

HHS Public Access

Author manuscript *Mod Pathol.* Author manuscript; available in PMC 2018 January 21.

Published in final edited form as:

Mod Pathol. 2017 October; 30(10): 1402-1410. doi:10.1038/modpathol.2017.64.

Computer-assisted measurement of primary tumor area is prognostic of recurrence-free survival in stage IB melanoma patients

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Abstract

Current staging guidelines are insufficient to predict which patients with thin primary melanoma are at high risk of recurrence. Computer-assisted image analysis may allow for more practical and objective histopathological analysis of primary tumors than traditional light microscopy. We studied a prospective cohort of stage IB melanoma patients treated at NYU Langone Medical Center from 2002-2014. Primary tumor width, manual area, digital area, and conformation were evaluated in a patient subset via computer-assisted image analysis. The associations between histologic variables and survival were evaluated using Cox proportional hazards model. Logistic regressions were used to build a classifier with clinicopathological characteristics to predict recurrence status. Of the 655 patients with stage IB melanoma studied, a subset of 149 patient tumors (63 recurred, 86 did not recur) underwent computer-assisted histopathological analysis. Increasing tumor width (HR 1.17, p=0.01) and digital area (HR 1.08, p<0.01) were significantly associated with worse recurrence-free survival, while non-contiguous conformation (HR 0.57, p=0.05) was significantly associated with better recurrence-free survival. The novel histopathological classifier composed of digital area, conformation, and baseline variables effectively distinguished recurrent cases from nonrecurrent cases (AUC 0.733, 95% CI 0.647-0.818), compared to the baseline classifier alone (AUC 0.635, 95% CI 0.545–0.724). Primary tumor cross-sectional area, width, and conformation measured via computer-assisted analysis may help identify high-risk patients with stage IB melanoma.

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Disclosure/Duality of Interest The authors have no duality of interest to declare.

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stage IB melanoma; computer-assisted analysis; tumor area; tumor width; tumor conformation

Introduction

Melanoma incidence continues to increase at a rate of 3–7% per year.^{1–3} Increased public awareness and screening efforts have led to substantial stage migration at diagnosis approximately 70% of newly diagnosed melanomas are now thin (Breslow thickness 1.00 mm), invasive lesions.⁴ While most patients diagnosed with thin primary melanoma have excellent outcomes, it is estimated that 3–5% of patients with thin melanoma, the majority of whom are classified as stage IB, will go on to develop metastatic disease.^{5–7} According to the 2009 American Joint Committee on Cancer (AJCC) staging system, stage IB melanomas are defined as those thinner than 1.00 mm with presence of either ulceration or at least one mitotic figure per mm², or as those with tumor thickness of 1.01 to 2.0 mm and without ulceration.^{8, 9} The 10-year survival rate associated with stage IB melanoma is approximately 85%.⁹ Given that a substantial portion of stage IB melanomas will progress, identification of clinical and pathologic variables associated with high-risk stage IB disease is crucial.

The current AJCC staging system is insufficient to predict which patients with stage IB melanoma are at highest risk of recurrence. The AJCC criteria described above allow for substantial histopathological heterogeneity within stage IB melanoma patients. Further stratification of stage IB patients based on histopathological phenotypes may help identify high- and low-risk patient subsets. Previous studies have demonstrated significant associations of increasing tumor volume with higher risk of sentinel lymph node metastases and worse survival outcomes in melanoma.^{10–13} Given the logistical constraints of measuring tumor volume, most of these studies estimated volume based on the mathematical approximation for a hemiellipsoid, which typically require serial measurements from multiple thin tumor sections.^{10, 12} These methods can be difficult to standardize and exceedingly time-consuming, thus limiting their practicality in the clinical setting.^{10–16} Recent advances in digital pathology software, however, may now allow more rigorous and efficient histopathological tumor analysis, providing an opportunity to re-evaluate multidimensional tumor measurements in thin primary melanomas.^{17–19}

To improve risk stratification of thin melanoma, we investigated the prognostic value of multidimensional primary tumor measurements (e.g. cross-sectional area and width) in a large, prospective cohort of stage IB melanoma patients. We postulated that multidimensional primary tumor measurements may be useful in predicting which patients with thin primary melanoma are at highest risk of recurrence. Additionally, the use of digital pathology software tools may allow for more practical and objective evaluation of multidimensional primary tumor measurements than those methods utilized in prior studies.

Materials and Methods

Study population

We studied a cohort of patients who presented to the NYU Langone Medical Center with AJCC stage IB melanoma from August 2002 to May 2014 and were prospectively enrolled in the Interdisciplinary Melanoma Cooperative Group, a prospective clinicopathological biospecimen database.²⁰ This study was approved by the NYU Institutional Research Board (Study #10362) and all patients provided informed, written consent at the time of enrollment. For patients referred from outside institutions, the diagnosis of primary melanoma was confirmed by the analysis of biopsy specimens by two NYU pathologists. All primary melanoma patients underwent surgical treatment of the initial primary tumor.

Patient demographics, clinical data, and primary tumor characteristics were obtained from clinical charts and pathology reports. Variables assessed included patient sex, age at stage IB diagnosis, primary tumor thickness, primary tumor ulceration, mitotic rate (per mm²), and histological subtype. Date of stage IB diagnosis was defined as the biopsy date of the primary tumor. Recurrence-free survival (recurrence-free survival) was defined as the time from the date of pathological diagnosis of stage IB melanoma to the first recorded date of regional or distant metastases. Overall survival (OS) was defined as the time from the date of pathological diagnosis of stage IB melanoma until death. Recurrence and survival information were collected by active, prospective follow-up of all patients enrolled in the database.

Computer-assisted histopathological analysis

A subset of patients was selected for further examination of additional histological variables via computer-assisted image analysis. This group consisted of all patients with stage IB melanoma whose disease later recurred, along with a subset of patients with non-recurrent disease, matched for age, gender, histopathological subtype, thickness, ulceration, and mitotic rate. Patients were excluded if the corresponding H&E-stained slide of their primary tumor was unavailable. Primary tumor slides were scanned into digital images and independently reviewed by the Interdisciplinary Melanoma Cooperative Group dermatopathologist (NB). Slides were examined using computer-assisted analysis (Aperio, Vista, CA USA) to confirm primary tumor characteristics, as well as assess tumor width, manual tumor area, digital tumor area, and tumor conformation. Tumors were considered "contiguous" if the majority of the tumor was present in one coalesced mass, while those divided into multiple, separate aggregates were considered "non-contiguous." Tumor width (mm) was measured in the invasive dermal component in the greatest horizontal dimension. In non-contiguous tumors, the sum of widths of individual dermal tumors was measured. Manual tumor area (mm²) was estimated by multiplying the Breslow thickness (mm) by the width (mm) of the invasive dermal component. To calculate digital tumor area (mm²), the invasive dermal component of the tumors were manually outlined by the dermatopathologist and the tumor area calculated by the software.

Statistical analysis

We tested the association between histologic variables and recurrence-free survival and overall survival using Cox univariate and multivariate analysis; only variables significant in univariate analysis were included in the multivariate model. Linear regression analysis was performed to assess the relationship between manual and digital area for all patients, including patients subdivided by conformation. Using the research cohort, we first examined the differences in clinicopathological characteristics between patients with recurrent melanoma and those with non-recurrent disease using Wilcoxon/Mann-Whitney nonparametric test for continuous variables and Fisher's Exact test for categorical variables. We then use logistic regressions to build classifier with clinicopathological characteristics to predict recurrence status. The candidate predictors of the baseline model were restricted to age, gender, histological subtype, thickness, mitoses (yes or no) and ulceration status. The candidate predictors of the novel histopathological model include the selected variables in the baseline model and the candidate from the novel histopathological predictors, including width, conformation status (contiguous or non-contiguous) and digital area or manual area. Both models were selected by minimizing Akaike's information criterion of the logistic regression. We used a 10-fold cross validation approach to obtain an unbiased estimate of the area under the receiver operating characteristic (ROC) curve (AUC) of the selected model and its 95% confidence interval. The linear combination of the selected model predictors weighted by regression coefficients was defined as the risk score. A risk score cutoff using the Youden Index of the ROC curve was chosen to separate patients into highand low-recurrence risk groups. Kaplan-Meier survival curves and log-rank tests were used to compare the recurrence-free survival of the two groups. All statistical tests were performed by in R including packages "survival", "ROCR", "cvAUC", and "Optimal Cutpoints".

Results

Patient characteristics

Table 1 displays the demographic and primary tumor characteristics of the patient cohort (n=655). Median follow-up was 4.4 years. The median age at stage IB diagnosis was 58.0 years and 57.1% of patients were male. The median primary tumor thickness and mitotic rate were 0.98 mm and 1.0 per mm², respectively. Ulceration was present in 4.3% of primary tumors. The most common histological subtype was superficial spreading (67.3%). Overall, 77 (11.8%) patients experienced a recurrence of their disease, while 578 (88.2%) of patients remained in remission throughout the study period.

Increasing thickness, mitotic rate, and ulceration are significantly associated with survival outcomes

We performed survival analysis to assess the relationship between patient baseline variables and survival outcomes. In univariate analysis, increasing Breslow thickness (HR 4.63, p<0.01), presence of ulceration (HR 2.27, p=0.05), and increasing mitotic rate (HR 1.14, p<0.01) were significantly associated with worse Interdisciplinary Melanoma Cooperative Group Interdisciplinary Melanoma Cooperative Group recurrence-free survival. Increasing age (HR 1.02, p=0.04), Breslow thickness (HR 3.26, p<0.01), and mitotic rate (HR 1.14,

p<0.01) were significantly associated with worse overall survival. In multivariate analysis, Breslow thickness (HR 4.13, p<0.01) and ulceration (HR 2.44, p=0.04) remained significantly associated with worse recurrence-free survival. Breslow thickness (HR 2.54, p<0.01) and increasing mitotic rate (HR 1.10, p=0.02) remained significantly associated with worse overall survival.

Primary tumor width, digital area, and conformation are prognostic of worse recurrencefree survival

A subset of 149 patients were examined for digital histopathological variables and associations with survival (Table 2). Fourteen of the 77 patients with recurrent disease were excluded from sub analysis due to unavailability of primary tumor slides for review. In this group, 63 (42.3%) patients had disease that later recurred with regional or distant metastases, while 86 (57.7%) patients had disease did not recur. There were no significant differences in sex, age at diagnosis, tumor thickness, ulceration, or mitotic rate between patients with recurrent melanoma and those with non-recurrent disease.

The digital histopathological variables evaluated are illustrated in Figure 1. Manual area was estimated by multiplying the Breslow thickness by the width of the tumor (Figure 1A), while digital area was calculated via the Aperio software algorithm (Figure 1B). Tumors were classified as contiguous or non-contiguous, as displayed in Figure 1B and 1C. Tumor width was measured as the cummulative sum of widths of the tumor's invasive dermal components (Figure 1D). respectively. The median width, digital area, and manual area measurements of the entire research cohort were 3.73 mm, 2.10 mm², and 3.97 mm², respectively (Table 2). With respect to tumor conformation, 63.8% of primary tumors were considered contiguous, while 36.2% were classified as non-contiguous. Primary tumors that later recurred were significantly larger in width (4.41 mm vs. 4.19 mm; p=0.03) and digital area (2.47 mm² vs. 1.90 mm²; p=0.01) than those that did not recur. However, no significant differences in manual area (p=0.14) or tumor conformation (p=0.14) was observed between recurrent and non-recurrent groups.

To assess the relationship between digital histopathological variables and survival, statistical analysis was performed by Cox proportional hazard ratios (Table 3). Multivariate analysis demonstrated that increasing tumor width (HR 1.20, p<0.01) and digital area (HR 1.08, p<0.01) were significantly associated with worse recurrence-free survival, while non-contiguous conformation (HR 0.55, p=0.05) was significantly associated with better recurrence-free survival. There was a trend towards worse recurrence-free survival with increasing manual area, but this association did not reach significance (HR 1.08, p=0.07). No digital histopathological variables demonstrated a significant association with overall survival.

Novel histopathological signature predicts recurrence

Among the 149 patients in the research cohort, three were removed due to missing ulceration status, and one was removed due to missing digital area measurement. Using the baseline variables, we first developed a baseline prognostic model of melanoma recurrence, using the parameters of age at stage 1b diagnosis, thickness, and mitoses (yes or no). We

then built the prognostic model with the addition of novel histopathological predictors. This model includes digital area and conformation status (contiguous or non-contiguous) in addition to the variables in the baseline model. The novel histopathological classifier displayed an area under the receiver operating characteristic (ROC) curve (AUC) of 0.733 (95% CI 0.647–0.818) estimated via the 10-fold cross validation approach, effectively distinguishing recurrent cases from nonrecurrent cases. In contrast, the baseline classifier alone achieved AUCs of 0.635 (0.545, 0.724).

We defined risk scores for individual patients as the linear combination of model predictors weighted by regression coefficients. Based on the cutoff value selected by the Youden Index, the sensitivity was 78.7% and the specificity 72.6% (Figure 2). For patients predicted to be negative for recurrence by the signature (negative predictive value), 17.6% (13/74) recurred and 82.4% (61/74) patients have not recurred, which supports the clinical sensitivity of the signature. For patients predicted to be positive for recurrence by the signature (positive predictive value), 67.6% (48/71) patients have recurred and 32.4% (23/71) patients have recurred.

To assess the ability of the signature to stratify patients by recurrence-free survival, highand low-risk groups were plotted in Kaplan-Meier survival curves and statistical analysis performed by log-rank tests (Figure 3). High- and low-risk groups displayed clear differences in progression-free survival (P < .001), but not significant for OR (P = 0.09). These analyses underscore the robustness of our classifier in predicting melanoma patient recurrence outcome.

We also built a prognostic model including manual area (substituted for digital area) and conformation status (contiguous or non-contiguous) in addition to the variables in the baseline model. This novel histopathological displayed an AUC of 0.691 (95% CI, 0.602– 0.780) estimated via the 10-fold cross validation approach. Therefore, we believe that the signature incorporating digital area is more effective.

Manual and digital primary tumor area are highly correlated, especially in contiguous tumors

We then investigated whether manual area may be an adequate surrogate for digital area in settings without access to digital pathology software. We evaluated the relationship between manual and digital area, which showed a significant correlation between the two measurements (p<0.01; Figure 4A). We then substratified patients into groups of either contiguous (n=95) or non-contiguous (n=54) tumors. Patients with contiguous tumors had a stronger correlation between manual and digital area measurements than did those with non-contiguous tumors (estimate 0.75 vs. 0.49, respectively; p<0.01; Figure 4B).

Discussion

Our study demonstrated the ability of computer-assisted histopathological analysis to identify novel prognostic variables in patients with stage IB melanoma. Using computer-assisted image analysis, we identified primary tumor intradermal width, digital area, and conformation as prognostic variables that may help predict which patients with stage IB

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melanoma are at highest risk of recurrence. Furthermore, our data demonstrated that a signature composed of digital tumor area and tumor conformation, combined with baseline variables, had superior prognostic capabilities compared to baseline variables alone and accurately stratified patients by recurrence-free survival. We also showed that manual tumor area (Breslow thickness \times width), which can be measured digitally or via light microscope, may potentially be a suitable substitute for digital area in clinical settings without access to digital pathology software.

Routine surveillance for patients with thin primary melanoma is cost-ineffective due to their generally low recurrence rate.²¹ Despite their favorable prognosis, thin melanomas comprise up to 70% of new melanoma diagnoses and account for over 25% of all melanoma deaths.⁴ Thus, targeted surveillance for high-risk thin primary melanoma patients has the potential to make a substantial impact on patient mortality. Defining prognostic variables that risk-stratify patients with thin primary melanoma, however, is difficult as these tumors tend to have similar and generally favorable histopathological primary tumor characteristics. Therefore, we utilized computer-assisted image analysis to create a predictive signature of baseline histopathological variables, cross-sectional tumor area, and conformation that effectively risk-stratifies stage IB melanoma patients by recurrence-free survival.

Our results highlight the prognostic utility of tumor cross-sectional area in patients with thin primary melanoma. Previous studies have also demonstrated the prognostic value of digitally calculated tumor area. Two studies of cross-sectional area in primary melanomas of all thicknesses (T1–T4) found that increasing tumor area was significantly associated with worse recurrence-free survival and overall survival.^{15, 16} However, in both studies, the prognostic value of cross-sectional area was highly correlated with thickness, and provided little additional prognostic information to the current paradigm. To mitigate the confounding effects of thickness encountered in prior studies, we restricted our study population to patients with stage IB melanoma, limiting the clinical and histopathological heterogeneity of our patient sample. Since there is still an inherent range of thicknesses (0.0 to 2.0 mm) in stage IB tumors, we then selected a research cohort composed of recurrent and non-recurrent cases matched for Breslow thickness (as well as age, gender, mitoses, and ulceration). Therefore, the association of increasing tumor area with worse recurrence-free survival reported in this study is less likely to be due to confounding with Breslow thickness.

In clinical settings lacking digital pathology software, manual area may be a suitable surrogate for digital area, particularly in patients with contiguous tumors. To extend the generalizability of our results, we also measured the manual area (Breslow thickness × intradermal width) of each tumor, which can be performed using traditional light microscopy. However, the associations between manual area and recurrence outcomes did not reach statistical significance. Similarly, the predictive model incorporating manual area did not demonstrate superior prognostic capability over that of the baseline model. This may be because manual area is directly dependent on thickness and, thus, was affected by our efforts to control for thickness. Therefore, we recognized that manual area might still provide useful prognostic information when digital pathology software is not available. Linear regression analysis of manual and digital area found a significant correlation between the two measurements in all patients. Further, when patients were substratified by their

Our novel analysis of tumor conformation demonstrated that patients with contiguous tumors had a worse recurrence-free survival than those with non-contiguous tumors. Additionally, tumor conformation was included in the novel histopathological signature that accurately stratified patients by recurrence-free survival, and was superior to that of the baseline model. This is the first study to report primary tumor conformation as a potential prognostic variable for melanoma patients. Our findings suggest that there may be underlying biological differences between contiguous and non-contiguous primary tumors. It is also likely, however, that primary tumor conformation is colinear with tumor area and cell proliferation markers, and thus may not offer additional prognostic information to the current staging paradigm. Further investigation of tumor conformation in multivariate models is needed to fully elucidate its prognostic value in thin primary melanoma.

Our study also found the histologic measurement of the primary tumor's intradermal width to be prognostic of recurrence and worse recurrence-free survival. Tumor surface-width (e.g. tumor diameter) measured *in vivo* is typically used as a screening tool in melanoma, where suspicious lesions with a diameter greater than 6 mm have a higher likelihood of being melanoma than those less than 6 mm.^{22–24} In regard to prognostic value, past studies have noted a correlation between the primary tumor diameter and survival.^{25–30} Breslow's landmark 1970 study found that no primary melanomas less than 5 mm in clinical diameter recurred or metastasized.³⁰ In our study, the median width of the invasive dermal component determined to be statistically significant in patients with recurrent disease was 4.19 mm, as compared to 3.41 mm in those who did not recur (p=0.03). We recognize that the width of the invasive component of melanoma is not an exact correlate of clinical diameter. While the clinical diameter can be accurately measured, the histologic determination of dermal width is potentially confounded by tissue shrinkage, grossing technique (i.e. the axis upon which the specimen was sectioned) and variability in the histologic planes of sections examined.

Survival analysis of the baseline variables yielded expected results that are consistent with those of the current AJCC guidelines. Interestingly, our analysis of mitotic rate as a continuous variable showed that increasing mitotic rate was independently associated with worse survival. This finding is consistent with multiple recent studies that also demonstrate the powerful independent effect of mitotic rate on patient outcomes, including a multi-institutional study by *Thompson et al.* of 13,296 patients with stage I and II melanoma.^{19, 33–36} Future staging guidelines should reconsider the use of mitotics as a binary criterion (present vs. absent), as there is sufficient evidence that higher mitotic rates are associated with worse outcomes.

There were several limitations to this study. First, we focused our investigation of thin melanoma on patients with AJCC stage IB disease because this patient population is at higher risk of poor outcomes and still lacks clear management guidelines. However, this limited our sample to patients who were histopathologically node-negative or who did not undergo sentinel lymph node biopsy at all, introducing possible sampling bias. This also prevented evaluation of sentinel lymph node positivity as an outcome measure. Second, we

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evaluated novel histopathological characteristics in a non-random subset of similar recurrent and non-recurrent patients in order to control for certain variables and increase the relative number of recurrence events. This resulted in a research cohort that was older in age, had thicker primary tumors with higher mitotic rates; however this is expected due to the increased prevalence of recurrence events. Third, the research cohort was histopathologically reviewed by only one dermatopathologist, thus inter-observer reliability could not be assessed. Finally, we did not split the data into discovery and test cohorts, as there were only 63 stage IB patients whose disease recurred and whose slides were available for review. We attempted to mitigate this limitation, however, by using cross-validation to obtain and unbiased AUC estimate. Therefore, independent validation of our model in a large prospective cohort of stage IB patients is necessary to assess the robustness of our prognostic signature prior to translation into the clinical setting.

Our study highlights the value of computer-assisted histopathological analysis to identify additional prognostic characteristics for patients with thin melanoma. We found primary tumor width, digital area, and conformation to be significantly associated with survival outcomes in stage IB patients, independent of Breslow thickness. Using this data, we developed a novel prognostic histopathological classifier that, in combination with baseline variables, effectively stratifies patients with stage IB melanoma by their risk of recurrence. With this prognostic model, 48.9% (71/145 in the studied cohort) stage IB patients are predicted at higher risk of recurrence, among whom 67.6% (48/71) patients are correctly predicted and 32.4% (23/71) patients are falsely classified for the observed follow-up time. The described signature represents a first step towards targeted surveillance—and therefore earlier intervention of recurrence—for high-risk thin primary melanoma patients. We believe such predictive models have the potential to improve survival outcomes for this growing subset of patients with thin melanoma.

Acknowledgments

This work was supported by the NYU Cancer Institute NCI Cancer Center Support Grant (P30 CA016087) and the Marc Jacobs Campaign to support melanoma research.

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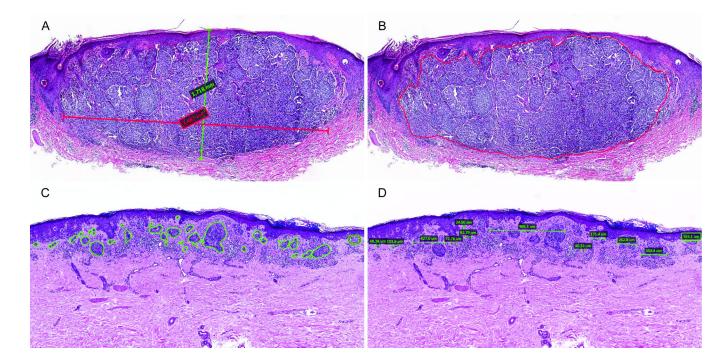
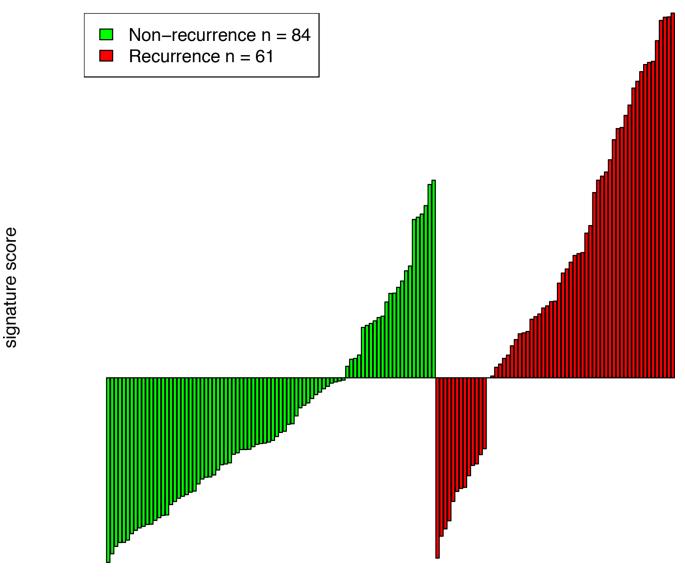


Figure 1.

Primary tumor H&Es demonstrating digital histopathological variables. (A) Manual area was estimated by multiplying the Breslow thickness by the width of the invasive dermal tumor (thickness 1.72 mm, width 3.49 mm, manual area 5.99 mm²). To measure digital area, the invasive dermal tumor was manually outlined and calculated by the Aperio software algorithm, as demonstrated in a (B) contiguous (digital area 4.99 mm²) and a (C) non-contiguous (digital area 0.36 mm²). (D) Tumor width was measured as the cumulative sum of widths of the tumor's invasive dermal components in the greatest horizontal dimension (width 2.75 mm).

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Sensitivity and specificity for prediction of recurrence using a risk score derived from novel histopathological predictive model.

PFS P < 0.001

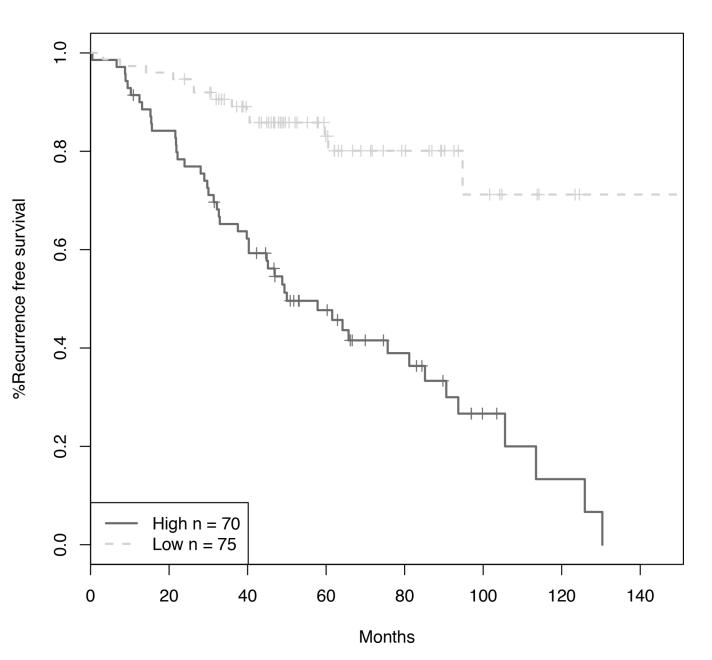


Figure 3.

Progression-free survival (PFS) of patients stratified by the novel histopathological predictive model. Using the risk score previously defined by the novel histopathological predictive model, high and low risk groups were plotted in a Kaplan-Meier survival curve and statistical analysis performed by the log-rank test.

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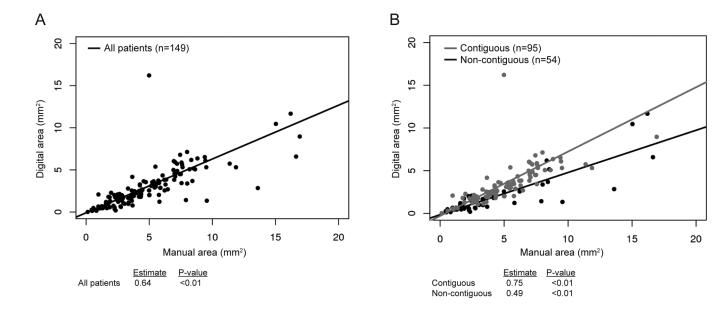


Figure 4.

Regression analysis of manual versus digital tumor area for (**A**) all patients (estimate 0.64, p<0.01) and (**B**) stratified into those with contiguous tumors and those with non-contiguous tumors (estimate 0.75 and 0.49, respectively; p<0.01).

Table 1

Baseline patient and tumor characteristics (n=655).

Characteristic	Total (<i>n=655</i>)
Sex, n (%)	
Male	374 (57.1)
Female	281 (42.9)
Age (years), median (SD)	58.0 (16.6)
Thickness (mm), median (SD)	0.98 (0.44)
Ulceration, n (%)	
Present	28 (4.3)
Absent	621 (95.7)
Mitotic rate (per mm ²), median (SD)	1.0 (2.3)
Mitoses, n (%)	
No	571 (88.1)
Yes	78 (11.9)
Histological subtype, n (%)	
Superficial spreading	379 (67.3)
Nodular	104 (18.5)
Other	80 (14.2)
Recurrence, n (%)	
Not Recurred	578 (88.2)
Recurred	77 (11.8)
Type of recurrence, n (%)	
Local	5 (6.5)
Only local	0 (0.0)
Regional	37 (48.1)
Only regional	11 (14.3)
Distant	52 (67.5)
Only distant	22 (28.6)
Distant + local	4 (5.2)
Distant + regional	25 (32.5)
Distant + regional + local	1 (2.3)
Unknown	14 (18.2)
Death, n (%)	
Alive	578 (88.4)
Dead	76 (11.6)

SD=standard deviation

Local=tumor recurrence within 2 cm of the surgical scar following definitive excision of a primary melanoma with sufficient surgical margins. Regional=tumor recurrence in the regional lymph node basinor that is intralymphatic (satellites or in transit metastases).

Distant= tumor recurrence in distant lymph nodes, skin or subcutaneous tissue >2 cm from the original primary tumor site, or distant organs.

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Table 2

Baseline patient and primary tumor characteristics of the patients that underwent digital histopathological analysis (n=149), only those whose disease recurred (n=63), and only those with non-recurrent disease (n=86).

Total	Recurred	Did not recur	
(n=149)	(n=63)	(n=86)	P-value
94 (63.1)	39 (61.9)	55 (64.0)	0.93
55 (36.9)	24 (38.1)	31 (36.0)	
60 (15.8)	58 (15.8)	62 (15.7)	0.16
1.14 (0.43)	1.10 (0.48)	1.15 (0.39)	0.88
14 (9.6)	6 (9.8)	8 (9.4)	1.00
132 (90.4)	55 (90.2)	77 (90.6)	
5 (5.8)	16 (25.4)	21 (14.1)	< 0.01
81 (94.2)	47 (74.6)	128 (85.9)	
2 (2.7)	3 (2.4)	2 (2.9)	0.73
89 (61.0)	37 (59.7)	52 (61.9)	0.50
47 (32.2)	19 (30.6)	28 (33.3)	
10 (6.8)	6 (9.7)	4 (4.8)	
3.73 (1.88)	4.19 (2.01)	3.41 (1.73)	0.03
3.97 (3.32)	4.38 (3.64)	3.70 (3.03)	0.14
2.10 (2.97)	2.47 (4.00)	1.90 (1.69)	0.01
95 (63.8)	45 (71.4)	50 (58.1)	0.14
54 (36.2)	18 (28.6)	36 (41.9)	
	(n=149) 94 (63.1) 55 (36.9) 60 (15.8) 1.14 (0.43) 14 (9.6) 132 (90.4) 5 (5.8) 81 (94.2) 2 (2.7) 89 (61.0) 47 (32.2) 10 (6.8) 3.73 (1.88) 3.97 (3.32) 2.10 (2.97) 95 (63.8)	(n=149) (n=63) 94 (63.1) 39 (61.9) 55 (36.9) 24 (38.1) 60 (15.8) 58 (15.8) 1.14 (0.43) 1.10 (0.48) 14 (9.6) 6 (9.8) 132 (90.4) 55 (90.2) 5 (5.8) 16 (25.4) 81 (94.2) 47 (74.6) 2 (2.7) 3 (2.4) 89 (61.0) 37 (59.7) 47 (32.2) 19 (30.6) 10 (6.8) 6 (9.7) 3.73 (1.88) 4.19 (2.01) 3.97 (3.32) 4.38 (3.64) 2.10 (2.97) 2.47 (4.00) 95 (63.8) 45 (71.4)	(n=149) $(n=63)$ $(n=86)$ 94 (63.1)39 (61.9)55 (64.0)55 (36.9)24 (38.1)31 (36.0)60 (15.8)58 (15.8)62 (15.7)1.14 (0.43)1.10 (0.48)1.15 (0.39)14 (9.6)6 (9.8)8 (9.4)132 (90.4)55 (90.2)77 (90.6)5 (5.8)16 (25.4)21 (14.1)81 (94.2)47 (74.6)128 (85.9)2 (2.7)3 (2.4)2 (2.9)89 (61.0)37 (59.7)52 (61.9)47 (32.2)19 (30.6)28 (33.3)10 (6.8)6 (9.7)4 (4.8)3.73 (1.88)4.19 (2.01)3.41 (1.73)3.97 (3.32)4.38 (3.64)3.70 (3.03)2.10 (2.97)2.47 (4.00)1.90 (1.69)95 (63.8)45 (71.4)50 (58.1)

SD=standard deviation

Table 3

Univariate and multivariate analyses of primary tumor width, manual area, digital area, and conformation with recurrence-free and overall survival (n=149).

	Univariate analysis		Multivariate analysis [*]	
	HR (95% CI)	P-value	HR (95% CI)	P-value
Recurrence-free survival				
Width (mm)	1.17 (1.04–1.32)	0.008	1.20 (1.06–1.37)	0.004
Manual area (mm ²)	1.06 (1.00–1.13)	0.069	1.08 (0.99–1.18)	0.069
Digital area (mm ²)	1.08 (1.03–1.14)	0.003	1.10 (1.04–1.17)	0.002
Conformation ¹	0.57 (0.32-0.99)	0.046	0.55 (0.30-1.00)	0.048
Overall survival				
Width (mm)	1.05 (0.91–1.21)	0.492	1.01 (0.87–1.17)	0.924
Manual area (mm ²)	1.02 (0.94–1.10)	0.690	0.98 (0.89–1.08)	0.641
Digital area (mm ²)	1.04 (0.97–1.11)	0.305	1.02 (0.93–1.13)	0.643
Conformation ¹	1.11 (0.60–2.06)	0.745	1.10 (0.57–2.10)	0.782

HR=hazard ratio; CI=confidence interval

¹ Reference value = contiguous

* Cox model adjusted for gender, age at diagnosis, thickness (mm), mitoses (yes/no) in addition to the variable of interest.