# ConfirmMDx Positive: GS 4+3=7

Confirm MDx for Prostate Cancer

Age: 68 I PSA: 6 ng/mL I DRE: Enlarged

Previous negative prostate biopsy > ConfirmMDx Positive > GS 4+3=7 prostate cancer diagnosed

ConfirmMDx Result: POSITIVE DNA METHYLATION

68 year old | PSA 6 ng/mL | DRE: Enlarged | No Family History

#### **HISTORY**

February 2016 **Negative Initial Biopsy Findings:** 

PSA Level: 6 ng/mL

Number of Cores Collected: 14

Histology Findings: Benign Prostatic Tissue

Complications from Bx: None DRE Results: Enlarged

#### **RESULTS**

September 2017 ConfirmMDx Results:

Following the initial negative biopsy results, the treating physician ordered a ConfirmMDx test.



### ConfirmMDx DNA Methylation Positive

At time of ConfirmMDx testing

PSA Level: 6 ng/mL DRE Results: Enlarged

#### **OUTCOME**

March 2018 **Prior to Repeat Bx:** 

PSA Level Prior to Repeat Bx: 6 ng/mL

DRE results: **Enlarged**Prostate Volume: **58** 

Comorbidities: HBP, Hyperlipidemia

Chronic Medications: Tamsulosin, Lisinopril

MRI Results: PI-RADS 5-Lesion left anterior

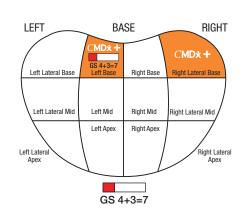
apical peripheral zone

June 2018 **Repeat Biopsy Results:** 

Pathology Results of Repeat Bx: Positive

Clinical Disease Stage: T1

Cancer Grade: G3



# Patient Report



#### **PATIENT**

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

PATH: Benign
PSA: 6 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



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** APC RASSF1 **Biopsy Site** Methylation Methylation 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: Negative Negative Negative Right Base: Negative **Negative** Negative Right Mid: Negative Negative Negative Right Apex: Right Lateral Base: **Positive** Negative Negative Negative **Negative** Negative Right Lateral Mid: Negative Right Lateral Apex: Negative **Negative** Negative **Negative** Negative Right Transition Zone:

# Distribution of DNA Methylation Diagram

