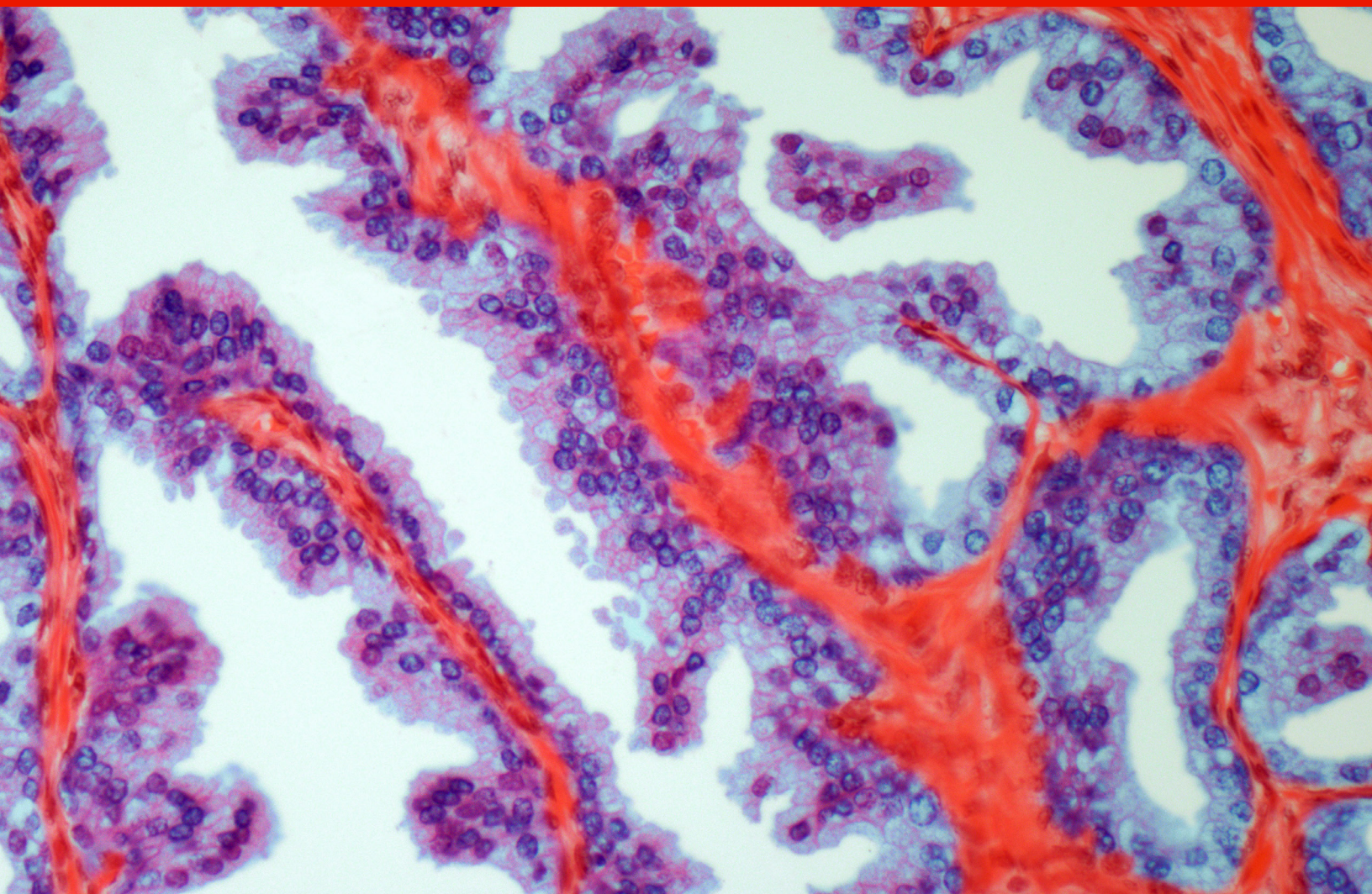


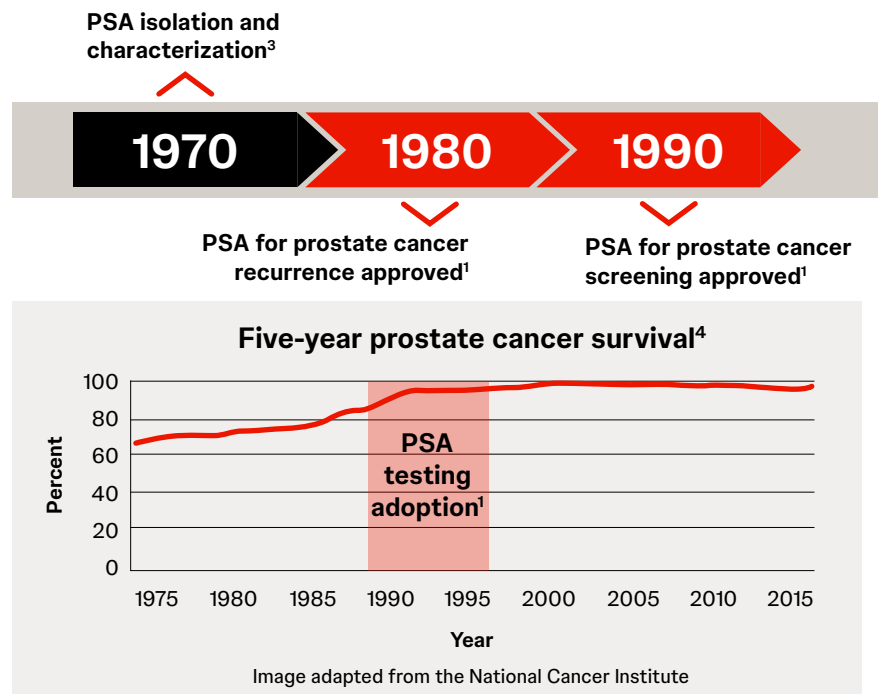
Prostate Cancer Molecular Biomarkers and Guideline Recommendations

J&J

Precision Medicine



Biomarkers have historically played an important role in the diagnosis of prostate cancer^{1,2}



Introduction of PSA screening led to earlier stage at diagnosis and a **reduction in prostate cancer mortality by 46.0%–63.7%**⁵

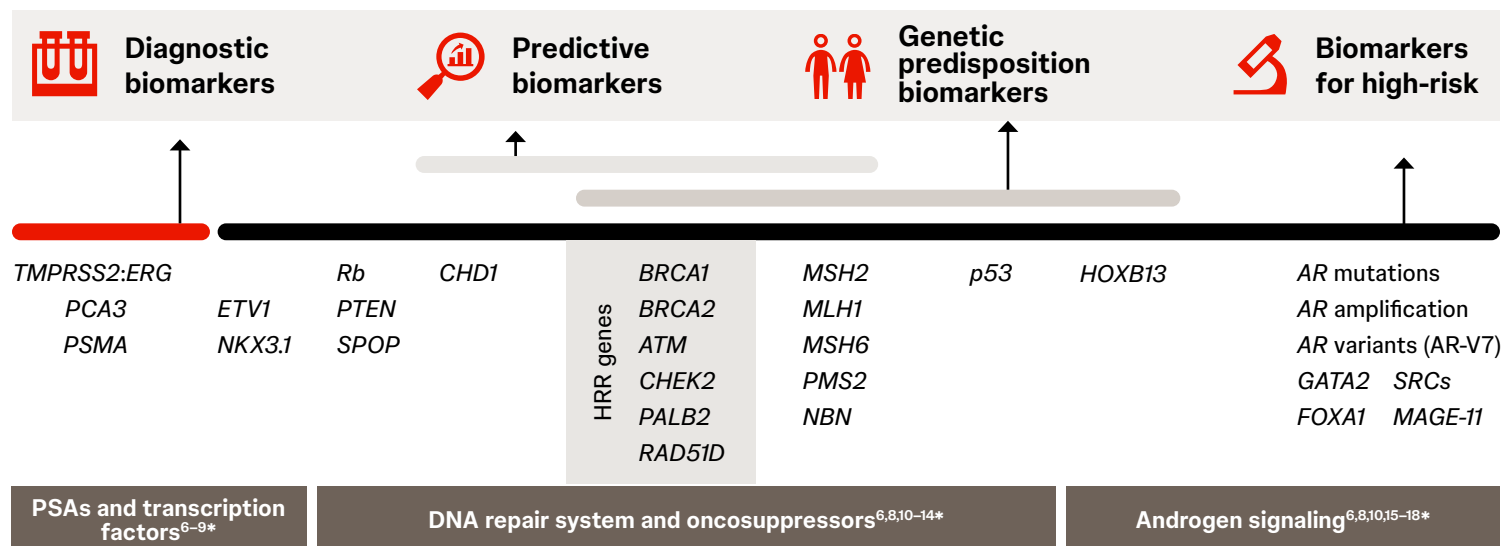
PSA is a valuable biomarker^{1,2,6*}

- For diagnosing prostate cancer
- For diagnosing recurrence
- As an indicator of treatment response

However, additional biomarkers are needed^{1,2}:

- To determine susceptibility for prostate cancer
- To improve risk stratification
- To inform personalized treatment strategies

Tumor-based molecular assays and germline testing are tools which can assist with disease management through risk stratification, diagnosis, and treatment selection^{6,7*}



Prostate cancer biomarkers may be identified through both germline and somatic testing^{6,19*}

Germline testing

Identifies biomarkers in DNA that are **inherited** and expressed in cells from anywhere in the body^{10,20}

 Blood

 Saliva

Somatic testing

Identifies new mutations in DNA that are **expressed by cancer cells**^{10,20}

 Tumor

 Blood (ctDNA)

There are limitations for performing germline or somatic testing alone:

Up to 50% of actionable DNA damage response (DDR) mutations may be missed by germline testing alone²¹

Up to 20% of patients will have mutations missed by somatic testing alone²⁰

Opportunity to optimize

Combining somatic and germline testing may be appropriate for assessing genetic disposition, prognosis, and predicting likelihood of treatment response in some patients with prostate cancer^{6,19*}

Prognostic biomarker tests evaluate gene expression to identify aggressive prostate cancer^{22†}

| | Oncotype Dx GPS ²²⁻²⁴ <i>Genomic Health</i> | ProMark ^{9,22,23} <i>Metamark</i> | Decipher ^{22,25,26} <i>Decipher Bioscience (formerly Genome Dx)</i> | Prolaris ^{22,23,27,28} <i>Myriad Genetics</i> |
|-----------|--|--|---|---|
| Evaluates | mRNA expression levels of 17 genes across 4 pathways in tumor tissue | Expression of 8 proteins in tumor tissue with multiplex immunofluorescence imaging | mRNA expression levels of 22 genes in tumor tissue | mRNA expression levels of 10 cell-cycle progression genes and 6 reference genes in tumor tissue |
| Indicates | Tumor aggressiveness | Tumor aggressiveness | Risk of metastasis and death | Likelihood of disease progression and tumor aggressiveness |
| | After positive biopsy ^{22,23,29} | | After definitive therapy ^{22,23,29} | |

Prognostic biomarker test results can help determine which patients require treatment^{6,23,29*}

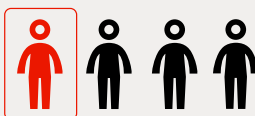
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†The presented list of biomarker tests for disease prognosis in prostate cancer is not exhaustive.²²

Genes with established or emerging potential clinical actionability may impact patient management^{10,30}



of metastatic castration-resistant prostate cancer (mCRPC) tumors **contain a potentially actionable mutation**³⁰

~ 1 in 4 patients



with advanced prostate cancer **have mutations** in **DDR genes**, which include the HRR and MMR pathways^{30,31}

Common MMR pathway biomarkers in prostate cancer^{32,33}

MMR pathway gene mutation

Key MMR biomarkers^{32,33}:

- *MLH1*
- *MSH2*
- *MSH6*
- *PMS2*

Predict treatment response

- dMMR can predict response to immune checkpoint inhibitors³⁴

Understand genetic disposition

- **Germline** mutations in these genes can also provide insights into both patient and familial risk for prostate cancer¹⁰

Mismatch repair deficit

Inform prognosis

- dMMR is associated with more aggressive disease and high Gleason scores³⁴

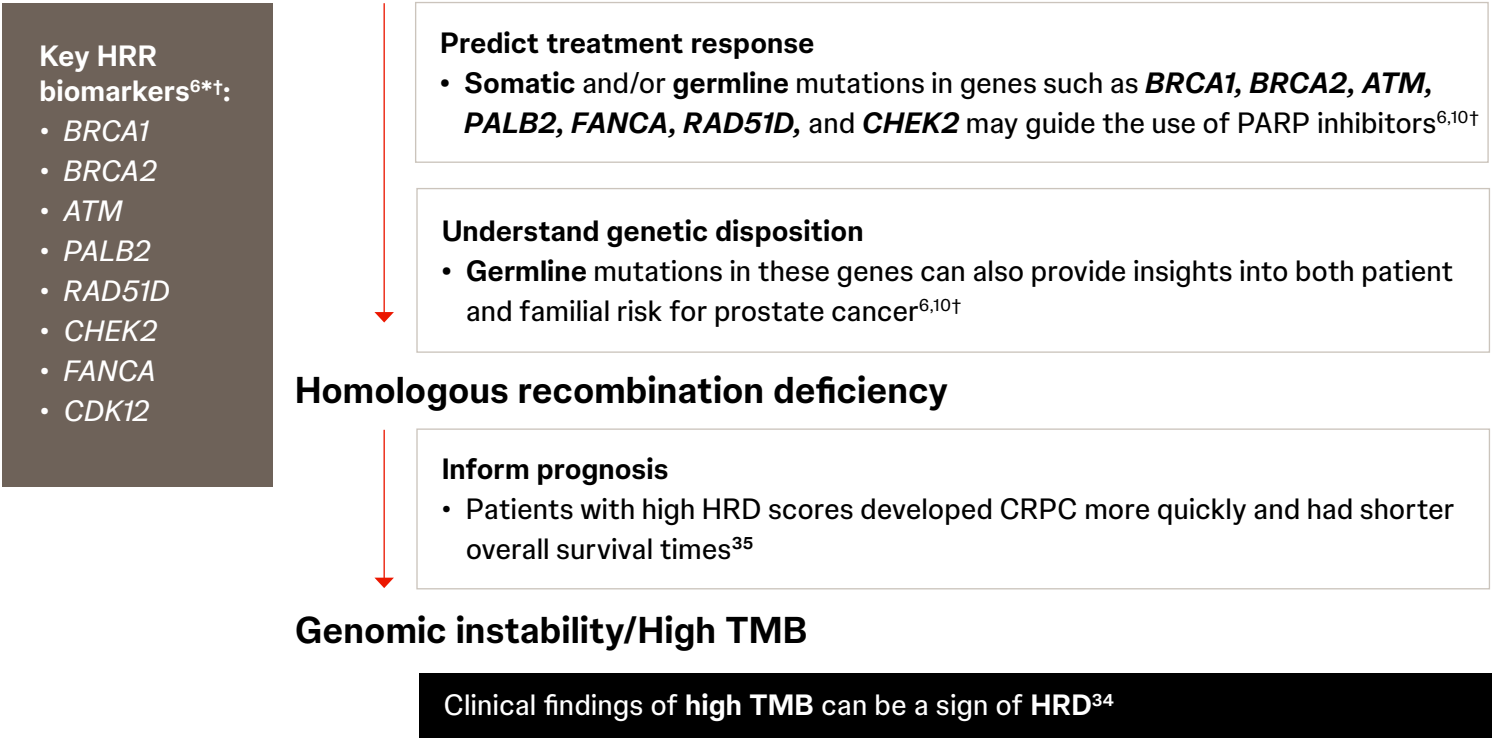
MSI or High TMB

Microsatellite instability (MSI)

- Used to evaluate dMMR pathway³³
- dMMR and MSI-H may be associated with high TMB³⁴


HRR pathway mutations are informative prostate cancer biomarkers^{10,35}

HRR pathway gene mutation




Among other HRR pathway genes, *BRCA* alterations are informative in prostate cancer^{6,36†}

| Percent of patients with <i>BRCA1/2</i> alterations ³⁶ | Germline | Somatic |
|---|-----------|---------|
| | Any stage | 7% |
| | mCRPC | 11% |




Understanding predisposition

7%–26% of males with germline *BRCA1* alterations and 19%–61% of males with germline *BRCA2* alterations will develop prostate cancer^{19,37†}



Prognostic implications

In one retrospective study of 1904 males diagnosed with prostate cancer, 12-year **survival rate in patients with a *BRCA2* alteration was >30% lower** compared with those without a *BRCA* alteration³⁸



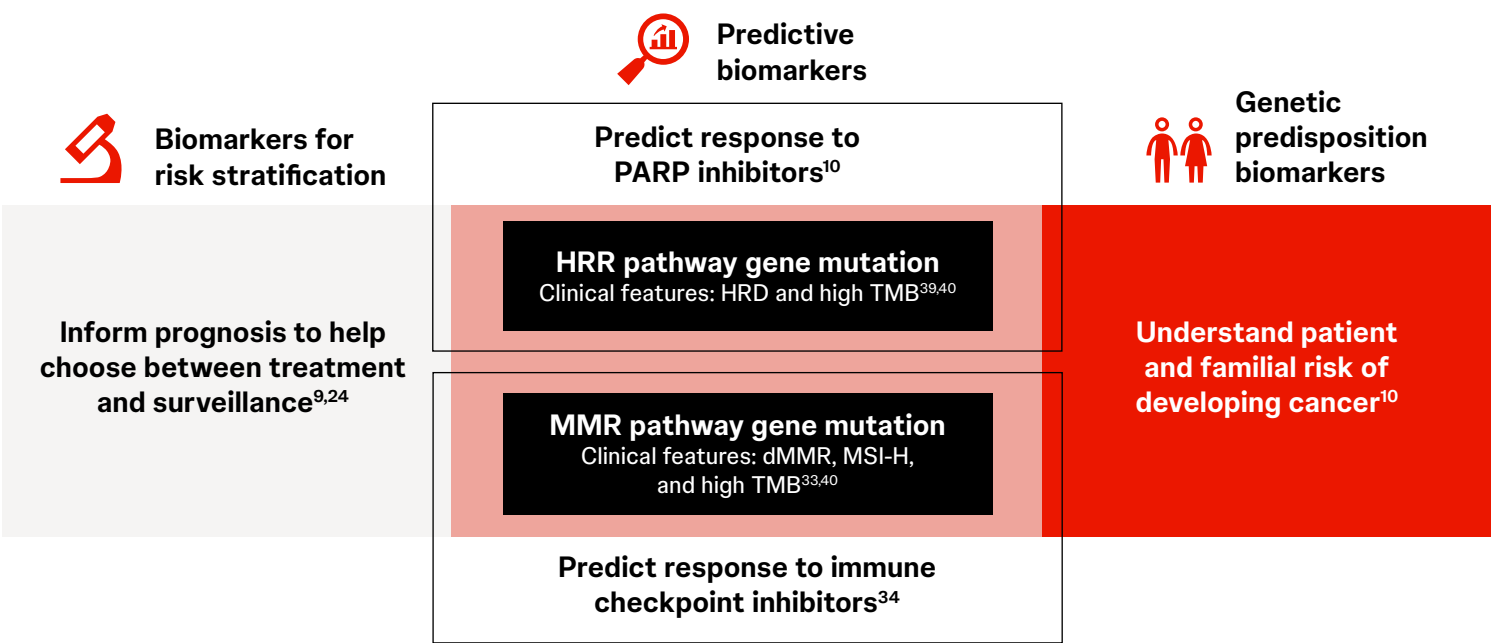
Predictive implications

BRCA alterations may predict response to PARPi^{6†}

Although mutations in DDR genes are associated with aggressive disease, identification of these biomarkers via both somatic and germline testing can inform treatment decisions and may improve prognosis^{6,34†}

*This is not a complete list of all HRR biomarkers in prostate cancer. †NCCN makes no warranties of any kind whatsoever regarding their content, use or application and disclaims any responsibility for their application or use in any way.

Germline and somatic testing for DDR gene mutations can help guide disease management approaches^{6,31,35*}



Actionable biomarkers in prostate cancer that can inform treatment decisions may be detected by different assays with unique capabilities^{41–47}


| Test type | Single-biomarker test ^{45,48} | | Broad-based panel ^{45,48} |
|--------------------------------|--|---|---|
| | Immunohistochemistry (IHC) ^{41,42} | Polymerase chain reaction (PCR) ^{42–44} | Next-generation sequencing (NGS) ^{42,45–47} |
| Alteration type(s) detected | Overexpression/amplification, protein expression | Single-nucleotide variant (SNV), copy number variant (CNV), insertions-deletions (indels), known gene fusions | SNV, CNV, indels, genomic rearrangements |
| Actionable biomarkers detected | MLH1, MSH2, MSH6, PMS2 | MSI, <i>MLH1</i> , <i>MSH2</i> , <i>MSH6</i> , <i>PMS2</i> | MSI, TMB, <i>MLH1</i> , <i>MSH2</i> , <i>MSH6</i> , <i>PMS2</i> , <i>ATM</i> , <i>BRCA1</i> , <i>BRCA2</i> , <i>CDK12</i> , <i>CHEK2</i> , <i>PALB2</i> , <i>FANCA</i> , <i>NBN</i> |

Opportunity to optimize


NGS is an efficient way to test for multiple alterations^{42,45}

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Several emerging biomarkers can be used for early prostate cancer detection^{49,50}

 PSA testing remains the first-line screening test of choice at this time,⁵¹ but may yield false negatives or overdiagnosis and overtreatment⁵²


New tests based on other biomarkers may help address these challenges⁵²



Extracellular vesicles containing nucleic acids, proteins, and metabolites may serve as highly sensitive biomarkers for the early diagnosis of prostate cancer^{49,50}



Analysis of plasma cell-free DNA (cfDNA) methylation is being explored for early detection and tumor classification⁴⁹



A panel of six multi-omics biomarkers, *TELO2*, *ZMYND19*, miR-143, miR-378a, and methylation status of two CpG loci, may be a biomarker for the early detection of recurrent prostate cancer⁴⁹




Many emerging prostate cancer biomarkers can be detected by liquid biopsy^{50,53}

Liquid biopsy can detect new biomarkers including ctDNA, ctRNA, CTCs, and EVPs^{50,53}

Liquid biopsy is a **non-invasive** tool that can aid in **early detection** and **diagnosis**, understanding **prognosis**, and guiding **treatment selection** for prostate cancer^{50,53}

| | Circulating tumor DNA (ctDNA) | Circulating tumor RNA (ctRNA) | Circulating tumor cells (CTCs) | Extracellular vesicles and particles (EVPs) |
|---|--|--|--|---|
| | Emerging biomarkers include DNA methylation, copy number variants, and rearrangements ^{54–56} | Primarily noncoding RNAs such as miRNAs ^{50,54} | EpCAM-positive cells circulating in blood that shed from primary/metastatic tumors ⁵³ | Extracellular vesicles and particles that harbor proteins and nucleic acids ⁵⁰ |
| Diagnostic ⁵⁰ | ✓ | ✓ | | ✓ |
| Distinguishes from BPH ⁵⁰ | | ✓ | | ✓ |
| Predicts metastasis ⁵⁰ | ✓ | | ✓ | ✓ |
| Prognostic ⁵⁰ | ✓ | ✓ | ✓ | ✓ |
| Predicts treatment response ⁵⁰ | ✓ | ✓ | | ✓ |

Germline testing guidelines for prognostic and predictive biomarkers and familial risk assessment

| | National Comprehensive Cancer Network® (NCCN®) ^{6,19*} | AUA/ASTRO ^{57†} (localized) | AUA/SUO ^{58†} (advanced) | ASCO ⁵⁹ (metastatic) |
|---|---|---|--------------------------------------|------------------------------------|
|  Who to test | Patients with prostate cancer and any of the following: | | | |
| | • Pathogenic somatic mutations in cancer-risk genes | ✓ | | ✓ |
| | • Metastatic prostate cancer | ✓ | ✓ | ✓ |
| | • (Strong) [‡] family history of (related) [‡] cancer | ✓ | ✓ | |
| | • Positive family history of germline variants | ✓ | ✓ | ✓ |
| | • Ashkenazi Jewish ancestry | ✓ | ✓ | |
| | • Strong personal history of related cancers | | ✓ | |
|  What to test | • High-risk [§] /adverse [‡] tumor characteristics | ✓ | ✓ | ✓ |
| | <i>BRCA1</i> and <i>BRCA2</i> | ✓ | ✓ | ✓ |
| | <i>ATM</i> and <i>CHEK2</i> | ✓ | ✓ | ✓ |
| | <i>PALB2</i> | | ✓ | ✓ |
| | <i>HOXB13</i> and <i>TP53</i> | ✓ | ✓ | |
| | <i>NBN</i> , <i>MLH1</i> , <i>MSH2</i> , <i>MSH6</i> , and <i>PMS2</i> | | ✓ | |
|  When to test | RAD51D | | ✓ | |
| | At initial diagnosis, where it is likely to inform treatment and clinical options | ✓ | | ✓ |
| | Offer germline testing to inform discussions around prognosis | | ✓ [#] | ✓ |
| Additional considerations | Germline testing should include appropriate genetic counseling | ✓ | ✓ ^{**} | ✓ |




There is consensus across clinical guidelines shown here that **germline testing may include appropriate genetic counseling**^{6,19,57–59*}

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†Where gaps in the clinical evidence exist, the AUA Panel provides guidance in the form of Clinical Principles or Expert Opinions.^{57,58} ‡AUA-specific language.






§NCCN-specific language. ¶AUA/SUO guideline notes HRR genes as examples and are not limited to those in the table.⁵⁸ ¶AUA/SUO guidelines note that germline testing to inform discussions around prognosis applies to mCRPC patients (if not already performed) and mCSPC patients.⁵⁸ *Germline testing is only offered after patient assessment of risk factors.⁵⁷ **AUA/SUO guidelines note that genetic counseling applies for mCSPC patients.⁵⁸

Somatic tumor testing guidelines for prognostic and predictive biomarkers

| | | NCCN ^{6*} | AUA/SUO ^{58†} (advanced) | ASCO ⁵⁹ (metastatic) |
|---|---|--------------------|--------------------------------------|------------------------------------|
|  Who to test | Patients with metastatic prostate cancer | ✓ | ✓ | ✓ |
| | Consider for patients with regional (N1) prostate cancer | ✓ | | |
|  Biomarkers to consider | Multigene testing for HRR genes [‡] <i>BRCA1, BRCA2</i> | ✓ | ✓ | ✓ |
| | <div>HRR genes noted by guidelines</div> <i>ATM, PALB2, RAD51D, CHEK2</i> | ✓ | ✓ | |
| | <i>FANCA, CDK12</i> | ✓ | | |
| | MSI-H or dMMR | ✓ | ✓ | ✓ |
| | TMB [§] | ✓ | ✓ | ✓ |
|  When to test | At metastatic diagnosis, if not previously tested | ✓ | | ✓ |
| | Consider retesting at time of progression | ✓ | | ✓ |

When considering testing, it is important to check historical records for prior germline and/or somatic testing to be aligned with guidelines^{6*}

Germline and somatic testing may be underutilized in practice^{60–63}

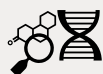
| Identified barriers to testing ⁶³ | | | | |
|---|--|--|---|---|
|  Access to genetic counselors |  Insurance coverage and cost |  Clinic workflow |  Time and space availability |  Access to resources for provider and patient education |

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†Where gaps in the clinical evidence existed, the AUA Panel provides guidance in the form of Clinical Principles or Expert Opinions.⁵⁸ ‡AUA/SUO guideline notes HRR genes as examples and are not limited to those in the table.⁵⁸ §TMB testing is recommended in patients with metastatic CRPC (mCRPC). A recommendation is not made for patients with regional (N1) prostate cancer.⁶

Summary



Advances in biomarker research have impacted multiple domains of prostate cancer management including **diagnosing** prostate cancer, predicting likelihood of **treatment response**, understanding **genetic disposition**, and identifying and managing **high-risk individuals**^{6,7*}



Current clinically actionable prostate cancer biomarkers are **primarily genes involved in DNA repair such as HRR genes** (*BRCA1*, *BRCA2*, *ATM*, *PALB2*, *RAD51D*, *CHEK2*, *FANCA*, and *CDK12*) and **MMR genes** (*MLH1*, *MSH2*, *MSH6*, and *PMS2*)^{6*}



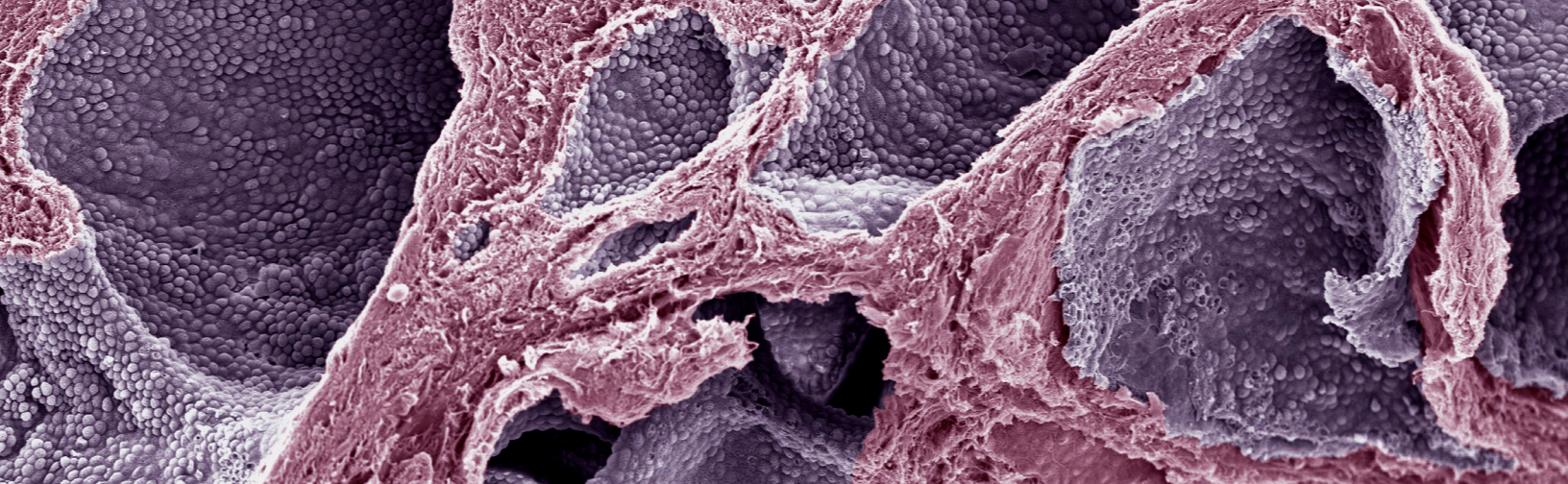
Combining both somatic and germline testing according to clinical guidelines **can offer a comprehensive view** for genetic disposition, prognosis, and predicting likelihood of treatment response^{6,19,57,58*}



Prostate cancer **biomarkers continue to emerge** with emphasis on biomarkers detectable by noninvasive liquid biopsy^{50,53}

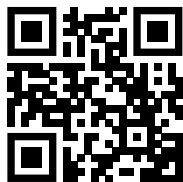
AR, androgen receptor; ASCO, American Society of Clinical Oncology; ASTRO, American Society for Radiation Oncology; *ATM*, ataxia-telangiectasia mutated; AUA, American Urological Association; BPH, benign prostatic hyperplasia; *BRCA*, breast cancer gene; *CDK12*, cyclin-dependent kinase 12; cfDNA, cell-free DNA; *CHEK2*, checkpoint kinase 2; CRPC, castration-resistant prostate cancer; CTC, circulating tumor cell; ctDNA/RNA, circulating tumor DNA/RNA; DDR, DNA damage response; dMMR, deficient mismatch repair; DNA, deoxyribonucleic acid; EVP, extracellular vesicles and particles; *FANCA*, FA complement group A; GPS, genomic prostate score; *HOXB13*, homeobox B13; HRD, homologous recombination deficiency; HRR, homologous recombination repair; mCRPC, metastatic castration-resistance prostate cancer; mCSPC, metastatic castration-sensitive prostate cancer; miRNA, microRNA; *MLH1*, MuTL homolog 1; MMR, mismatch repair; mRNA, messenger RNA; *MSH2/6*, MutS homolog2/6; MSI, microsatellite instability; MSI-H, microsatellite instability-high; *NBN*, nibrin gene; NCCN, National Comprehensive Cancer Network®; *PALB2*, partner and localizer of *BRCA2*; PARPi, poly-ADP ribose polymerase inhibitor; *PMS2*, postmeiotic segregation increased 2; PSA, prostate-specific antigen; PSMA, prostate-specific membrane antigen; SUO, Society of Urologic Oncology; TMB, tumor mutational burden.

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