

DECIPHER RADICAL PROSTATECTOMY REPORT

PATIENT DETAILS

Patient Name:
 MRN/Patient ID:
 Date of Birth:
 Date of Prostatectomy:
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 Pathology Laboratory:
 Pathologist:
 Address:

ORDER INFORMATION

Order Date:
 Specimen Received Date:
 GenomeDx Accession ID:
 Specimen ID: S
 Ordering Physician:
 Clinic/Hospital Name:
 Clinic/Hospital Address:
 Additional Physician:

CLINICAL DETAILS

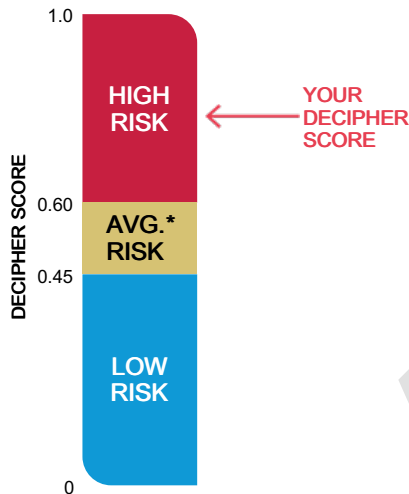
Pre-operative PSA (ng/mL): **21.8**

Specimen Type: **Radical Prostatectomy**

Grade Group: **5**

- SM+
 EPE
 SVI
 LNI
 BCR
 Tertiary Gleason 5
 Other: Preoperative PSA (≥ 20 ng/mL), Perineural Invasion, Capsular Invasion

YOUR DECIPHER RESULT: GENOMIC HIGH RISK



DECIPHER SCORE: 0.78	
Risk at RP - Percent Likelihood	
5-Year Metastasis	25.5%
10-Year Prostate Cancer Specific Mortality	15.3%
INTERPRETATION References on reverse	
<p>Clinical studies concluded that Decipher high risk men with adverse pathology have a poor prognosis overall.^{1-3,8} These men may benefit from adjuvant or early salvage radiotherapy and consideration for clinical trials.^{4,6}</p> <p>Relevant findings from published clinical studies: Patients with Decipher high risk had 77.5% 5-year metastasis free survival and 70.0% 10-year cause specific survival.^{1,2,3,9} For these patients there was improved metastasis-free survival favoring adjuvant and early salvage post-operative radiotherapy compared to post-operative observation.^{5,6,7}</p> <p>In patients with PSA rise or biochemical recurrence after surgery that received salvage radiotherapy, only 66.9% remained metastasis free after 5 years.⁴</p> <p>Additional Comment: Tumor heterogeneity exists in most cancers, including prostate cancer. To ensure that the most aggressive tumor tissue was evaluated, analysis was repeated on two different areas of the tumor. The higher Decipher risk score was reported.</p>	

*Average clinical risk refers to the average cohort risk of metastasis at 5 years post radical prostatectomy (RP). The average cumulative incidence of metastasis was 6.0% at 5 years post RP, as reported by Karnes et al., 2013 from analysis of a cohort of 1,010 men with intermediate and high risk clinical features who received RP as first line treatment at the Mayo Clinic between 2000 and 2006.¹

Five-year probability of metastasis endpoint: Decipher uses the genomic risk score to predict the 5-year probability of metastasis from the time of RP. Probabilities were generated from a Cox proportional hazards model based upon a cohort of 1,010 men with intermediate and high risk clinical features with a median 6.9 years of follow up.¹ Decipher had an AUC of 0.76-0.85 in multiple clinical validation studies for prediction of metastasis.^{1,6,9,10} Percent likelihood for this endpoint ranges from 0.3-67%.

Ten-year probability of prostate cancer specific mortality (PCSM) endpoint: Decipher uses the genomic risk score to predict the 10-year probability of PCSM from the time of RP. Probabilities are generated from a logistic regression analysis based upon a cohort of 557 patients with 112 prostate cancer deaths within 10 years post RP. These probabilities are adjusted for a PCSM cumulative incidence of 5% at 10 years post RP.¹¹ All non-PCSM patients in the study had at least 10 years of follow-up. Decipher had an AUC of 0.73 in predicting PCSM.^{2,11,13} Percent likelihood for this endpoint ranges from 0.7-30.5%.

GenomeDx Medical Director (Signature)
 Medical Directors: Timothy J. Triche, MD, PhD | Doug Dolginow, MD

Report Date

Disclaimer: The Decipher test was developed and its performance characteristics were determined by GenomeDx. GenomeDx Biosciences Laboratory is certified under the Clinical Laboratory Improvement Amendments of 1988 (CLIA '88) as qualified to perform high complexity clinical laboratory testing. This test is used for clinical purposes and should not be regarded as investigational or for research. This test has not been cleared or approved by the U.S. Food and Drug Administration. Summary of surgical pathology report is provided for the convenience of the Ordering Physician. Please refer to Referring Pathologist's original pathology report to guide treatment decisions.

DECIPHER RADICAL PROSTATECTOMY REPORT

TEST DESCRIPTION

Decipher uses oligonucleotide microarrays to measure 22 RNA expression biomarkers, extracted from formalin fixed paraffin embedded (FFPE) primary prostate adenocarcinoma specimens, to derive a Decipher score and corresponding probability of:

- 5-year probability of clinical metastasis
- 10-year prostate cancer specific mortality

The Decipher score ranges from 0 to 1.0.

INTENDED USE

Results from Decipher are intended for use by the physician and patient as an adjunct to conventional clinical variables and models currently used for determining prognosis and treatment of prostate cancer patients after radical prostatectomy. Decipher is intended for use in those patients who present with specific risk factors for the recurrence of prostate cancer after radical prostatectomy: (1) stage T2 disease with positive surgical margins, or (2) stage T3 disease, or (3) rising prostate-specific antigen (PSA) levels after initial PSA nadir when PSA is undetectable, or (4) pre-operative PSA of 20 ng/mL or higher, or (5) lymph node involvement (LNI), or (6) high Gleason Score of 8-10.

CONFIDENCE INTERVALS

- Probability of 5-year metastasis reported here has a 95% confidence interval of 16.8% to 33.4%
- Probability of 10-year Prostate Cancer Specific Mortality reported here has a 95% confidence interval of 7.8% to 22.7%

DEFINITIONS

Clinically High Risk: These men are at high risk of clinical metastasis as defined in the Karnes, et al. study cohort inclusion criteria, which was any of: pre-operative Prostate-Specific Antigen (PSA) > 20 ng/mL; pathologic Gleason score > 8; Seminal Vesicle Invasion; GPSM nomogram > 10.¹⁰

Average Clinical Risk: Refers to the average cohort risk of metastasis at 5 years post RP. The average cumulative incidence of metastasis was 6.0% at 5 years post RP as reported by Karnes et al., 2013 from analysis of a cohort of 1,010 men with intermediate and high risk clinical features who received RP as first line treatment at the Mayo Clinic between 2000 and 2006.¹

Genomic Low or High Risk: Based on the individualized genomic risk of metastasis identified by Decipher, these men have significantly higher (Decipher result > 0.6) or lower (Decipher result < 0.45) risk than the average clinical risk as defined above. These Decipher risk categories were selected by optimizing both the partial likelihood and hazard ratios in a series of Cox models. The categories were trained using data from the Karnes, et al. study and validated in Ross, et al. study.^{1,2}

Clinical Metastasis: Regional (e.g., to regional lymph nodes) or distant (e.g., to bones) spread of cancer from the prostate as confirmed by positive CT and/or bone scan.

REFERENCES

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