

VALIDATION OF A CELL CYCLE PROGRESSION SCORE FOR FIVE-YEAR MORTALITY RISK IN PATIENTS WITH STAGE I LUNG ADENOCARCINOMA

Takashi Eguchi, MD, PhD¹; Kyuichi Kadota, MD, PhD¹; Brent Evans, MS²; Kay See Tan, PhD¹; Camelia S. Sima, MD, MS¹; Thaylon Davis, BS²; Stephanie A. Hamilton, MBA²; Kraig Yager, PhD²; Joshua T. Jones, PhD²; Anne-Renee Hartman, MD³; Prasad S. Adusumilli, MD¹

1. Memorial Sloan Kettering Cancer Center, New York, NY 2. Myriad Genetic Laboratories, Inc., Salt Lake City, UT
3. Myriad Genetics, Inc., Salt Lake City, UT

PRESENTED AT THE 2015 AMERICAN SOCIETY OF CLINICAL ONCOLOGY ANNUAL MEETING:
MAY 29 – JUNE 2, 2015, CHICAGO, ILLINOIS

BACKGROUND

- Although surgery is the standard treatment for patients with early stage (I-IIIa) lung cancer [1], survival rates remain relatively low.
- There is evidence that some patients may benefit from post-surgical chemotherapy; however, identification of patients who would benefit from adjuvant chemotherapy is under investigation [2,3].
- A 46-gene panel assessing cell cycle progression (CCP) gene expression has been developed for pathologic stage I and II lung adenocarcinoma patients [4,5].

OBJECTIVE

- The aim of this study was to validate the CCP gene signature (CCP score) and a Prognostic Score [(PS) combination of CCP and pathological stage] (IA or IB) that identify stage I non-small cell lung cancer patients with a higher risk of cancer-related death after surgical resection.

METHODS

PATIENTS

- A set of 1200 consecutively treated patients at the Memorial Sloan Kettering Cancer Center was identified.
- Patients were included in the study if they had NSCLC with adenocarcinoma histology, Stage I disease using 7th edition International Association for the Study of Lung Cancer criteria and a complete resection of the primary lung tumor. Patients who recurred and then received chemotherapy and/or radiation were acceptable.

GENE EXPRESSION TESTING

- Formalin-fixed paraffin-embedded surgical tumor samples were analyzed for the expression levels of 31 CCP genes and 15 house-keeping genes by quantitative RT-PCR [6].
- The CCP score is an un-weighted average of 31 cell cycle genes normalized by the average of 15 house-keeping genes.
- The formula for the prognostic score (PS) is $20 \times (0.33 \times \text{CCP} + 0.52 \times \text{stage}) + 15$, where CCP score is rounded to the nearest tenth, and stage is treated as a numerical variable (stage IA = 1, stage IB = 2). The PS was rounded to the nearest integer.

STATISTICAL ANALYSIS

- Primary endpoint: 5-year lung cancer mortality (death from lung cancer within 5 years of surgery).
- The linear association of the CCP score and prognostic score with 5-year lung cancer mortality was evaluated with Cox proportional hazards regression.
- A PS threshold was used to define a binary variable for patients who have a “low” PS (< 28) or a “high” PS (≥ 28), as previously established [5].
- The log-rank test was employed to evaluate whether the 5-year lung cancer survival was significantly more favorable for patients in the low prognostic score group than in the high prognostic score group.

RESULTS

- 1137/1200 patients were evaluable based on available follow-up status and adequate tumor tissue quality to generate a CCP score.
- Demographic and other baseline characteristics for these evaluable patients are shown in Table 1.
- The baseline characteristic of morphology grade is derived from predominant histologic subtype (Table 2).

TABLE 2. PREDOMINANT HISTOLOGY AND MORPHOLOGY GRADE

Predominant Histologic Subtype	n (%)	Morphology Grade	n (%)
Lepidic	121 (10.6)	Low	160 (14.1)
Adenocarcinoma in situ (AIS)	1 (0.1)		
Minimally invasive adenocarcinoma (MIA)	38 (3.3)		
Acinar	464 (40.8)	Intermediate	712 (62.6)
Papillary	248 (21.8)		
Micropapillary	65 (5.7)	High	265 (23.3)
Solid with mucin production	153 (13.5)		
Invasive mucinous adenocarcinoma*	39 (3.4)		
Colloidal adenocarcinoma*	8 (0.7)		

TABLE 1. PATIENT CLINICAL CHARACTERISTICS AND MULTIVARIATE COX PROPORTIONAL HAZARDS REGRESSION ANALYSIS

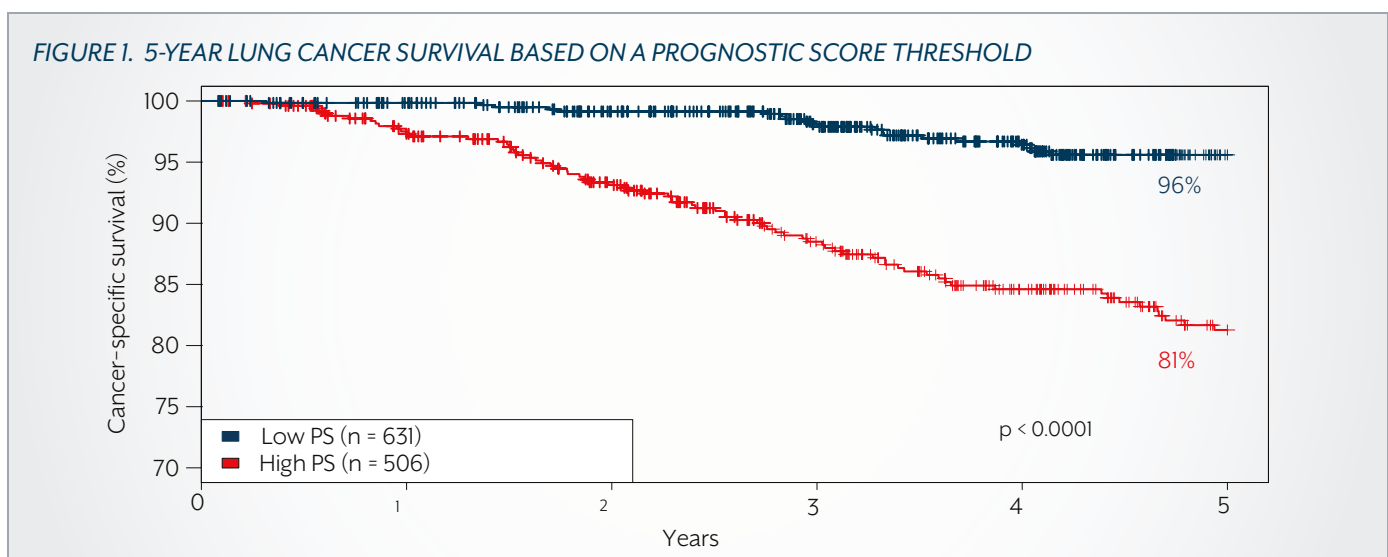
Characteristic	Statistic/ Category	Evaluable Analysis Set (N=1,137)	CCP Score		Prognostic Score	
			Hazard Ratio* (95% CI)	p-value	Hazard Ratio* (95% CI)	p-value
			1.54 (1.10, 2.15)	0.011	1.76 (1.14, 2.72)	0.011
Age at diagnosis (years)	n	1,137	1.03 (1.01, 1.06)	0.007	1.03 (1.01, 1.06)	0.007
	Mean (SD)	68.2 (10.00)				
	Median	69				
	Min, Max	23, 96				
Gender	Male	440 (38.7%)	1	0.188	1	0.188
	Female	697 (61.3%)	0.76 (0.50, 1.15)		0.76 (0.50, 1.15)	
Smoking Status	Never	200 (17.6%)	1	0.649	1	0.646
	Former	775 (68.2%)	1.35 (0.70, 2.94)		1.35 (0.70, 2.95)	
	Current	162 (14.2%)	1.46 (0.62, 3.63)		1.47 (0.62, 3.63)	
Tumor size (cm)	n	1,137	1.54 (1.17, 2.01)	0.003	1.54 (1.17, 2.00)	0.003
	Mean (SD)	2.2 (0.97)				
	Median	2.0				
	Min, Max	0.3, 5.0				
Pathological Stage	IA	810 (71.2%)	1	0.748	1	0.196
	IB	327 (28.8%)	0.88 (0.40, 1.92)		0.56 (0.23, 1.35)	
Pleural invasion	PLX/PL0	953 (83.8%)	1	0.138	1	0.140
	PL1	164 (14.4%)	1.81 (0.86, 3.81)		1.81 (0.86, 3.81)	
	PL2	20 (1.8%)	2.54 (0.87, 6.60)		2.53 (0.86, 6.57)	
Lymphatic invasion	Absent	769 (67.6%)	1	0.013	1	0.012
	Present	368 (32.4%)	1.79 (1.13, 2.86)		1.80 (1.14, 2.86)	
Vascular invasion	Absent	830 (73.0%)	1	0.055	1	0.055
	Present	307 (27.0%)	1.58 (0.99, 2.51)		1.58 (0.99, 2.51)	
Surgical procedure	Lobectomy/ Bilobectomy/ Pneumonectomy	851 (74.8%)	1	<0.0001	1	<0.0001
	Segmentectomy	98 (8.6%)	3.23 (1.59, 6.13)		3.23 (1.59, 6.12)	
	Wedge	188 (16.5%)	4.55 (2.71, 7.56)		4.54 (2.70, 7.54)	
Morphology grade	Low	160 (14.1%)	1	0.012	1	0.012
	Intermediate	712 (62.6%)	2.18 (0.77, 9.10)		2.18 (0.77, 9.11)	
	High	265 (23.3%)	3.78 (1.28, 16.16)		3.79 (1.29, 16.21)	

Abbr: Max = maximum, Min = minimum, SD = standard deviation, CI = confidence interval.

*Hazard ratios for CCP score and Prognostic score are per the interquartile range of each score, respectively.

RESULTS -CONTINUED-

- In the multivariate analysis, the CCP score was an independent significant prognostic marker ($p = 0.011$) of 5-year lung cancer mortality with a hazard ratio (HR) of 1.54 (95% CI = [1.10, 2.15]) (Table 1).
- Other significant variables included age at diagnosis ($p = 0.007$), tumor size ($p = 0.003$), lymphatic invasion ($p = 0.013$), surgical procedure ($p < 0.0001$) and morphology grade ($p = 0.012$) (Table 1).
- In a separate multivariate analysis, the prognostic score was an independent significant prognostic marker ($p = 0.011$) of 5-year lung cancer mortality with a hazard ratio (HR) of 1.76 (95% CI = [1.14, 2.72]) (Table 1).
- Other significant variables included age at diagnosis ($p = 0.007$), tumor size ($p = 0.003$), lymphatic invasion ($p = 0.012$), surgical procedure ($p < 0.0001$) and morphology grade ($p = 0.012$) (Table 1).
- The 5-year lung cancer survival rate was 96% for low prognostic score patients and 81% for high prognostic score patients (Figure 1).
 - The difference in survival rates between these two groups was significant ($p < 0.0001$).



CONCLUSIONS

- This study validates both the CCP score and PS as independent prognostic markers of lung cancer death in patients with Stage I lung adenocarcinoma treated with surgery.
- The CCP score and PS provide quantitative risk information above that captured by current NCCN high risk features.
- Patients with a high PS had nearly five times the rate of disease-related mortality than patients with a low PS (19% vs. 4%).
- Patients with resected stage I lung adenocarcinoma and a high CCP score and PS may be candidates for adjuvant therapy to reduce cancer related mortality.

REFERENCES

1. National Comprehensive Cancer Centers. NCCN clinical practice guidelines in oncology (NCCN Guidelines): Non-small cell lung cancer v3.2014. Available at http://www.nccn.org/professionals/physician_gls/pdf/nscl.pdf.
2. Pignon JP, Tribodet H, Scagliotti GV, et al. *J Clin Oncol*. 2008;26:3552–3559.
3. Subramanian J, Simon R. *J Natl Cancer Inst*. 2010, 102(7):464-474.
4. Wistuba II, Behrens C, Lombardi F, et al. *Clin Cancer Res*. 2013;19(22):6261-6271.
5. Bueno R, Hughes E, Wagner S, et al. *J Thorac Oncol*. 2015;10(1):67-73.
6. Cuzick J, Swanson GP, Fisher G, et al. *Lancet Oncol*. 2011;12(3):245-255.