

Confirm mdx positive: GS 4+4=8

Age: 77 | PSA: 9 ng/mL | DRE: Not Suspicious | No family history

Previous negative prostate biopsy

Confirm mdx positive

GS 4+4=8 prostate cancer diagnosed

History

09/09/2016 Initial biopsy findings:

PSA Level: 9 ng/mL

Number of Cores Collected: 12

Histology Findings: Benign Prostatic Tissue

Complications from Bx: None DRE Results: Not Suspicious

Results

06/23/2017 Confirm mdx test results:



DNA Methylation Positive

At time of Confirm mdx testing

PSA Level: 9 ng/mL

DRE Results: Not Suspicious

Outcome

01/05/2018 Prior to repeat biopsy

PSA Level: 18.62 ng/mL DRE results: Enlarged Prostate Volume: 40.38

Comorbidities: **A-Fib, Hypertension, DM**Chronic Medications: **Eliquis, Glimepiride**

MRI Results: PI-RADS 5
Date of MRI: 01/28/18

02/04/2018 Repeat biopsy results

Pathology Result: Positive Clinical Disease Stage: T1c

Cancer Grade: G4



Patient Report



PATIENT

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

PATH: Benign
PSA: 9 ng/mL
DRE: Not Suspicious

DNA Methylation Status Table

SPECIMEN

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

Specimen Type: Prostate Tissue Block

MDxH Accession#:

ACCOUNT

100%

Physician: Account: Address:

City/State/Zip:

Patient Result: DNA Methylation Positive

The DNA methylation positive test result for this patient indicates an 41% likelihood of detecting prostate cancer, with a 22% probability for low-grade disease (GS \leq 6) versus a 19% probability of high-grade disease (GS \geq 7), on repeat biopsy.

Likelihood of prostate cancer upon repeat biopsy

41%

Likelihood of detecting

Gleason score ≤ 6 cancer

19%

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 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

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 Negative Negative Left Apex: Negative Negative Negative Left Transition Zone: **Negative** Negative Right Base: **Positive** Negative Negative Right Mid: Negative Negative Negative Negative Right Apex: Right Lateral Base: Negative Negative Negative Right Lateral Mid: Negative Negative Negative

Posterior View Base Right Transition Zone Rosterior View Right Right Transition Zone

Distribution of DNA Methylation Diagram

Negative

Negative

Right Lateral Apex:

Right Transition Zone:

Negative

Negative

Negative

Negative