

Confirm mdx positive: GS 4+3=7

Age: 71 | PSA: 7.1 ng/mL | DRE: Enlarged | No family history

Previous negative prostate biopsy

Confirm mdx positive

GS 4+3=7 prostate cancer diagnosed

History

11/16/2016 **Negative initial biopsy findings:**

PSA Level: 7.1 ng/mL

Number of Cores Collected: 12

Histology Findings: Benign Prostatic Tissue

Complications from Bx: None

DRE Results: Normal

Results

10/20/2017 Confirm mdx test results:



DNA Methylation Positive

At time of Confirm mdx testing

PSA Level: **7.1 ng/mL** DRE Results: **Normal**

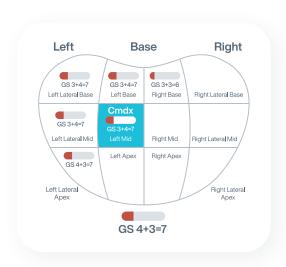
Outcome

11/02/2017 Prior to repeat biopsy

PSA Level: 1.56 ng/mL DRE results: Enlarged Prostate Volume: 35 MRI Results: PI-RADS 5 Date of MRI: 11/2/17

12/18/2017 Repeat biopsy results

Pathology Result: Positive Clinical Disease Stage: T1 Cancer Grade: G3 (GS 4+3)



Patient Report



PATIENT

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

PATH: Benign
PSA: 7.1 ng/mL
DRE: Not Suspicious

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 60% likelihood of detecting prostate cancer, with a 27% probability for low-grade disease (GS \leq 6) versus a 33% 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 α -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: Negative **Negative** Negative Left Mid: Negative **Positive Positive** Left Apex: Negative Negative Negative Left Transition Zone: **Negative** Negative Right Base: Negative Negative **Negative** Negative Right Mid: **Negative** Negative Negative Right Apex: Right Lateral Base: Negative Negative Negative Right Lateral Mid: Negative **Negative** Negative Negative Negative Negative Right Lateral Apex: Right Transition Zone:

Distribution of DNA Methylation Diagram

