

## Confirm mdx positive: GS 4+3=7

Age: 68 | PSA: 5.65 ng/mL | DRE: Enlarged | No family history

Previous negative prostate biopsy

Confirm mdx positive

GS 4+3=7 prostate cancer diagnosed

## **History**

02/22/2016 **Negative initial biopsy findings:** 

PSA Level: 6.02 ng/mL Number of Cores Collected: 14

Histology Findings: Benign Prostatic Tissue

Complications from Bx: None DRE Results: Enlarged

### **Results**

9/12/2017 Confirm mdx test results:



## **DNA Methylation Positive**

At time of Confirm mdx testing PSA Level: 5.65 ng/ml

PSA Level: 5.65 ng/mL DRE Results: Enlarged

### **Outcome**

03/14/2018 Prior to repeat biopsy

PSA Level: 6.48 ng/mL DRE results: Enlarged Prostate Volume: 58

Comorbidities: HBP, Hyperlipidemia

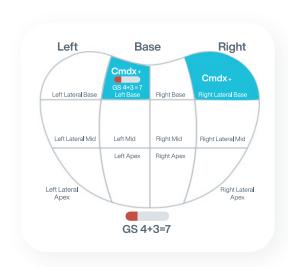
Chronic Medications: Tamsulosin, Lisinopril

MRI Results: PI-RADS 5 - Lesion left anterior apical

peripheral zone

## 06/11/2018 Repeat biopsy results

Pathology Result: Positive Clinical Disease Stage: T1 Cancer Grade: G3



# Confirm Dx for Prostate Cancer

## Patient Report

**PATIENT** 

Patient Name: Date of Birth: MRN/Patient#:

PATH: Benign
PSA: 5.65 ng/mL
DRE: Normal

### **SPECIMEN**

Specimen#:
Collection Date:
Received Date:
Report Date:

Specimen Type: Prostate Tissue Block

MDxH Accession#:

### ACCOUNT

Physician: Account: Address:

City/State/Zip:

## Patient Result: DNA Methylation Positive

The DNA methylation positive test result for this patient indicates an 36% likelihood of detecting prostate cancer, with a 21% probability for low-grade disease (GS  $\leq$  6) versus a 15% probability of high-grade disease (GS  $\geq$  7), on repeat biopsy.

## Likelihood of prostate cancer upon repeat biopsy

0% 100%

21%

Likelihood of detecting Gleason score ≤ 6 cancer

36%



Likelihood of detecting Gleason score ≥ 7 cancer

The ConfirmMDx test result indicating the likelihood of GS  $\leq$  6 and GS  $\geq$  7 prostate cancer being detected on repeat biopsy is calculated by incorporating DNA methylation intensity with clinical risk factors, including PSA, DRE, age, and histopathology of the previous biopsy, based on a clinical model that yields an area under the curve (AUC) of 0.762 (95% CI: 0.679-0.844). Performance is based on the presence of all relevant data elements; if all data are not available, or 5 $\alpha$ -reductase inhibitors (5ARI) have been administered to decrease serum PSA values, results should be interpreted with caution since the AUC of the test may vary. Cancer association with DNA methylation of the ConfirmMDx gene markers has been reported on ~4,500 patients. 1-55

DNA Methylation Status Table			
Biopsy Site	GSTP1 Methylation	APC Methylation	RASSF1 Methylation
Left Lateral Base:	Negative	Negative	Negative
Left Lateral Mid:	Negative	Negative	Negative
Left Lateral Apex:	Negative	Negative	Negative
Left Base:	Positive	Negative	Negative
Left Mid:	Negative	Negative	Negative
Left Apex:	Negative	Negative	Negative
Left Transition Zone:			
Right Base:	Negative	Negative	Negative
Right Mid:	Negative	Negative	Negative
Right Apex:	Negative	Negative	Negative
Right Lateral Base:	Positive	Negative	Negative
Right Lateral Mid:	Negative	Negative	Negative
Right Lateral Apex:	Negative	Negative	Negative
Right Transition Zone:	Negative	Negative	Negative

## Distribution of DNA Methylation Diagram Posterior View

