi-squared or Fischer's exact tests. Differences in continuous variables were evaluated by Mann-Whitney U Test. A multivariate logistic regression model was used to identify independent predictors of nodal metastasis. Variables that were found to be statistically significant in the univariate analysis were included in the multivariate analysis. Overall survival was determined from the time of esophagectomy until death—as documented in the clinical record or Social Security Death Index—or the time of last known clinical follow up. Time to recurrence was defined as the time from esophagectomy to first recurrence or the last clinical evaluation for recurrence. Local, regional and distant recurrences were all included. The survival estimates for tumor budding groups were characterized using Kaplan-Meier curves, with statistical differences tested using the log-rank test. Cox proportional hazards analysis was used to control for other known predictors of survival in multivariate analysis. Patients who died as a result of complications from surgery (survival < 3 months) were excluded from the analysis. All tests were 2-sided and statistical significance was defined as a *p*-value <0.05.

RESULTS

Patient Characteristics

The median age of patients included in the study was 66 years (interquartile range 60–74) and the majority (n=179; 85%) were men. Most patients (n=176; 85%) underwent minimally invasive esophagectomy³⁰ and the remainder underwent an open (n=22; 11%) or hybrid procedure (n=8; 4%), most often a transhiatal esophagectomy (n=15; 7%); the operative approach was not specified in the available medical record for 4 patients. Margins of resection were negative in all cases. There were 125 (60%) tumors located at the gastroesophageal junction and 85 (40%) tumors located in the esophagus (79 lower

esophagus, 6 middle esophagus). Barrett's esophagus was present in 92% (n=193) of patients.

Overall Pathologic Characteristics

Two thirds of the superficial esophageal adenocarcinomas (n=138, 66%) invaded into the submucosa (T1b), almost equally divided between T1b-superficial and T1b-deep (Table 1). The remaining intramucosal adenocarcinomas (T1a) were almost equally divided between T1a-superficial and T1a-deep. Our assessment of T stage (T1a vs. T1b) agreed with the original diagnosis in 199/210 (95%) of cases. On review, 6 cases were upstaged (T1a to T1b) and 5 cases were downstaged (T1b to T1a).

Nearly two thirds (138/210) of cases were graded as moderately differentiated and 15% (32/210) were graded poorly differentiated (Table 1). Angiolymphatic invasion was identified in 36/210 cases (17%) (Table 1). The median tumor size in our cohort was 1.95 cm (interquartile range, 1.0–3.0 cm).

Lymph node metastases were present in 20% (n=41) of all cases. Of those with node metastases, the majority of cases (26/41, 63%) had 1–2 positive lymph nodes (stage pN1). The median number of lymph nodes examined at esophagectomy was 20 (interquartile range, 13–28) and did not significantly differ between the node positive and the node negative groups (Table 1).

Some degree of tumor budding was seen in 93/210 cases. It was focal (1–2 20X fields) in 16% (n=34) and extensive (≥ 3 20X fields) in 28% (n=59).

Association between Other Tumor Characteristics and Tumor Budding

Ninety-five percent of superficial esophageal adenocarcinoma with extensive tumor budding (≥ 3 20X budding fields) invaded either the superficial or deep submucosa compared to only 54% of those with none or focal budding ($p<0.001$, Table 2). Likewise, tumors with extensive tumor budding were more likely to be poorly differentiated, angioinvasive and larger than 2 cm ($p<0.001$ for all, Table 2). The association with tumor grade is noteworthy. Though extensive tumor budding was more common in high grade tumors (present in 23/32), it was also present in 36/178 well to moderately differentiated tumors graded according to College of American Pathologists/World Health Organization criteria (Table 2; examples in Figure 2).

Extensive Tumor Budding and Angiolymphatic Invasion are Independent Risk Factors for Lymph Node Metastasis

We next evaluated the association of primary tumor characteristics with the prevalence of nodal metastasis. Submucosal invasion ($p<0.001$), higher tumor grade ($p=0.003$), extensive tumor budding ($p<0.001$), angiolymphatic invasion ($p<0.001$), and tumor size ≥ 2 cm ($p<0.001$) all were associated with a significantly higher prevalence of nodal metastasis (Table 1). For instance, 41% (24/59) of tumors with extensive tumor budding were metastatic to regional lymph nodes, compared to 10% (12/117) of tumors with no tumor budding and 15% (5/34) of tumors with focal tumor budding (Table 1). The difference in

nodal metastasis between tumors with none versus focal tumor budding (1–2 fields) was not significant ($p=0.538$). We therefore used extensive tumor budding (at least 3 tumor budding fields) as a cutoff in the subsequent multivariate analysis as this was associated with a significantly higher frequency of nodal metastasis compared to all tumors without extensive budding ($p<0.001$, Table 1).

We also further evaluated depth of invasion and found that there was no difference in the prevalence of nodal metastasis when comparing superficial intramucosal (T1a) esophageal adenocarcinoma to deep T1a esophageal adenocarcinoma (Table 1). Given the limited number of cases with metastases in this group ($n=3$), this finding was not unexpected. However, depth of invasion into the submucosa did show a weak association with the prevalence of node metastasis ($p=0.049$) as 37% of esophageal adenocarcinoma with deep submucosal invasion had node metastasis compared to 22% of superficial submucosal esophageal adenocarcinoma (Table 1). Given these findings we categorized depth of invasion for the subsequent multivariate analysis as T1a, T1b superficial and T1b deep.

Because of the significant association between tumor budding and other variables known to predict lymph node metastasis, we performed a multivariate logistic regression analysis to determine which risk factors were independent predictors of nodal metastasis. In the univariate logistic regression, the presence of extensive tumor budding increased the odds of nodal metastasis 5.4-fold (95% confidence interval 2.6–11.1, $p<0.001$, Table 3). Similarly, the odds of nodal metastasis were significantly increased in the presence of submucosal invasion (either superficial or deep), high tumor grade, angiolymphatic invasion, and tumor size ≥ 2 cm (Table 3). In the multivariate logistic regression, extensive tumor budding (≥ 3 budding fields) was independent of other risk factors and associated with a 2.5-fold (95% confidence interval 1.1–6.3, $p=0.039$) increased risk of nodal metastasis (Table 3). Angiolymphatic invasion ($p=0.049$) also remained an independent predictor when controlling for other variables. Tumor size ≥ 2 cm showed a trend toward increased risk of nodal metastasis ($p=0.061$), but depth of invasion ($p=0.298$) and high tumor grade ($p=0.716$) did not.

The Effect of Tumor Budding on the Risk of Lymph Node Metastasis in Combination with Other Risk Factors

To illustrate how tumor budding modifies the risk of nodal metastasis associated with other predictors, we classified tumors using combinations of risk factors (Figure 3). When tumors are grouped based on tumor grade and tumor budding, 15/36 (42%) of low grade (well to moderately differentiated) tumors with extensive tumor budding were node positive compared to only 14/142 (10%) of low grade tumors without extensive budding ($p<0.001$, Figure 3). Interestingly, we found no difference in the rate of node metastasis between low grade tumors with extensive tumor budding and high grade tumors ($p=0.726$).

While superficial submucosal invasion (T1b-superficial) was associated with a 21% rate of nodal metastasis overall (Table 1), the rate of nodal metastasis is significantly influenced by tumor budding. Eight of 16 (50%) T1b-superficial cancers with extensive tumor budding were metastatic versus only 7/56 (13%) of those without extensive budding ($p=0.003$, Figure

3). Intramucosal cancers seldom metastasized; consequently, there was insufficient statistical power for comparison of subgroups.

In cases without angiolymphatic invasion, 13/37 (35%) with tumor budding were node positive, compared to only 11/137 (8%) with neither risk factor ($p<0.001$, Figure 3). Lastly, extensive tumor budding was associated with a significantly increased rate of nodal metastasis in small T1 esophageal adenocarcinoma (≤ 2 cm) and large T1 esophageal adenocarcinoma (> 2 cm, Figure 3).

Tumor Budding is a Poor Prognostic Factor with Respect to Overall Survival and Time to Recurrence

Out of the entire cohort of 210 patients, 194 were included in the survival analysis. There were 16 patients excluded because they died within 3 months of surgery ($N=13$) or had fewer than 3 months clinical follow-up recorded ($N=3$). Among the included patients, there were 72 deaths. The median follow-up interval was 44 months.

Because of the association with nodal status, tumor budding is expected to be a poor prognostic factor. When patients were stratified based on the extent of tumor budding, there was a significant difference in survival associated with the extent of tumor budding. Seventy-nine percent of patients with no tumor budding were alive at 5 years, compared to 71% of patients with focal budding and 37% of those with extensive tumor budding ($p<0.0001$, Figure 4A). In univariate Cox proportional hazards analysis extensive tumor budding, submucosal invasion (T stage), higher N stage, higher patient age and undergoing an operation other than minimally invasive esophagectomy were all significant adverse prognostic factors (data not shown). We performed a multivariate analysis to determine whether the negative effect of tumor budding on survival was independent of these other factors. Extensive tumor budding was associated with a 3.3-fold increased risk of death (95% confidence interval 1.5–7.4, $p=0.004$), after controlling for these other prognostic variables in the multivariate analysis.

Time to recurrence is a more specific indicator of aggressive tumor behavior in superficial gastroesophageal cancer because of its relatively favorable prognosis. In the 194 patients, there were 31 recurrences (25 with distant metastatic recurrence, 2 with only regional lymph node recurrence and 4 with only local anastomotic recurrences). Median time to last clinical evaluation for recurrence was 30 months. In patients with a documented recurrence, the median time to recurrence was 14 months (range 4–50 months). At 24 months, only 5% of patients without tumor budding had recurrence. In comparison, 19% of those with focal budding and 36% of those with extensive tumor budding had developed recurrence by 24 months ($p<0.0001$, Figure 4B). When patients with focal budding were compared to those with no tumor budding, the modest difference in time to recurrence was not significant ($p=0.096$). In univariate analyses, both submucosal invasion (T stage) and higher N stage were risk factors for tumor recurrence (data not shown), so we also performed a multivariate Cox proportional hazards analysis to control for these factors. Extensive tumor budding was associated with a 3.2-fold increased risk of recurrence (95% confidence interval 1.4–7.0, $p=0.005$), independent of T and N stage in the multivariate analysis.

DISCUSSION

Predicting nodal metastasis is of critical importance in superficial esophageal adenocarcinoma because this risk assessment is the primary determinant of the therapeutic approach. Therapy can range from endoscopic resection of the tumor with close follow-up for low risk tumors to neoadjuvant chemotherapy and radiation followed by esophagectomy for those thought to possess a high metastatic risk. The results of our study show that tumor budding is an independent predictor of nodal metastasis in superficial esophageal adenocarcinoma. In addition, it is a poor prognostic factor for overall survival and time to recurrence independent of other important survival factors.

Previously, tumor budding has been extensively studied in colorectal adenocarcinoma, where it is an important prognostic feature associated with lymph node metastasis, local recurrence, and cancer-related death particularly in American Joint Committee on Cancer stage 1^{25, 31–39} and stage 2^{21, 37, 40–43} disease. In a widely cited study, tumor budding along with high tumor grade and vascular invasion were independent prognostic features associated with lymph node metastasis in submucosally invasive colorectal adenocarcinoma.²⁵ Tumor budding has also been shown to be an independent prognostic factor associated with overall survival in esophageal squamous carcinomas.^{23, 44}

With regard to tumor budding in esophageal adenocarcinoma, one prior study found tumor budding to be a prognostic factor for overall survival in a cohort of patients with esophageal cancer which included 287 adenocarcinomas and 69 squamous cell carcinomas.²⁴ This association with outcome persisted even when controlling for other prognostic factors in a multivariate analysis. However, that study did not evaluate the association of tumor budding with nodal metastasis and included only 38 T1 tumors. A more recent study published in abstract form showed that tumor budding was associated with increased likelihood of tumor recurrence and nodal metastasis in 42 T1 esophageal adenocarcinomas.²⁸ Although that study assessed superficial esophageal adenocarcinomas for other histologic risk factors (tumor size, grade, angioinvasion and depth of invasion), it was not sufficiently powered to evaluate whether tumor budding was an independent prognostic factor when controlling for other predictors of nodal metastasis and recurrence.

One challenge in studying the prognostic significance of tumor budding is the absence of a universally accepted definition that is both qualitative (small clusters of cells at the invasive front) and quantitative (how much budding is needed to increase the risk). The most common method proposed by Ueno uses a cutoff of 5 or more individual buds in a 20X field (area of 0.785 mm²) at the invasive front.²⁵ Other studies have proposed more labor-intensive quantitative scoring systems, requiring precise numerical scoring of multiple high power fields followed by averaging of tumor bud counts.^{32, 45–48} Although this is likely to produce an accurate description of the extent and variability of tumor budding, we feel that a simpler approach is required for routine clinical practice.

We therefore selected a semi-quantitative method previously used to evaluate ampullary adenocarcinomas that relies on counting the number of 20X fields with more than 5 tumor buds.²² In that study, tumors with 3 or more 20X tumor budding fields were more likely to

metastasize than those with fewer. The general utility of this scoring method and quantitative cutoff is supported by our finding in superficial esophageal adenocarcinoma that tumors with focal tumor budding (1 to 2 budding fields) did not have a greater risk of lymph node metastasis than those with no budding, but tumors with 3 or more budding fields had a significantly higher risk of nodal metastasis and worse survival outcomes. It is worth noting that focal tumor budding did show a trend toward a higher risk of recurrence, but the difference in survival compared to those with no tumor budding was not significant. It may be prudent to regard focal tumor budding as an indeterminate risk factor in T1 esophageal adenocarcinoma.

The issue of whether to score tumor budding on the basis of H&E stained sections or a broad spectrum keratin immunostain has been discussed.⁴⁵ In preparation for this study we did review serial H&E sections along with sections stained with pancytokeratin to confirm our impression of tumor budding (data not shown). This anecdotal experience convinced us that in the large majority of cases, a keratin stain would not be required to identify and score tumor budding and it is unlikely in clinical practice that pathologists would prefer to obtain keratin immunostains in all cases as a condition for scoring tumor budding. We, therefore, elected not to use a keratin stain in this study. We would echo the practical suggestion made by Mitrovic et al. that a pankeratin immunostain be reserved for difficult cases (i.e. in the presence of abundant inflammation) to confirm the histologic identification and quantitation of tumor budding.⁴⁹

The College of American Pathologists/World Health Organization criteria for grading adenocarcinomas of the gastrointestinal tract are based on glandular architecture. Using these criteria, extensive tumor budding may increase the tumor grade (i.e. if the tumor budding pattern represented over half of the tumor it would be graded as poorly differentiated). However, only 38% of the cases in our series with extensive tumor budding were poorly differentiated by College of American Pathologists/World Health Organization criteria. The remainder were low grade (well or moderately differentiated), illustrating that a tumor can show prominent glandular architecture and have extensive tumor budding by the Ohike criteria.²² In fact, in our subgroup analysis we show that low grade tumors with tumor budding have a similar risk of nodal metastasis as poorly differentiated adenocarcinomas. Because tumor budding and architectural grade are interrelated, future studies should evaluate the utility of a grading system that combines glandular architecture and tumor budding.

It may be surprising to some that depth of invasion was not an independent risk factor for nodal metastasis because it is generally regarded as an essential predictor of nodal metastasis in routine clinical practice. Our data confirms numerous studies that show submucosal invasion is a risk factor for nodal metastasis^{4, 8, 11, 16, 50–54} when analyzed as a single variable. Because of the number of cases in our series, we were able to control for the most common additional risk factors and show that the risk associated with submucosal invasion is significantly influenced by other variables. In other words, tumors that invade into the submucosa are more likely to be larger than 2 cm, to be poorly differentiated and to show angiolymphatic invasion.^{3, 4} We also show that they are also more likely to have extensive tumor budding. These observations beckon for a system of risk stratification that accounts

for multiple variables in addition to depth of invasion. Such an approach could be clinically useful, as suggested by studies showing it is possible to safely treat carefully selected submucosal esophageal adenocarcinoma based on the absence of other high risk pathologic features.⁵⁵

We noted that extensive tumor budding is almost never the only risk factor for nodal metastasis in a given tumor, yet our multivariate analysis demonstrates that its predictive significance is independent of other commonly accepted risk factors. We illustrate how the relationship between other risk factors and nodal metastasis is modified by the extent of tumor budding such that T1b-superficial, low grade tumors and tumors without angiolymphatic invasion had dramatically higher rates of nodal metastasis in the presence of extensive tumor budding.

We chose to focus on the subset of patients with T1 esophageal adenocarcinoma in this study because predicting nodal metastasis in this patient population is of utmost clinical relevance. The significance of tumor budding may be different in more advanced esophageal adenocarcinoma with a significantly higher likelihood of nodal metastasis. Tumor budding and other pathologic risk factors for nodal metastasis could be assessed prior to definitive therapy in endoscopic resection specimens or in biopsies. Intratumoral budding identified in biopsy specimens of rectal carcinoma was recently shown to associate with poor response to neoadjuvant treatment.⁵⁶ Our study did not address this question and future studies to evaluate the predictive ability of these pathologic features in the pre-operative setting are warranted.

Lastly, in addition to tumor budding, we focused this report on the most commonly cited pathologic predictors of nodal metastasis in superficial esophageal adenocarcinoma. We did not evaluate all previously reported prognostic factors such as human epidermal growth factor receptor 2 amplification, epidermal growth factor receptor amplification, inflammatory host response, and perineural invasion, among others. To incorporate additional variables into a multivariate analysis would require a larger number of cases.

In summary, we have shown that tumor budding is an independent risk factor for nodal metastasis in superficial esophageal adenocarcinoma when evaluated by a semi-quantitative scoring method on routine histologic sections. Tumor budding should be evaluated in superficial esophageal adenocarcinoma as part of a comprehensive prognostic assessment.

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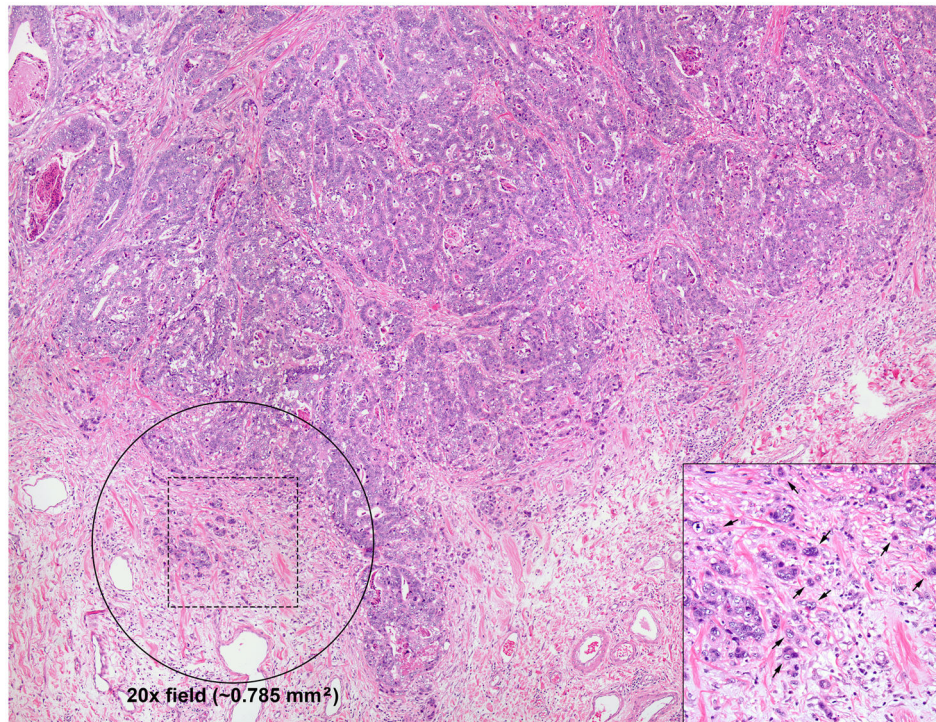
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**Figure 1. Scoring of tumor budding**

This figure illustrates the semi-quantitative scoring criteria used to assess tumor budding in each case. Even at relatively low magnification, areas that are likely to represent tumor budding are apparent due to the indistinct border between tumor and stroma at the advancing edge of the tumor. The circle represents the approximate size of a single 20X high power field (0.785 mm^2). For this study we counted the number of 20X fields with more than 5 tumor buds (“tumor budding fields”). Individual tumor buds were defined as clusters of fewer than 5 tumor cells lacking gland lumen formation at the advancing edge of the tumor. Several of the individual tumor buds in the hashed square are indicated by arrows (inset). As detailed in the methods, cases were classified as no tumor budding (no 20X fields with more than 5 tumor buds); focal tumor budding (1–2 20X budding fields); or extensive tumor budding (≥ 3 20X budding fields) according to Ohike et al. (reference 22).

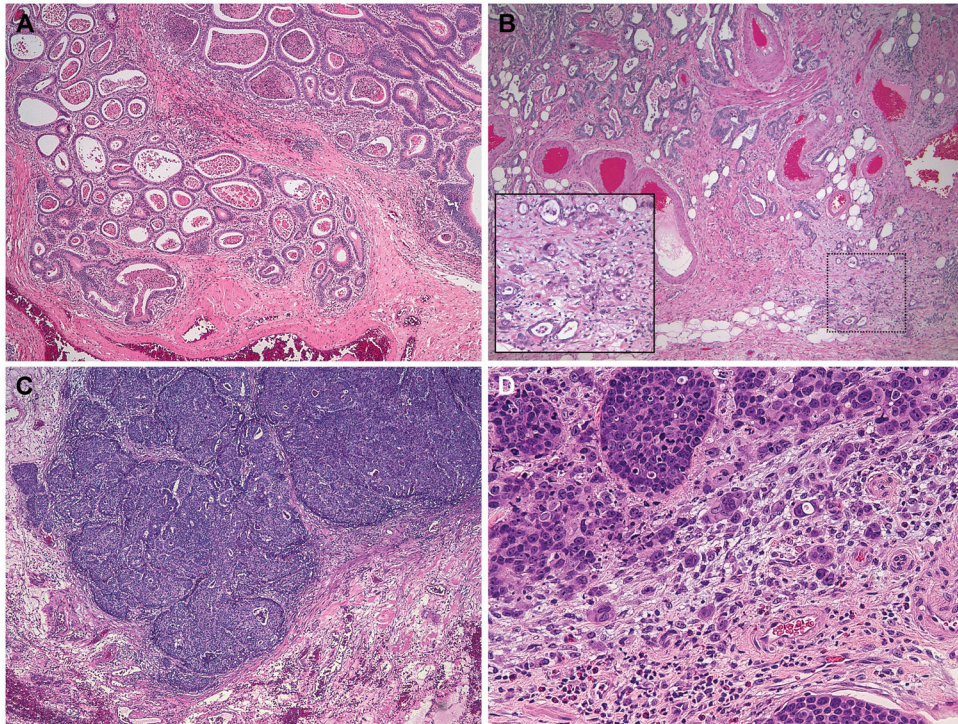


Figure 2. Representative examples of tumor grade and budding

(A) Low grade tumor without tumor budding. The well-formed glands with luminal necrosis invade the submucosa and are surrounded by a mild inflammatory infiltrate at low power (~40X magnification). (B) A predominantly low grade adenocarcinoma with tumor budding. Budding is best seen in the inset at the advanced edge of the tumor as it infiltrates the submucosal fat (40X magnification, inset 200X magnification). (C) A poorly differentiated adenocarcinoma that lacks tumor budding. There is a fairly distinct “pushing” interface between the advancing edge of the tumor and the surrounding stroma (40X magnification). (D) A poorly differentiated adenocarcinoma shows prominent tumor budding in this high power field (200X magnification).

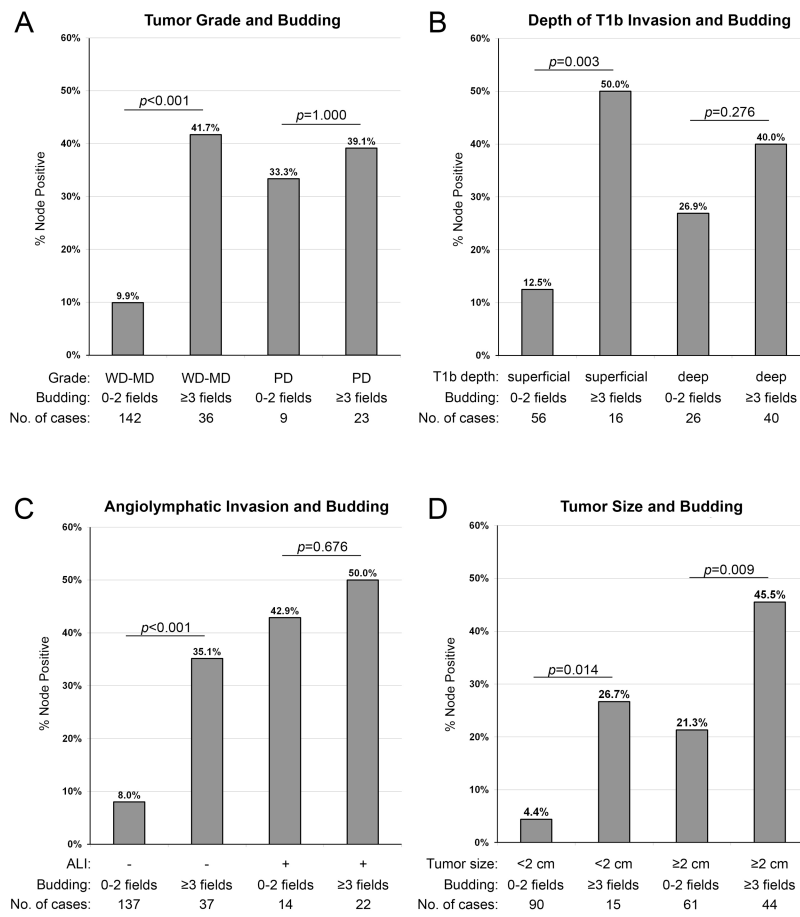


Figure 3. Bar graphs illustrating the prevalence of node metastasis in cases classified based on the presence of tumor budding and other risk factors.

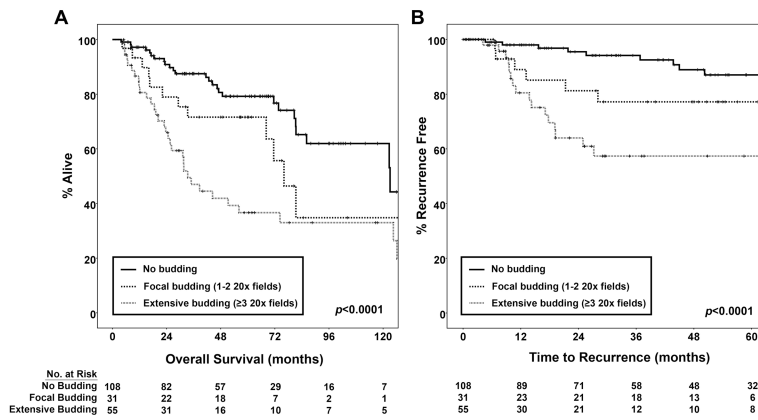


Figure 4. Kaplan-Meier curves for overall survival and time to recurrence in patients stratified by the extent of tumor budding

The graphs demonstrate significant differences in the survival functions. Extensive tumor budding was associated with worse overall survival (A) and accelerated time to recurrence (B).

Table 1

Lymph node status associated with pathologic features.

	All Cases N	Node negative N (%)	Node positive N (%)	<i>p</i> -value
Tumor Stage				
T1a (intramucosal)	72	69 (96)	3 (4)	<0.001
T1b (submucosal)	138	100 (72)	38 (28)	
Depth of Invasion, Intramucosal Only				
T1a superficial	35	34 (97)	1 (3)	1.000
T1a deep	37	35 (95)	2 (5)	
Depth of Invasion, Submucosal Only				
T1b superficial	72	57 (79)	15 (21)	0.049
T1b deep	66	43 (65)	23 (35)	
Lymph Node Stage				
N0 (0 positive lymph nodes)	169	169 (100)	0 (0)	N/A
N1 (1–2 positive lymph nodes)	26	0 (0)	26 (100)	
N2 (3–6 positive lymph nodes)	10	0 (0)	10 (100)	
N3 (>6 positive lymph nodes)	5	0 (0)	5 (100)	
Lymph nodes examined (median, interquartile range)	20 (13–28)	19 (14–28)	20 (13–28)	0.969
WHO Tumor Grade				
Well differentiated	40	38 (95)	2 (5)	0.003
Moderately differentiated	138	111 (80)	27 (20)	
Poorly differentiated	32	20 (62)	12 (38)	
Tumor Budding				
None	117	105 (90)	12 (10)	<0.001
Focal (1 to 2 budding fields)	34	29 (85)	5 (15)	
Extensive (≥ 3 budding fields)	59	35 (59)	24 (41)	
Tumor Budding				
None or Focal (< 3 budding fields)	151	134 (89)	17 (11)	<0.001
Extensive (≥ 3 budding fields)	59	35 (59)	24 (41)	
Angiolymphatic Invasion				
Absent	174	150 (86)	24 (14)	<0.001
Present	36	19 (53)	17 (47)	
Tumor Size				
< 2 cm	105	97 (92)	8 (8)	<0.001
≥ 2 cm	105	72 (69)	33 (31)	

Table 2

Pathologic features according to tumor budding status.

	Tumor Budding		<i>p</i> -value
	None or Focal (0–2 budding fields) N (%)	Extensive (≥ 3 budding fields) N (%)	
Depth of Invasion (T stage)			
Intramucosal (T1a), superficial	34 (23)	1 (2)	< 0.001
Intramucosal (T1a), deep	35 (23)	2 (3)	
Submucosal (T1b), superficial	56 (37)	16 (27)	
Submucosal (T1b), deep	26 (17)	40 (68)	
WHO Tumor Grade			
Well differentiated	39 (26)	1 (2)	< 0.001
Moderately differentiated	103 (68)	35 (59)	
Poorly differentiated	9 (6)	23 (39)	
Angiolymphatic Invasion			
Absent	137 (91)	37 (63)	< 0.001
Present	14 (9)	22 (37)	
Tumor Size			
< 2 cm	90 (60)	15 (25)	< 0.001
≥ 2 cm	61 (40)	44 (75)	

Table 3

Univariate and multivariate analysis of histologic predictors of node metastasis

	Univariate Analysis		Multivariate Analysis	
	Odds Ratio (95% confidence interval)	<i>p</i> -value	Odds Ratio (95% confidence interval)	<i>p</i> -value
Depth of Invasion (T stage)				
Intramucosal (T1a)	Reference		Reference	
Submucosal (T1b), superficial	6.4 (1.8 – 23.2)	0.001	2.8 (0.7 – 11.6)	0.329
Submucosal (T1b), deep	13.0 (3.3 – 41.6)		2.6 (0.6 – 12.1)	
Tumor Grade				
Well or Moderately Differentiated	Reference		Reference	
Poorly Differentiated	3.1 (1.4 – 7.0)	0.007	1.3 (0.5 – 3.5)	0.582
Angiolymphatic Invasion				
Absent	Reference		Reference	
Present	5.6 (2.6 – 12.2)	< 0.001	2.5 (1.0 – 5.6)	0.049
Tumor Budding				
None or focal (0–2 fields)	Reference		Reference	
Extensive (≥ 3 fields)	5.4 (2.6 – 11.1)	< 0.001	2.5 (1.1 – 6.3)	0.039
Tumor Size				
< 2 cm	Reference		Reference	
≥ 2 cm	5.5 (2.4 – 12.7)	< 0.001	2.5 (1.0 – 6.6)	0.061