

A computed tomography-based score indicative of lung cancer aggression (SILA) predicts lung adenocarcinomas with low malignant potential or vascular invasion

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Abstract.

BACKGROUND: Histologic grading of lung adenocarcinoma (LUAD) is predictive of outcome but is only possible after surgical resection. A radiomic biomarker predictive of grade has the potential to improve preoperative management of early-stage LUAD.

OBJECTIVE: Validate a prognostic radiomic score indicative of lung cancer aggression (SILA) in surgically resected stage I LUAD ($n = 161$) histologically graded as indolent low malignant potential (LMP), intermediate, or aggressive vascular invasive (VI) subtypes.

METHODS: The SILA scores were generated from preoperative CT-scans using the previously validated Computer-Aided Nodule Assessment and Risk Yield (CANARY) software.

RESULTS: Cox proportional regression showed significant association between the SILA and 7-year recurrence-free survival (RFS) in a univariate ($p < 0.05$) and multivariate ($p < 0.05$) model incorporating age, gender, smoking status, pack years, and extent of resection. The SILA was positively correlated with invasive size (spearman $r = 0.54$, $p = 8.0 \times 10^{-14}$) and negatively correlated with percentage of lepidic histology (spearman $r = -0.46$, $p = 7.1 \times 10^{-10}$). The SILA predicted indolent LMP with an area under the receiver operating characteristic (ROC) curve (AUC) of 0.74 and aggressive VI with an AUC of 0.71, the latter remaining significant when invasive size was included as a covariate in a logistic regression model ($p < 0.01$).

CONCLUSIONS: The SILA scoring of preoperative CT scans was prognostic and predictive of resected pathologic grade.

Keywords: Lung adenocarcinoma, vascular invasion, radiomic biomarkers, SILA, indolent lung cancer

1. Introduction

Lung cancer (LC) is the deadliest cancer in the United States (U.S.) with an estimated 238,340 new cases and 127,070 deaths in 2023 [1]. However, lung cancer mortality has begun to decline in part due to declining rates of cigarette smoking and more recently

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the widespread implementation of low-dose computed tomography (CT) screening programs that have led to detection at earlier stages where curative surgery is possible [1]. Despite the mortality reduction associated with LC screening, CT screening results in an increase in overdiagnosis leading to higher morbidity, financial burden, and stress among patients [2,3]. Accurate pre-surgical prognostic markers are needed to personalize the management of early stage LC. Indolent tumors may be able to be treated with non-surgical approaches such as stereotactic body radiation therapy (SBRT), cryoablation, microwave ablation, or radiofrequency ablation (RFA) [4]. New clinical trials also indicate that a subset of early stage LC can be adequately managed with sublobar resection rather than standard of care lobectomy [5,6]. On the other hand, patients with aggressive disease and high risk of recurrence may benefit from adjuvant or neoadjuvant systemic therapy, which is not standard of care for stage I disease [7,8]. Tumor histopathology is highly prognostic, but it requires comprehensive histologic examination that is only possible after complete surgical excision [9]. Small biopsies, such as those obtained via bronchoscopy or CT-guided biopsy, are able to establish a diagnosis of LC and distinguish between LC subtypes, but cannot reliably provide the same level of prognostic information as resected specimens due to limited sampling, tumor heterogeneity, and crush artifact [10]. Widespread validation and clinical implementation of machine learning approaches that can predict prognostic histologic patterns and features from CT scans are an important approach to improve clinical management of early-stage tumors.

Lung adenocarcinoma (LUAD) is the most common subtype of LC overall, accounts for virtually all cases among light and never smokers, and is heterogeneous in its histologic patterns, features and prognosis [9]. In the National Lung Screening Trial (NLST), overdiagnosis was high (79%) among a subset of LUAD historically termed “bronchoalveolar carcinoma” (BAC), which comprised 27% of all LUAD detected by CT-screening [3,11,12]. Since the NLST, BAC has been discontinued as a diagnostic entity and replaced with adenocarcinoma in situ (AIS) and minimally invasive adenocarcinoma (MIA) which exhibit 100% disease-free survival (DFS) after excision, but together make up only ~5% of stage I LUAD, substantially less than BAC in the NLST [13].

Recently, a proposed histopathology classification of stage I LUADs as low malignant potential (LMP) with 100% DFS that includes AIS and MIA, accounted for

23% of stage I LUAD, reflecting a similar proportion of cases as was reported as overdiagnosed stage I BAC in NLST [14]. In contrast to LMP, there are tumor invasive characteristics that are associated with poor prognosis. Vascular invasion (VI), a pathological hallmark of cancer pre-metastasis and a strong predictor of recurrence, cancer specific and overall mortality in patients with early-stage LUAD, even among tumors < 2 cm invasive size, has been shown to be more prognostic than the highest World Health Organization (WHO) grade [15, 16,17,18,19].

We sought to evaluate the ability of a previously published CT scan-based method to distinguish between stage I LUAD classified as indolent (AIS/MIA/LMP), aggressive (VI), and intermediate grade (NST-no special type) at the time of resection. Computer Aided Nodule Assessment and Risk Yield (CANARY) is a software for automated risk assessment of adenocarcinoma based on of the clustering of voxel density histograms into nine clusters or exemplars named after colors [20]. Multidimensional scaling showed these nine exemplars clustered into three groups that visually corresponded to ground-glass appearance, solid appearance, and intermediate density. CANARY was originally designed and validated to distinguish invasive adenocarcinomas from AIS/MIA [20,21]. Subsequently, three CANARY risk groups were defined and association with patient outcomes were validated, independent of histology, in two retrospective surgical lung adenocarcinoma cohorts, including the NLST [22,23]. The good risk group among pathologic stage I adenocarcinoma was associated with 100% disease specific survival (DSS) in both cohorts. Interestingly, the good risk group represented 17% and 18% of pathologic stage I tumors in these cohorts, far exceeding the expected rate of AIS/MIA (~5% combined). The latter finding implies that CANARY can predict a proportion of invasive lung adenocarcinomas beyond AIS/MIA that behave in an indolent fashion. Subsequent studies transformed the output of CANARY into a score indicative of lung cancer aggression (SILA) based on the prediction of invasive size and outcome [24,25]. Here, we further validate the association of CANARY and the corresponding SILA with prognosis in a retrospective cohort of pathologic stage I LUAD treated by surgical excision in an urban safety-net hospital setting. We also show that CANARY/SILA is predictive of WHO-2021 grade and our novel histopathologic grade, indicating that it detects histopathologic characteristics of LUAD invasion beyond invasive size.

2. Materials and methods

2.1. Clinical samples and pathology review

A retrospective cohort of 161 patients who were treated with surgery between 2005–2014 for pathologic stage I/0 LUAD were included in this study, representing a subset of a previously reported cohort [14,18]. Tumors measuring > 4 cm total size were not included, as subsets of these patients were given adjuvant therapy within this historic cohort. Cases were reviewed from Boston Medical Center (BMC), an urban safety-net hospital, after IRB approval (BU/BMC IRB H-37859 12/11/2018) in which patient consent was waived as this retrospective study posed no more than minimal risk of harm to subjects and involved no procedures for which written consent is normally required. The study was performed in accordance with the Declaration of Helsinki. Preoperative CT scans were obtained for all patients between December 2004 and November 2015. The median time from preoperative CT scan acquisition to surgery was 30 days. All matching pathology cases were reviewed by an experienced board-certified thoracic pathologist (EJB). Vascular invasion (VI) was defined as luminal invasion of a muscular artery or vein either within or adjacent to the tumor. Tumors were assessed for proportion of lepidic, acinar, papillary, micropapillary, and solid patterns in 5% increments with distinction of simple tubular acinar from complex and cribriform acinar patterns. Adenocarcinoma in situ (AIS) was rendered for purely lepidic tumors ≤ 3 cm whereas minimally invasive adenocarcinoma (MIA) was diagnosed when non-lepidic foci measured ≤ 0.5 cm as per WHO criteria [26]. WHO-2021 grade was defined as G1, lepidic predominant with < 20% high-grade patterns; G2, acinar or papillary predominant with < 20% high-grade patterns; and G3, $\geq 20\%$ high-grade patterns (solid, micropapillary and/or complex glands) [26,27]. Low malignant potential adenocarcinoma (LMP) was assigned as previously described [14]. LMP tumors were non-mucinous adenocarcinoma measuring ≤ 3 cm in total size, exhibiting $\geq 15\%$ lepidic growth, and lacking nonpredominant high-grade patterns ($\geq 10\%$ cribriform, $\geq 5\%$ micropapillary, $\geq 5\%$ solid), > 1 mitosis per 2 mm², vascular, lymphatic or visceral pleural invasion, STAS or necrosis. Given the identical behavior (100% 10-year DSS) to AIS/MIA, these were analyzed together, except where reported separately. All other tumors not classified as VI or LMP are referred to as no special type (NST). Pathologic stage assignments were retrospec-

tively made based upon the 8th edition of the AJCC. In older editions, tumors up to 5 cm were classified as stage IB and may have been recommended for adjuvant therapy. In the 8th edition, only tumors up to 4 cm are classified as stage I.

2.2. CANARY analysis

CANARY Plus software version 1.0 was licensed from Mayo Clinic. CANARY has previously been demonstrated to have low inter-observer variability for segmenting and analyzing LUAD CT scans [28]. All CT scans were reviewed by an experienced board-certified thoracic radiologist at the time of clinical diagnosis. CT scans were acquired using a variety of scanners, with the majority (96.3%) acquired on one scanner. As part of this retrospective study, we collaborated with an experienced board-certified thoracic surgeon (KS) who confirmed that the nodule location on the CT scan matched the resected nodule on the original clinical report, and that adequate masking was performed by the CANARY nodule detection algorithm. The SILA and associated exemplars were generated by CANARY and exported for further analysis. The nine exemplars were previously named based on nine arbitrary colors: blue (B), cyan (C), green (G), yellow (Y), pink (P), violet (V), indigo (I), red (R), and orange (O) [20].

2.3. Statistical analysis

All statistical analysis was performed with R version 4.2.1. Tables were created with the tableone package. Comparisons of distributions of count data were tested with `chisq.test`. Correlations were performed using spearman correlation with `stat_cor` or `cor.test`. Comparisons of distributions of continuous data were tested with `wilcox.test` or `t.test`, as specified. *P*-values were converted to false-discovery rate (FDR) values by `p.adjust` using the bonferonni method. Survival analysis used recurrence-free survival (RFS) as an endpoint, which was defined as the time from surgery to recurrence or last follow up. Univariate and multivariate Cox regression was performed using the survival package version 3.5.3. Kaplan-Meier plots were created using the `survminer` package version 0.4.9 and groups compared using the log-rank test. Area under the curve (AUC) calculations and receiver operating characteristic (ROC) plots were created using the `pROC` package version 2.3.0 [29]. All statistical tests were two-tailed and *p* values < 0.05 were considered significant.

Table 1
Clinical and pathologic characteristics of 161 patients with resected stage I LUAD included in the study

	Overall
<i>n</i>	161
Age (mean (SD))	67.30 (9.59)
Gender	
Female	97 (60.2)
Male	64 (39.8)
Race	
Asian	13 (8.1)
Black/African American	50 (31.1)
Hispanic/Latino	5 (3.1)
Unknown	8 (5.0)
White	85 (52.8)
Pack years (mean (SD))	38.76 (33.67)
Smoking Status	
Never	22 (14.2)
Former	73 (47.1)
Current	60 (38.7)
Procedure	
Lobe	103 (64.0)
Segment	8 (5.0)
Wedge	50 (31.1)
Invasive size (mean (SD))	1.39 (0.79)
Total size (mean (SD))	1.86 (0.84)
WHO 2021 grade	
AIS/MIA	4 (2.5)
G1	28 (17.4)
G2	43 (26.7)
G3	80 (49.7)
M	6 (3.7)
Novel grade	
LMP	27 (16.8)
NST	87 (54.0)
VI	47 (29.2)
Recurrence	20 (12.4)
Follow-up years (mean (SD))	5.95 (3.42)

Note: The data are shown as the number and (%) unless otherwise indicated. Abbreviations: AIS, adenocarcinoma in situ; MIA, minimally invasive adenocarcinoma; G1, grade 1; G2, grade 2; G3, grade 3; M, mucinous, LMP, low malignant potential; NST, no special type; VI, vascular invasion.

3. Results

3.1. Patient and tumor characteristics

Table 1 shows the clinical and pathologic characteristics of 161 patients with resected stage I LUAD included in the study. The mean age was 67.3 years. Most patients were female (60%), self-identified as white (53%), were former (47%) smokers, and were treated with lobectomy (64%). The patients in the study had an overall 7-year RFS of 88% with a mean follow-up time of 5.95 years. Kaplan-Meier estimation showed a significant difference in both RFS and DSS among grades from both the WHO 2021 grading ($p < 0.05$) and the novel grading classifications ($p < 0.001$) (Fig. S1A-B).

AIS/MIA, WHO G1, WHO G2, and WHO G3 had 7-year RFS of 100%, 96%, 95%, and 81%, respectively. LMP, NST, and VI grades had 7-year RFS of 96%, 95%, and 65%, respectively. A single LMP recurred after wedge-resection with a positive surgical margin. The tumor recurred at the staple line and was treated with SBRT with prolonged survival (> 10 years) without recurrence or metastasis. VI grade was associated with patients that identified as male ($p < 0.01$), Black or African American ($p < 0.05$), and were current smokers ($p < 0.05$), as previously reported (Table S1) [18]. No patients received adjuvant therapy.

3.2. The SILA is associated with recurrence-free survival

The SILA scores were binned into good ($n = 12$), intermediate ($n = 94$), and poor ($n = 55$) subgroups using the cutoffs established in the original manuscript (Fig. 1A) [24]. The mean SILA in each subgroup was 0.26, 0.54, and 0.75, respectively. Detailed results are shown in Table 2. Kaplan-Meier estimation revealed a significant difference in outcome among the three subgroups, with the good, intermediate, and poor subgroups having 7-year RFS of 100%, 91%, and 73%, respectively ($p < 0.05$) (Fig. 1B). The SILA was significantly predictive of RFS in univariate analysis (hazard ratio (HR) = 2.07, $p < 0.05$) (Fig. 1C). In a multivariate analysis including pack years, smoking status, gender, age, and surgical procedure, the SILA remained significant for RFS (HR = 1.84, $p < 0.05$) (Fig. 1D).

3.3. The SILA is associated with pathologic grade at resection

Given that the SILA has previously been reported as linearly increasing with invasive size (non-lepidic tumor size) at resection [24], we sought to validate this in our cohort and examine associations with other pathology features observable in the resected tumor. The SILA was positively correlated with invasive size at resection ($R = 0.54$, $p = 8.0 \times 10^{-14}$) (Fig. 2A) and negatively correlated with the percentage of lepidic growth pattern ($R = -0.46$, $p = 7.1 \times 10^{-10}$) (Fig. 2B). The SILA increased with grade in both the novel grading system and WHO 2021 grades but was not significantly different between tumors classified as AIS/MIA and WHO grade 1 ($p = 0.19$), AIS/MIA and LMP ($p = 0.27$), or tumors classified as WHO grade 2 and WHO grade 3 ($p = 0.93$) (Fig. 2C-D). Given the inverse correlation of the SILA with percentage of lepidic growth pattern, we

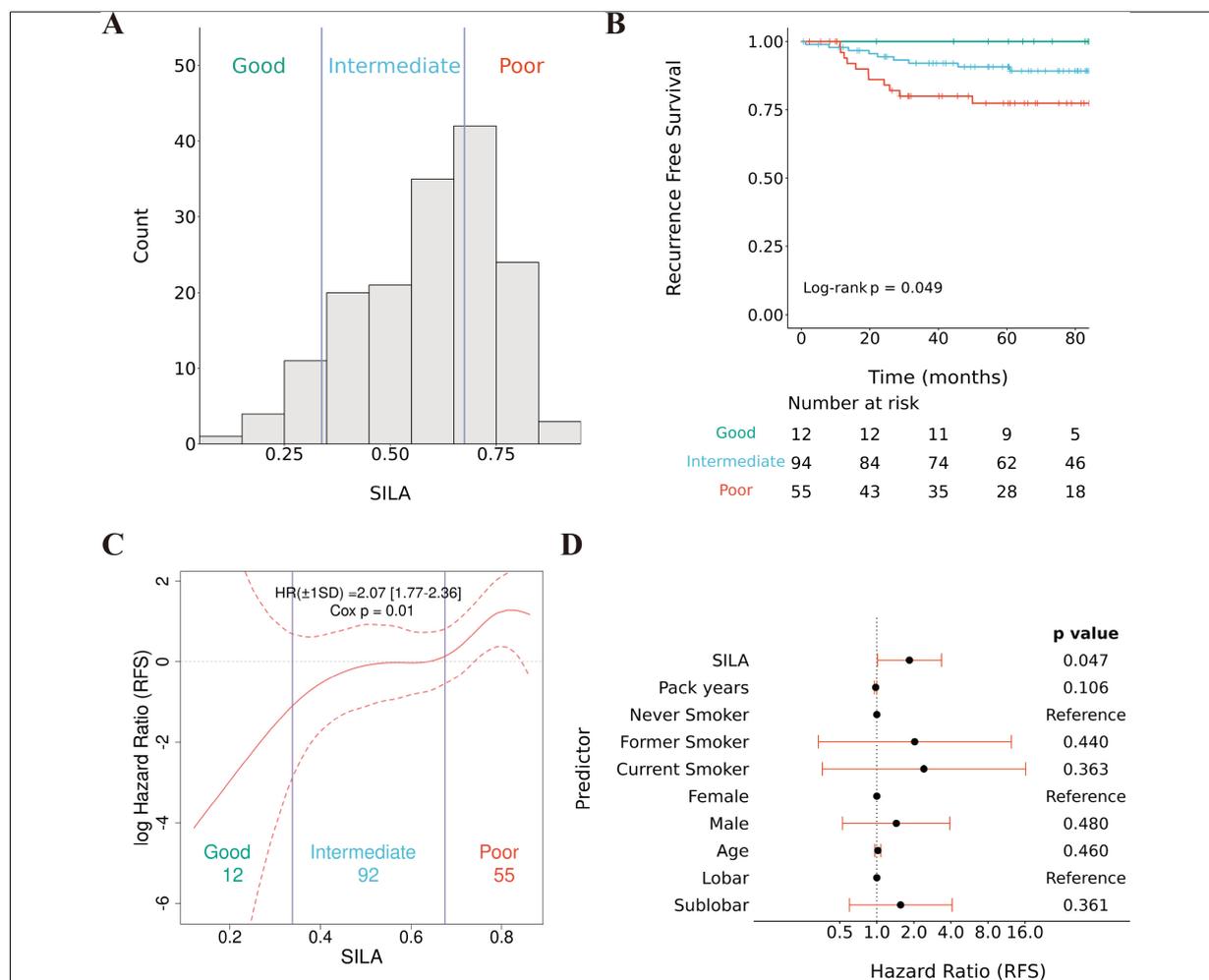


Fig. 1. The SILA is associated with recurrence-free survival in a cohort of resected stage I LUAD. (A) Distribution of the SILA by prognostic subgroup using previously established cutoffs (Varghese et al., 2019). (B) Kaplan Meier curve of the SILA prognostic subgroups with 7-year RFS. (C) Univariate cox proportional hazard model of the SILA predicting 7-year RFS. (D) Multivariate cox proportional hazard model of the SILA predicting 7-year RFS, with pack years, smoking status, gender, age, and surgical procedure as covariates.

263 evaluated the relationship between percentage of lepidic
 264 growth pattern and grade which significantly decreased
 265 between each grade group (Fig. S2). The SILA separated
 266 the combined category of AIS/MIA/LMP tumors
 267 from non-AIS/MIA/LMP tumors with an AUC of 0.74
 268 and tumors with VI with an AUC of 0.71 (Fig. 3A–B).
 269 The SILA was also significantly associated with VI in
 270 a logistic regression model even after controlling for
 271 invasive size ($p < 0.01$) or percentage of lepidic pattern
 272 ($p < 0.05$). A model for predicting VI containing
 273 the SILA and invasive size had significantly less error
 274 than a model containing invasive size alone (LRT $p =$
 275 0.004) but not less than a model containing the SILA
 276 alone (LRT $p = 0.52$), indicating that the SILA mediates
 277 the association of VI and invasive size. Examples

278 of CANARY masks of LMP and VI cases are provided
 279 (Fig. S3). The SILA distinguished indolent cancer as
 280 classified by WHO 2021 AIS/MIA grade with an AUC
 281 of 0.84 but showed lower performance in the prediction
 282 of WHO grade 3 (AUC 0.60) from other grades
 283 (Fig. S4A–B).

3.4. The CANARY red exemplar is associated with VI at resection

284 Given that the SILA was weakly associated with
 285 WHO grade 3 tumors containing aggressive histologic
 286 patterns, we sought to determine whether any of the
 287 nine CANARY exemplars were associated with percent-
 288 ages of different growth patterns. Correlation analysis
 289
 290

Table 2

Clinical and pathologic characteristics of resected stage I LUAD classified by the SILA prognostic subgroups

	Good	Intermediate	Poor	<i>p</i> value
<i>n</i>	12	94	55	
SILA (mean (SD))	0.26 (0.06)	0.54 (0.10)	0.75 (0.05)	< 0.001
Age (mean (SD))	65.58 (11.04)	67.48 (9.43)	67.36 (9.68)	0.81
Gender (%)				0.03
Female	11 (91.7)	58 (61.7)	28 (50.9)	
Male	1 (8.3)	36 (38.3)	27 (49.1)	
Race (%)				0.646
Asian	1 (8.3)	10 (10.6)	2 (3.6)	
Black/African American	2 (16.7)	27 (28.7)	21 (38.2)	
Hispanic/Latino	1 (8.3)	2 (2.1)	2 (3.6)	
Unknown	1 (8.3)	4 (4.3)	3 (5.5)	
White	7 (58.3)	51 (54.3)	27 (49.1)	
Pack years (mean (SD))	24.58 (18.37)	38.77 (36.78)	42.01 (30.22)	0.272
Smoking Status (%)				0.009
Never	3 (25.0)	17 (18.9)	2 (3.8)	
Former	2 (16.7)	46 (51.1)	25 (47.2)	
Current	7 (58.3)	27 (30.0)	26 (49.1)	
Procedure (%)				0.815
Lobe	6 (50.0)	60 (63.8)	37 (67.3)	
Segment	1 (8.3)	4 (4.3)	3 (5.5)	
Wedge	5 (41.7)	30 (31.9)	15 (27.3)	
Invasive size (mean (SD))	0.63 (0.39)	1.21 (0.65)	1.87 (0.82)	< 0.001
Total size (mean (SD))	1.33 (0.60)	1.76 (0.82)	2.14 (0.85)	0.002
WHO 2021 grade (%)				< 0.001
AIS/MIA	2 (16.7)	2 (2.1)	0 (0.0)	
G1	3 (25.0)	22 (23.4)	3 (5.5)	
G2	5 (41.7)	19 (20.2)	19 (34.5)	
G3	1 (8.3)	46 (48.9)	33 (60.0)	
M	1 (8.3)	5 (5.3)	0 (0.0)	
Novel grade (%)				< 0.001
LMP	7 (58.3)	17 (18.1)	3 (5.5)	
NST	3 (25.0)	61 (64.9)	23 (41.8)	
VI	2 (16.7)	16 (17.0)	29 (52.7)	

Note: The data are shown as the number and (%) unless otherwise indicated. Abbreviations: SILA, score indicative of lung cancer aggression; AIS, adenocarcinoma in situ; MIA, minimally invasive adenocarcinoma; G1, grade 1; G2, grade 2; G3, grade 3; M, metachronous, LMP, low malignant potential; NST, no special type; VI, vascular invasion.

291 followed by unsupervised clustering revealed that non-
 292 lepidic patterns clustered separately from the exemplars
 293 (Fig. 4A), suggesting they are not major drivers of the
 294 SILA. The red exemplar had the highest performance
 295 for predicting VI (AUC of 0.69) (Fig. 4B). When all
 296 exemplars were included in a logistic regression model
 297 for predicting VI, only the red exemplar was significant
 298 ($p < 0.05$). Furthermore, after performing stepdown
 299 Akaike information criterion (AIC) analysis, the lowest
 300 AIC was obtained for a model that included only the red
 301 exemplar, suggesting that the red exemplar is primarily
 302 responsible for SILA's ability to predict VI. Finally,
 303 LMP was classified equivalently by multiple CANARY
 304 exemplars (Fig. 4C). The lowest AIC was obtained for
 305 a model that included the indigo ($p < 0.01$), blue ($p <$
 306 0.01), and green ($p < 0.10$) exemplars, suggesting that

there are multiple radiologic aspects of the nodule that
 contribute to the prediction of LMP.

4. Discussion

This study evaluated the association between CA-
 NARY, a well-described algorithm for preoperative pre-
 diction of indolent and aggressive LUAD [20,22,24,25,
 28], and histologic grade in an urban safety-net hospital
 for the first time. In this cohort, the low, medium, and
 high CANARY SILA prognostic groups were associ-
 ated with 100%, 91%, and 73% 7-year RFS respec-
 tively, and the SILA was significantly associated with
 RFS even after correction for other clinical factors. The
 SILA prognostic subgroups were originally identified
 by association with linear extent of histologic invasion

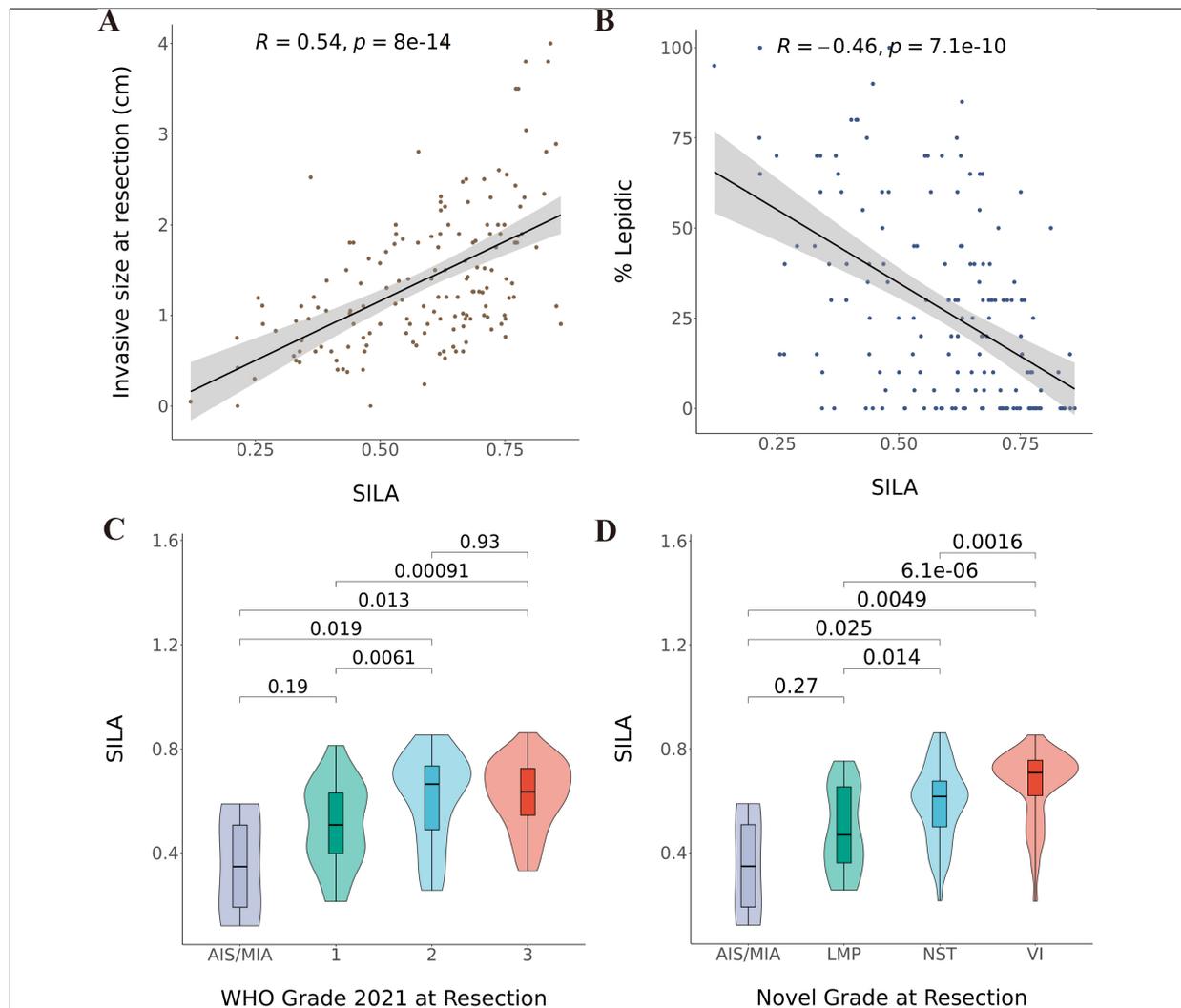


Fig. 2. The SILA is associated with pathologic grade at resection. (A) The SILA correlation with invasive size at resection. (B) The SILA correlation with percentage of lepidic growth pattern, measured at resection. (C) The SILA association with WHO 2021 grading criteria. (D) The SILA association with novel pathology grading criteria.

321 and showed prognostic stratification in both an internal
 322 and external cohort of predominantly (83%) clinical
 323 stage I LUAD; exhibiting 100%, 79%, and 58% 5-year
 324 DSS [24]. Our improved outcomes among intermediate
 325 and poor SILA risk groups likely reflect the restriction
 326 of our analysis to pathologic stage I LUAD. As the
 327 SILA has been previously validated in a cohort derived
 328 from a subset of the NLST containing 94% white pa-
 329 tients, the validation of CANARY and the SILA for
 330 predicting prognosis in a cohort containing patients of
 331 diverse racial and ethnic identity (47% non-white) is en-
 332 couraging given that both LUAD incidence and LUAD
 333 aggressiveness at diagnosis is higher for non-Hispanic
 334 black patients [18,30,31]. Additionally, our cohort cap-

335 tures the diverse etiology that is known about LUAD,
 336 with 14% of patients being never-smokers.

337 There remains no clinically accepted approach to
 338 preoperatively predict tumor aggressiveness among sur-
 339 gically operable LCs, which are managed uniformly
 340 by clinical stage, potentially resulting in over-treatment
 341 of indolent lesions. While the SILA has been shown
 342 to accurately predict AIS and MIA stage I LUAD, we
 343 have previously shown that tumors designated as LMP
 344 more closely match the proportion of overdiagnosed
 345 cases in the NLST [14] and Burks et al. in this edition.
 346 In our cohort, the SILA “Good” group ($n = 12$) had
 347 100% RFS, and the SILA achieved an AUC of 0.74 for
 348 classifying the larger group of LMP tumors ($n = 27$),

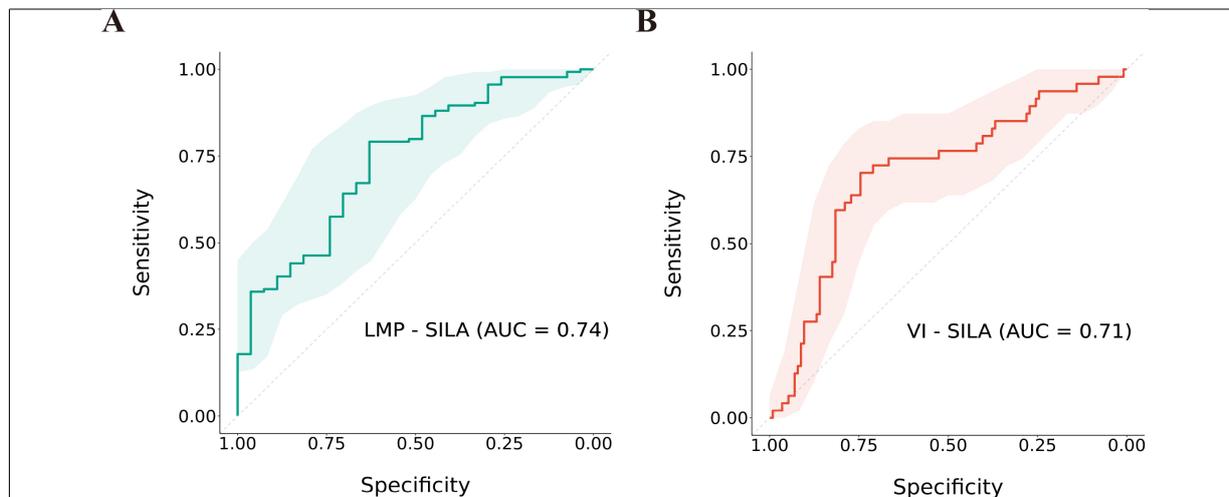


Fig. 3. The SILA performance for predicting LMP and VI stage I LUAD tumors. (A) ROC curve of the SILA predicting cases of LMP (including AIS/MIA) (Wilcoxon $p = 8.0e^{-05}$). (B) ROC curve of the SILA predicting cases of VI (Wilcoxon $p = 4.2e^{-05}$).

349 which had 100% DFS. Future studies may therefore
 350 seek to improve the SILA's identification of indolent
 351 stage I LUAD by incorporating other features such as
 352 serum proteomics, biopsy pathology, liquid biopsy and
 353 mutational or transcriptomic profiling into multimodal
 354 predictive models [32,33]. Although data from large
 355 clinical trials including JCOG0802/WJOG4607L and
 356 CALGB140503 suggest that lobectomy does not offer
 357 a survival benefit over limited resection, additional
 358 data is needed to determine whether patients identified
 359 with indolent disease may in the future have similar
 360 outcomes when treated with non-surgical approaches
 361 such as SBRT and RFA [4,5,6].

362 Tumor grading by microvascular invasion, the his-
 363 tologic representation of tumor intravasation, has been
 364 shown to be more strongly associated with post-surgical
 365 outcome in stage I LUAD than grading that takes into
 366 consideration the proportion of the aggressive LUAD
 367 histologic patterns – solid and micropapillary [18]. Ret-
 368 rospective analysis shows that patients with VI who
 369 undergo sublobar resection have poorer outcomes [34].
 370 This underscores the growing importance of identify-
 371 ing individuals with more aggressive disease prior to
 372 surgery. Using the SILA to predict VI preoperatively,
 373 potentially in combination with other biomarker modal-
 374 ities, may therefore offer opportunities to guide preci-
 375 sion surgery, but prospective studies are needed. Tum-
 376 ors exhibiting VI may also identify candidates who
 377 would benefit from adjuvant or neoadjuvant therapy. In
 378 this study, the SILA predicted VI with an AUC of 0.71
 379 and was associated with VI independently of invasive
 380 size at resection. A previous study of stage IA LUAD

381 nodules found that the ratio of the length of nodule con-
 382 solidation to nodule diameter in preoperative CT scans
 383 predicted combined lymphatic and/or blood vessel inva-
 384 sion [35]. While lymph vessel invasion is often reported
 385 interchangeably with angioinvasion, it may not be as
 386 strong of an independent prognostic factor [18,19]. Ex-
 387 amination of the CANARY exemplars showed that the
 388 red exemplar, which corresponds to the most visually
 389 solid tumor areas on CT, [20] was most responsible for
 390 the performance of the SILA for predicting VI. This is
 391 the first analysis showing CANARY to be predictive
 392 of a specific type of pathologic invasion; prior studies
 393 assessed CANARY and/or the SILA for predicting the
 394 size of any type of invasion. Others have identified the
 395 violet, indigo, red, and orange CANARY exemplars,
 396 visually corresponding to varying degrees of solid tu-
 397 mor CT appearance, as being associated with a lower
 398 likelihood of EGFR-mutated LUAD, which might be
 399 expected since these tumors frequently are rich in lep-
 400 idic histology and ground glass CT-appearance [36].
 401 Additionally, other studies have shown a lower preva-
 402 lence of EGFR-mutated cases among both LUAD and
 403 NSCLC with VI [37,38].

404 Future efforts to improve the preoperative prediction
 405 of VI from CT images may take advantage of convolu-
 406 tional neural network-based extraction of perinodular
 407 features to add additional context from the surrounding
 408 lung microvascular architecture [39]. A study seeking
 409 to preoperatively predict VI positive hepatocellular car-
 410 cinoma from CT scans achieved an AUC of 0.89 in a
 411 validation set using a radiomic model incorporating per-
 412 itumoral features [40]. In contrast to our findings with

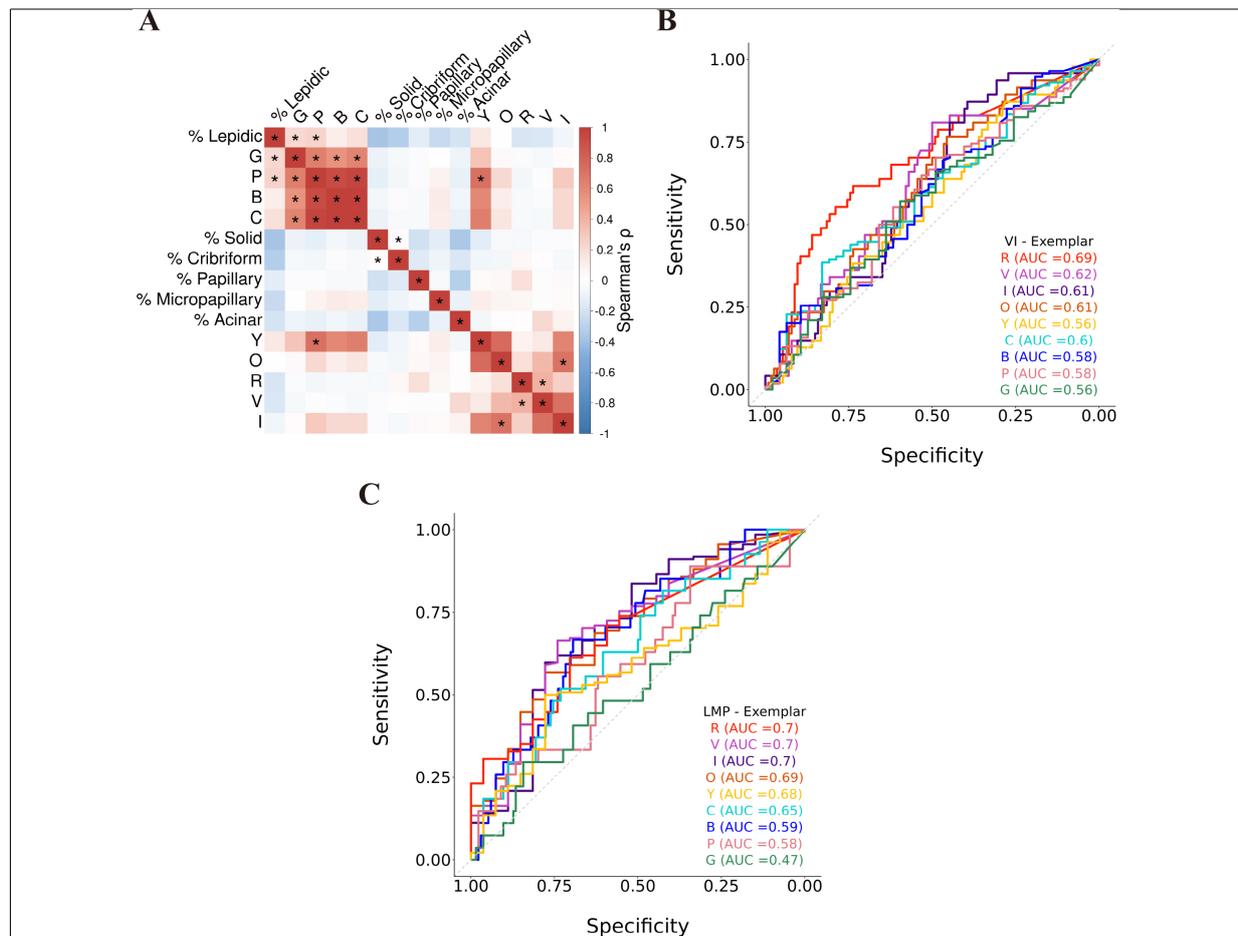


Fig. 4. The CANARY red exemplar is associated with VI at resection. (A) Correlation matrix of CANARY exemplars with percentages of different LUAD histologic growth patterns (* indicates FDR < 0.05). (B) ROC curves of individual CANARY exemplars predicting VI cases. (C) ROC curves of individual CANARY exemplars predicting LMP cases.

413 VI, the SILA and the CANARY exemplars were not
 414 associated with aggressive LUAD histologic patterns,
 415 which may explain the poor predictive performance we
 416 reported for WHO grade 3 tumors. Other studies have
 417 demonstrated the feasibility of building radiomic clas-
 418 sifiers that may predict solid and micropapillary histol-
 419 ogy, suggesting that other nodule features than those
 420 extracted by the SILA may be more representative of
 421 these high-grade patterns [41,42].

422 There were several limitations present in this study.
 423 Despite the robust validation of the SILA in a new
 424 cohort, the cohort was collected over 11 years and there
 425 may be variability due to changes in standard of care
 426 and practice patterns. Additionally, the low number of
 427 AIS/MIA cases included ($n = 4$) did not allow for a
 428 robust validation of the SILA to distinguish between
 429 indolent and invasive LUAD as defined by the WHO
 430 2021 grading scheme [25]. Finally, because most of the

431 CT scans used in our study were acquired with the same
 432 scanner manufacturer, we cannot rule out variability in
 433 the SILA due to scanner type. However, CANARY and
 434 the SILA were both derived on and have since been
 435 validated across a variety of scanner types.

436 5. Conclusion

437 The SILA derived from preoperative CT scans was
 438 prognostic and predictive of resected pathologic grade
 439 in stage I LUAD patients from a diverse cohort of pa-
 440 tients. New strategies are necessary to minimize over-
 441 diagnosis in this clinical setting and identify aggres-
 442 sive tumors that may benefit from precision surgery,
 443 adjuvant and/or neoadjuvant treatment. Ultimately, the
 444 SILA should be prospectively validated and bench-

marked against pathology review of biopsies to identify both LMP and VI tumors preoperatively.

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Author contributions

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Supplementary data

The supplementary files are available to download from <http://dx.doi.org/10.3233/CBM-230456>.

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