# PROGNOSTIC AND CLINICAL UTILITY CAPABILITIES OF CELL CYCLE PROGRESSION TESTING, PROSTATE IMAGING-REPORTING AND DATA SYSTEM SCORING, AND CLINICOPATHOLOGIC DATA IN MANAGEMENT OF LOCALIZED PROSTATE CANCER

David Morris, MD<sup>1</sup>; J. Scott Woods, FNP-BC<sup>1</sup>; Lauren Lenz, MS<sup>2</sup>; Jennifer Logan, PhD<sup>2</sup>; Todd Cohen, PhD<sup>2</sup>; Steven Stone PhD<sup>2</sup> 1.Urology Associates, PC, Nashville TN 37209 2.Myriad Genetics, Inc., Salt Lake City, UT 84108

# BACKGROUND

- For men with newly diagnosed, localized prostate cancer (PrCa), determining whether it is safe to pursue active surveillance (AS) requires precise risk stratification.
- Multiparametric magnetic resonance imaging (mpMRI) with Prostate Imaging and Reporting and Data System (PI-RADS) scoring and the cell cycle progression (CCP) molecular prognostic test both have emerged as important tools for improving PrCa risk discrimination.
- We compared the prognostic and clinical utility capabilities among CCP testing, mpMRI with PI-RADS, and clinicopathologic data in selected medical management scenarios. We assessed:
- Distributions of CCP scores, clinical cell-cycle risk (CCR) scores, and clinicopathologic data relative to PI-RADS.
- Ability to predict tumor grade post-radical prostatectomy.
- Impact on the decision to pursue AS or curative therapy.

## METHODS

#### COHORT

- This was a retrospective, observational analysis of data from sequential patients (N=223, across two cohorts) from a single Urology community practice (January 2015-June 2018).
- Inclusion criteria: diagnosed with localized PrCa; had a PI-RADS version 2 score derived from mpMRI-ultrasound fusion targeted biopsy; and a concomitant biopsy CCP test result.
- Cohort 1 (n=157): Men newly diagnosed with localized PrCa, either with or without a previous negative biopsy.
- Cohort 2 (n=66): Men with localized PrCa who had initiated AS without CCP testing, but who subsequently received the test, with medical management informed by the result.

#### ANALYSIS

- The CCP test measured the expression of 31 CCP genes and 15 housekeeper genes in FFPE tissue using RT-PCR.
- The CCP score was calculated as the normalized expression of 31 CCP genes and was combined in a validated model with the UCSF Cancer of the Prostate Risk Assessment (CAPRA) score (0.57×CCP + 0.39×CAPRA) (Cuzick et al., Br J Cancer, 2015).
- Likelihood-ratio tests were used to determine predictor significance in both univariate and multivariate models.

Table 1.

Age at Colle

CCP

#### CAPR/

Low

Inter (3 –

High

### CCR

Belo

ADOV thres

#### **PI-RAD**

2/3 (| interr 4/5 ( very

(cm<sup>2</sup>

### PSA (

AS, active surveillance; CAPRA, UCSF Cancer of the Prostate Risk Assessment; CCP, cell cycle progression; IQR, interquartile ratio; PI-RADS, Prostate Imaging and Reporting Data System; PSA, prostate-specific antigen

Patient C	hara	cteristics					Table 2. Prediction of Gleason Score Category: Multivariate Analysis						Table 3. Impact on M	anagement Selection (Active Sur	<i>veillance vs.</i>	
	Cohort 1 Newly Diagnosed			Cohort 2	Con	nbined Cohorts	Post-RP			Diagnostic Biopsy			Definitive Treatment) Among Newly Diagnosed Patients (Cohort 1) (N=151)			
				On AS	1+2		Cohort 1, Newly Diagnosed			Cohort 1, Newly Diagnosed		Predictor	Odds Ratio (95% CI)	p-value		
	Median (IQR)		N	Median (IQR)	N	Median (IQR)	(n=56/157)			(n=157)			Univariate Models			
		or Frequency		or Frequency		or Frequency	Predictor	Odds Ratio (95% CI)	p-value	Predictor	Odds Ratio (95% CI)	p-value	CCP	2.47 (1.45, 4.45)	6.3 x 10 <sup>-4</sup>	
Biopsy	157	68 (61 72)	66	60 (63 75 73)	223	68 (62 72)		4.39			4.09		CAPRA	1.43 (1.15, 1.80)	8.7 x 10 <sup>-4</sup>	
ion	107	00 (01, 72)	00	00 (00.70, 70)	220	00 (02, 72)	CCP	(1.62, 14.81)	2.8 x 10 <sup>-3</sup>	CCP	(2.23, 8.07)	<b>1.7 x 10</b> -6	CCR	2.31 (1.51, 3.73)	<b>6.4 x 10</b> -5	
	157	-0.40 (-0.90, 0.00)	66	-0.60 (-1.18, -0.10)	223	-0.50 (-0.90, 0.00)	CAPRA	2.06 (1.24, 3.82)	<b>3.9 x 10</b> -3	PI-RADS	1.94 (0.97, 4.03)	0.060	PI-RADS	1.39 (0.80, 2.45)	0.244	
													CCP, CAPRA, PI-RAI	DS Multivariate Model		
							PI-RADS	0.43	0.23	PSA	1.01	0.778	CCP	1.98 (1.11, 3.67)	0.020	
) – 2)	47	29.9%	34	51.5%	81	36.3%	Combined	(0.09, 1.07) Cohorts (n=68/	223)	Combined	(0.95, 1.07) Cohorts (n=222	<b>5</b> )	CAPRA 1.31 (1.03, 1.69) 0.030			
ediate	82	52 2%	28	12 1%	110	19 3%		Odds Ratio	-		Odds Ratio		PI-RADS	0.98 (0.53, 1.79)	0.938	
	02	JZ.Z /0	20	<b>ΤΖ.Τ</b> /0	110	-J.J /0	Predictor	(95% CI)	p-value	Predictor	(95% CI)	p-value	CCR, PI-RADS Multivariate Model			
6 – 10)	28	17.8%	4	6.1%	32	14.3%	CCP	4.06	30x10 <sup>-3</sup>	CCP	3.05	1.1 x 10 <sup>-5</sup>	CCR	2.33 (1.49, 3.84)	<b>1.3 x 10</b> -4	
	157	1.10	66	0.80	223	1.00		(1.57, 12.69)			(1.83, 5.29)		PI-RADS	0.97 (0.53, 1.77)	0.92	
	101	(0.55, 1.94)		(0.21, 1.38)		(0.38, 1.63)	CAPRA	2.43 (1 50 4 45)	<b>1.1 x 10</b> -4	PI-RADS	1.99	0.029	CAPRA, UCSF Cancer of the Prostate Risk Assessment; CCP, cell cycle progression; CI, confidence interval; PI-RADS, Prostate Imaging and Reporting Data System Multivariate models adjusted for CCP, CAPRA, CCR, and PI-RADS.			
AS	58	36.9%	33	50.0%	91	40.8%		0.36			1.04					
010 (≤0.0)							PI-RADS	(0.079, 1.32)	0.13	PSA	(0.99, 1.10)	0.12				
AS old (>0.8)	99	63.1%	33	50.0%	132	59.2%	CAPRA, UCSF Cancer of the Prostate Risk Assessment; CCP, cell cycle progression; CI, confidence interval;							alysis, both CCP and CCR were si	gnificant and	
) <b>S</b>							PI-RADS, Prostat	e Imaging and Reporting	ng Data System;	PSA, prostate specific antigen; RP, radical prostatectomy			Each one-unit increase in score corresponded to an approximately two-			
w/							Figure 1 [	Distribution of (	CCP CAPR	A and CCR	Scores Across	PI-RADS	fold greater likeliho	od of selecting curative therapy (7	able 3).	
ediate)	50	31.8%	24	36.4%	74	33.2%	Score Gro	ups for Patient	s in Combi	elow the AS threshold significantly	reduced the					
gh/	107	68.2%	42	63.6%	149	66 8%	Combined Cohorts						probability of selecting curative therapy over AS [OR 0.29 (95% CI 0.14,			
gh)	107	00.270	12				0.59), p=6.1x10 <sup>-4</sup> ].									
te volume	157	41	66	35.8 (29.0, 50.4) 6.6 (4.9, 8.8)	223 223	39.5 (29.4, 50.5) 7.5 (5.4, 11.7)	호 이 이 이 이 이 이 이 이 이 이 이 이 이 이 이 이 이 이 이									
	157	(31, 50.3)	66				A/5 /ery						(Table 3).			
g/mL)		8.2 (5.8.12.4)					ligh∧							CONCLUSIONS		
		(3.0, 12.4)												GUNGLUSIUNS		

• On multivariate analysis, CCP was a significant predictor of higher-grade tumor (Gleason score  $\geq$ 4+3) after radical prostatectomy, with the resected tumor approximately four times more likely to harbor a higher-risk Gleason score with every one-unit increase in CCP score (Table 2).

• In combined Cohorts 1+2, weak but significant correlations were observed between PI-RADS and CCP, CAPRA, or CCR, suggesting that much of the prognostic information captured by these measures is independent (Figure 1).

ZA ZA

# RESULTS



Dotted line shows the validated CCR active surveillance (AS) threshold of 0.8 (Lin et al., Urol Oncol, 2018). CAPRA, UCSF Cancer of the Prostate Risk Assessment; CCP, cell cycle progression; CCR, clinical cell-cycle risk; PI-RADS, Prostate Imaging and Reporting Data System

Presented at SUO on December 4, 2019.

• In multiple scenarios, the CCP test was an independent and accurate prognostic measure that aided in risk stratification and medical management of localized PrCa.

• The CCP score was a better predictor of both tumor grade (at biopsy and after radical prostatectomy) and treatment selection than PI-RADS scores.

• A broad portfolio of measures, including targeted biopsy, clinicopathologic measures and molecular biomarker information, remains essential to ensure the most accurate and precise risk assessment to inform treatment selection.