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An appraisal of genetic testing for prostate cancer susceptibility

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Most criteria for genetic testing for prostate cancer susceptibility require a prior diagnosis of prostate cancer, in particular cases with metastatic disease are selected. Advances in the field are expected to improve outcomes through tailored treatments for men with advanced prostate cancer with germline pathogenic variants, although these are not currently offered in the curative setting. A better understanding of the value of genetic testing for prostate cancer susceptibility in screening, for early detection and prevention is necessary. We review and summarize the literature describing germline pathogenic variants in genes associated with increased prostate cancer risk and aggressivity. Important questions include: what is our ability to screen for and prevent prostate cancer in a man with a germline pathogenic variant and how does knowledge of a germline pathogenic variant influence treatment of men with nonmetastatic disease, with hormone-resistant disease and with metastatic disease? The frequency of germline pathogenic variants in prostate cancer is well described, according to personal and family history of cancer and by stage and grade of disease. The role of these genes in aggressive prostate cancer is also discussed. It is timely to consider whether or not genetic testing should be offered to all men with prostate cancer. The goals of testing are to facilitate screening for early cancers in unaffected high-risk men and to prevent advanced disease in men with cancer.

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INTRODUCTION

Increasing numbers of men are encouraged to undergo germline genetic testing for prostate cancer. This is due in large part to better understanding of inherited prostate cancer risk and progression and an expansion of genetic services for cancer patients and for those at high risk of developing cancer.

Genetic testing for cancer has emerged over the last decade into panel-based testing whereby numerous genes are examined simultaneously for deleterious pathogenic/likely pathogenic variants or mutations. For example, most hereditary cancer panels include genes for breast, ovarian and colon cancer – and includes the five main genes for prostate cancer discussed here (BRCA1, BRCA2, CHEK2, ATM, PALB2). Most cancer susceptibility genes are associated with a predisposition to develop tumors at multiple sites. As a result, selecting individual genes for analysis is no longer a primary concern for the physician because patients are offered a 'comprehensive cancer panel' regardless of the indication.

Prostate cancer is the primary form of hereditary cancer which is specific to males. The main risk factors for prostate cancer are age, family history and race. Men of African descent are at greater risk for prostate cancer than Caucasian men, while men of Asian and Hispanic descent are at lower risk¹. Men of African ancestry have an earlier age at diagnosis of prostate cancer, a higher incidence and increased prostate-specific mortality when compared with Caucasian men^{1,2}. A genetic basis for the racial difference has not been established.

A man with a first-degree relative with prostate cancer has approximately twice the risk of a man who does not; this risk is

higher if the relative is diagnosed under age 60^{3–5}. This increased familial risk of prostate cancer is attributable in part to the additive effects of multiple low-risk germline variants (SNPs) and higher risk germline pathogenic variants in cancer susceptibility genes⁶. Polygenic risk scores (PRS) generated through combining a number of low-risk of prostate cancer susceptibility variants may provide discrimination in risk both for carriers of high-risk germline variants and for men in the general population^{7–9}.

Our primary focus here is on moderate-to-high-risk hereditary cancer susceptibility genes. We discuss screening and prevention in men at high risk and the over-representation of these genes in aggressive prostate cancer. The individual genes are discussed below. Finally, we consider whether all men with prostate cancer should be offered genetic testing.

Testing prostate cancer patients

There are several circumstances under which a man might benefit from genetic testing. For unaffected men, the information may help to specify his lifetime prostate cancer risk and aid in prevention and screening. For affected men, the information may be used to personalize cancer treatment. It is rare that an unaffected man will be the first to receive genetic testing in a family. For unaffected men, the most common scenario is that he is approached to initiate testing after a woman in his family is found to carry a *BRCA1* or *BRCA2* mutation (or a mutation in another gene). Many men describe being tested 'for the sake of their daughters' rather than for their own health¹⁰. For a prostate cancer patient, the opportunity to be tested will depend on his family history of cancer and his clinical presentation. To our

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knowledge, there is no guideline that suggests that all men with prostate cancer should be eligible for testing. Factors that determine eligibility are based on the probability of a mutation being detected (i.e. the prevalence) and include the presence of metastatic disease, high-grade disease, intraductal/cribriform histology, a family history of prostate cancer, a family history of breast or ovarian cancer, or a known mutation in the family¹¹. If we consider the *utility* of testing as the primary goal of germline genetic testing we should address the following five questions:

- 1. What is our ability to prevent prostate cancer in men with a high risk of cancer?
- 2. What is the value of screening for prostate cancer among men at high risk?
- 3. Does a germline mutation influence prognosis?
- 4. How does the knowledge of a germline mutation influence treatment of men with nonmetastatic cancer?
- 5. How does the knowledge of a germline mutation influence treatment of men with metastatic cancer or with hormoneresistant disease?

For women, genetic testing for *BRCA1* and *BRCA2* mutations is now widely endorsed and accepted. Its value has been demonstrated through surgical prevention of breast cancer with prophylactic mastectomy, surgical prevention of ovarian cancer with bilateral salpingo-oophorectomy, the improved sensitivity of MRI over mammography screening, the benefit of MRI screening on breast cancer mortality and the benefit of specific chemotherapies, such as platinum-based neoadjuvant therapy for women with breast cancer and poly-ADP ribose polymerase (PARP) inhibitors (i.e. Olaparib) for treatment of women with ovarian cancer. It is now recommended that all women with epithelial ovarian cancer undergo genetic testing for susceptibility genes¹¹. The clinical benefits of genetic testing for men for susceptibility genes is emerging but has not yet been established.

What is the ability to offer prevention to unaffected men at high risk of prostate cancer?

Currently, there is no approved drug for the prevention of prostate cancer. Numerous randomized controlled trials have been performed on potential agents, including 5-alpha reductase inhibitors (which act to block the action of androgens in the prostate and include Dutasteride and Finasteride), medications, which manipulate the hormonal axis (e.g. Selective estrogen receptor modulator Toremifene), and non-steroidal anti-inflammatory drugs (e.g. Refocoxib). Several nutritional supplements have also been evaluated (e.g. selenium, vitamin E and soy products). Results of these trials have been disappointing, with one trial stopped because of cardiovascular toxicity (e.g Refocixib)¹², and another showed an increase in prostate cancer risk (e.g. Vitamin E)¹³ and two trials, which showed a decline in overall prostate cancer risk in the treatment arm, but an increased risk of high-grade disease (PCPT and REDUCE trials)^{14,15}. Given the lack of compelling evidence, none of these agents is currently recommended for prostate cancer prevention among individuals at average risk or high-risk. There are no trials underway specifically for chemoprevention in high-risk men, including carriers of highrisk pathogenic variants.

What is the value of screening for prostate cancer among men at high risk?

The value of screening for prostate cancer in the general population is controversial. Three large randomized controlled trials of PSA screening and mortality produced mixed results. The European ERSPC study and the Gotenberg study found a 20–30% and 52% reduction in prostate cancer mortality respectively^{16,17}, whereas the US PLCO trial showed no mortality difference between the treatment and control arms¹⁸. The US Preventative

Services Task Force (USPSTF) recommends a discussion of the risks and benefits of PSA screening in men aged between 55 and 69 (and against screening for men over 70)¹⁹. The NCCN recommends that men with a germline mutation that increases the risk for prostate cancer consider annual PSA screening from age 40 (NCCN Guidelines V 2.2021 Prostate Cancer Early Detection)²⁰. The NCCN also recommends annual screening in men with a family history and men of African descent from age 40, but notes that the evidence does not show a mortality benefit associated with earlier screening^{2,20}. The American Cancer Society recommends starting a discussion about screening at the age of 40 for men at high risk, including those with more than one first-degree relative who had prostate cancer at an early age (a very small group)²¹.

The IMPACT study targets PSA screening specifically for male *BRCA1* and *BRCA2* mutation carriers (and controls) between ages 40 and 69. A biopsy is offered for those with a PSA > 3.0 ng/ml. A preliminary analysis demonstrated that men who carry a *BRCA2* mutation have a higher incidence of prostate cancer, a younger age at diagnosis and more clinically significant tumours than noncarrier controls. The IMPACT study group concluded that men who carry a *BRCA2* mutation should be offered annual PSA screening, but further follow-up is needed for *BRCA1* carriers²².

To some extent, the impact of PSA screening is limited by the sensitivity of the test and the possibility of diagnosing indolent disease. Alternative screening modalities include multiparametric magnetic resonance imaging (MRI). Segal et al. screened 188 BRCA1 and BRCA2 carriers with no previous prostate biopsy with PSA and MRI²³. Sixteen prostate cancers (8.5%) were diagnosed at the initial round of screening. MRI screening only missed one cancer (6%), while the PSA test alone missed five (31%). Further follow-up is needed to determine if there is an effect on mortality. The utility of screening in carriers will be determined when a large number of men who are identified to be at high risk are screened and followed.

Does a germline mutation influence prognosis?

It is a central question whether or not the presence of a mutation predisposes men to more aggressive disease. There are several indicators of prostate cancer aggressivity that can be used in the evaluation. These include:

- 1. Gleason score (Grade group) at presentation.
- 2. The presence of metastatic disease at presentation.
- 3. Time to biochemical recurrence.
- 4. Treatment response.
- 5. Prostate-specific survival.
- 6. Time to death.
- Mutation prevalence among men with metastatic disease versus localized disease.

It is important to ask if the prognosis of a man with an inherited pathogenic variant is influenced by the presence of the variant, independent of stage, grade and PSA level. It should also be noted that a pathogenic variant may not increase the risk of prostate cancer overall but may influence the probability of high-grade disease and prognosis to metastatic disease²⁴. Based on limited studies to date, it appears that men with *BRCA2*, *ATM* and *PALB2* pathogenic variants face increased risks of progression and death that cannot be accounted for by traditional clinical parameters²⁵. Further work is needed in this area.

How does the knowledge of a germline mutation influence treatment of men with nonmetastatic cancer?

Prostate cancer accounts for 10% percent of all cancer deaths in men and its behavior varies from indolent to aggressive disease²⁶. In a study of prostate cancer incidence and survival in the United States, localized, regional and distant-stage prostate cancer accounted for 77%, 11% and 5% of cases, respectively (cases

with unknown stage accounted for 7%)²⁷. Ten-year survival for localized prostate cancer was 100%, but was 96.1% and 78.1% for regional and unknown stage respectively. The five and 10-year survival for distant-stage prostate cancer was much lower at 30.7% and 18.5%, respectively. Current practice is to offer treatment according to recurrence risk, based on a combination of Grade group, PSA level and extent of disease. Germline mutation status is most often not considered in clinical practice guidelines at present but represents an opportunity to personalize treatment. For some genes (BRCA2, PALB2 and ATM), germline mutations have been associated with aggressive disease and relatively poor outcomes²⁵. Inherited variants in these DNA-repair genes are likely to play an increasingly important role in deciding the choice of treatment including the option of active surveillance^{28–30}. Other genes have been less well studied or have been shown not to influence outcome

Traditional treatments for localized prostate cancer include surgery (radical prostatectomy) and radiotherapy (external beam or brachytherapy). In general, surgery is favoured in younger men (<55 years old), radiotherapy is favoured in men over age 70 (in men between age 55 and 70 the decision is based on patient preference and tumour characteristics).

There are few studies that compare outcomes after surgery and radiotherapy for men with and without germline mutations. Castro et al. examined the tumor features and outcomes of 2,019 patients with prostate cancer, which included 18 BRCA1 and 61 BRCA2 carriers in a retrospective study²⁹. They found that BRCA2 mutation carriers were more likely to be diagnosed with high-risk disease (Gleason 8-10), have advanced clinical stage disease (T3/ 4), and/or have involvement of local lymph nodes or metastatic disease at diagnosis. Five-year cancer-specific survival was significantly worse for BRCA carriers than noncarriers (82% vs. 96%; $p = 9 \times 10^{-8}$). In a subsequent publication, Castro et al. compared survival among 67 BRCA carriers and 1235 noncarriers who received either radiotherapy or surgery for localized or locally advanced prostate cancer³¹. The groups were not directly comparable because there were a greater number of high-risk patients in the radiation therapy group than the radical prostatectomy group (69% vs. 34%). The 10- year metastasesfree survival (MFS) was significantly lower for carriers than noncarriers in both the radical prostatectomy group (63% vs. 91%) and in the radiation therapy group (39% vs. 80%). Prostatespecific survival of carriers vs. noncarriers who underwent radical prostatectomy at 10 years was 79% vs 95% (log rank test p = 0.6). There was a significant difference in outcomes between carriers and noncarriers who had radiation therapy, with 10-year survival rates of 47% vs 81% respectively (log rank test p < 0.001). It is important that we compare the relative benefits of radical prostatectomy and radiation therapy directly among carriers with similar risk profiles in future studies.

ATM and the risks of radiotherapy

A study of prostate cancer patients with heterozygous *ATM* mutations found an increased risk of late complications of external beam radiotherapy^{32,33}. Others have demonstrated that there is potential for increased therapeutic efficacy of radiotherapy among *ATM* carriers, but care must be taken to minimize radiation dose to prevent toxicity and second malignancies³⁴. At this time, the NCCN guidelines state that carrying a single *ATM* mutation should not lead to a recommendation to avoid radiation therapy¹¹. There is little evidence regarding late toxicity and second malignancy of radiotherapy for carriers of other germline mutations.

Active surveillance for low-risk disease

A historical overview of treatment outcomes of men with screendetected, low-grade prostate cancer has led to the adoption of active surveillance for some low-risk prostate cancer patients. The choice of active surveillance relies on a combination of factors, including the PSA level, Grade group, clinical stage, and biopsy results (number of cores involved, percentage of involvement of dominant core, etc.). Active surveillance consists of PSA test every 3–6 months, digital rectal exam once per year, a confirmatory biopsy within 6–12 months, then serial biopsy a minimum of every 3–5 years thereafter with or without mpMRl 35 . In the event of a rising PSA or grade progression (e.g., Gleason 6 to \geq 7) active treatment is offered. Between 36% to 73% of patients will transition to treatment over 10 years but the development of metastatic disease is rare 36,37 .

There is little data on the safety or efficacy of active surveillance for men diagnosed with a high-risk germline mutation. An important question is "Does the presence of a mutation predict for a worse outcome (in particular metastatic disease) after adjusting for the traditional clinical parameters (grade, PSA, screening history, etc.)? If so, active surveillance may not be appropriate for men with mutations. Carter et al. followed 1211 men with prostate cancer (from two cohorts) under active surveillance, including 26 with a mutation in BRCA1, BRCA2 or ATM^{28} . Eleven of 26 men with a mutation in one of the three genes experienced grade progression compared to 278 of 1185 noncarriers (adjusted hazard ratio = 2.0; (p = 0.05). This association was significant for BRCA2 carriers (adjusted HR = 2.7; p = 0.01). The authors suggest that the mutation status should be considered in the decision to enter active surveillance.

How does the knowledge of a germline mutation influence treatment of a man with metastatic cancer or with hormoneresistant disease?

After surgery or radiotherapy, patients may experience prostate cancer recurrence. Biochemical recurrence refers to a rising PSA level. Approximately 30-40% of patients will develop biochemical recurrence after radical treatment (e.g. surgery or radiotherapy)^{38–41}. Biochemical recurrence is an early indicator of metastatic disease. The majority of men who experience a biochemical recurrence after surgery will develop metastatic prostate cancer within 10 years⁴². Common treatments for biochemical recurrence include radiotherapy and/or androgen deprivation (after surgery) and androgen deprivation therapy or surgery (after radiotherapy). Androgen deprivation therapy is generally not considered until a biochemical recurrence is documented. As previously discussed, individuals with a germline mutation in several genes (BRCA2, ATM, PALB2) are at increased risk for biochemical recurrence and prostate cancerspecific mortality. These men may potentially be candidates for earlier initiation of androgen deprivation therapy. Other options include cisplatin-based chemotherapy or early use of PARP inhibitors. While recent evidence has shown that these agents can be effective for women with breast cancer and a BRCA mutation⁴³ and men with metastatic prostate cancer, more evidence is needed for high-risk men with localized prostate cancer.

Approximately 5–10% of men with prostate cancer present with metastatic disease¹. The 5-year survival for men with metastatic prostate cancer is approximately 30%⁴⁴, but recent advances in treatment are extending survival. The conventional treatment for metastatic prostate cancer is androgen axis inhibition, chemotherapy or radioligand treatments. Response to androgen deprivation therapy has led to the distinction between castrate-sensitive metastatic prostate cancer (responds to androgen blockage) and castrate resistant (continued progression of disease). Ten to twenty percent of patients with metastatic prostate cancer become castrate-resistant within 5-years⁴⁴. Carriers of germline mutations are at greater risk of progressing from castrate-sensitive to castrate-resistant metastatic disease than noncarriers^{45–47}.

PROREPAIR-B is an ongoing prospective study to evaluate the outcomes of patients with metastatic castrate-resistant prostate cancer with and without germline ATM/BRCA1/BRCA2/PALB2



mutations⁴⁶. They have demonstrated that patients with mutations in *BRCA2* have worse outcomes.

There is increasing interest in cisplatin-based chemotherapy and PARP inhibitors for metastatic prostate cancer⁴⁸. Two PARP inhibitors (Olaparib and Rucaparib) are approved by the US Food and Drug Administration for men with metastatic castrate resistant prostate cancer (mCRPC) and a mutation in one of 15 homologous recombination repair genes (*BRCA1*, *BRCA2*, *ATM*, *BRIP1*, *BARD1*, *CDK12*, *CHEK1*, *CHEK2*, *FANCL*, *PALB2*, *PPP2R2A*, *RAD51B*, *RAD51C*, *RAD51D*, and *RAD54L*)^{49,50}.

Men with a mutation in one of the MMR genes or with microsatellite instability-high/MMR-deficient tumours may benefit from anti-programmed cell death protein 1 (PD-1) antibody therapy⁵¹. Pembrolizumab was approved by the US Food and Drug Administration for the treatment of microsatellite instability-high (MSI-H) or mismatch repair–deficient (dMMR) solid tumors, including prostate cancer.

Summary

There is currently no evidence for chemoprevention for prostate cancer.

-There is emerging evidence that screening unaffected men with a pathogenic mutations may have clinical utility and further research this area is required.

-There is emerging evidence that the presence of a mutation in *BRCA2, ATM* or *PALB2* puts the patient at increased risk for biochemical recurrence and metastatic disease.

-It is anticipated that knowing gene carrier status will allow more personalized discussions about prognosis, immediate management and management of metastatic and hormone resistant disease.

Historical perspective

BRCA2 was one of the first genes implicated in hereditary prostate cancer and is responsible for a significant proportion of hereditary prostate cancer cases around the world. Other genes implicated in the development of prostate cancer include those involved in Homologous Recombination DNA repair: *BRCA1* in 1994⁵², *CHEK2* in 2003^{53,54}, *ATM* in 2004⁵⁵, and *PALB2* in 2008⁵⁶. The DNA mismatch repair genes that are associated with Lynch Syndrome

(MLH1, MSH2, MSH6 and PMS2) have also been implicated⁵⁷. Pathogenic variants in additional genes have been found at lower frequency including HOXB13 and RAD51D⁵⁷. These genes are associated with a range of cancer types. The exception is HOXB13 which, to date has only been associated with prostate cancer.

Frequency of germline mutations in men with prostate cancer

While germline mutations in genes that predisposes to prostate cancer have relatively low mutation prevalence individually, together they account for a significant proportion of prostate cancer cases. The prevalence depends on the population studied (e.g., family history, clinical presentation, means of selection) and the number of genes surveyed. In a cohort of 53,105 men with no diagnosis of cancer (general population from the Exome Aggregation Consortium), the frequency of DNA-repair gene mutations was 2.7%. Among 499 men with localized prostate cancer in the Cancer Genome Atlas Research Network the frequency of germline mutations was 4.6%. In a recent study of localized prostate cancer, 4% of men with European ancestry carried a DNA repair gene mutation compared 2.8% of unaffected controls⁵⁸. Of 351 men with African ancestry and localized prostate cancer, 1.4% had a mutation. It is important to note that this study excluded men who later progressed to metastatic disease.

The prevalence rates are higher than this in men with metastatic disease and/or a family history of prostate cancer. In a study of 692 men with metastatic prostate cancer, 82 men (11.8%) were found to carry a mutation in one of 16 cancer susceptibility genes^{56,59,60} (See Table 1 for mutation frequency by gene). The frequencies of mutations in *ATM* and *BRCA1* and *BRCA2* were compared in 313 men who died of prostate cancer and 486 men with localized prostate cancer⁶¹. The mutation rate was much higher among those with lethal prostate cancer (6.1% vs 1.4%, p = 0.0007). For those who died below age 60, 10% carried a mutation. Of those who died within five years of diagnosis 12.3% carried a mutation.

In the UK Genetic Prostate Cancer Study (UKGPCS), 191 men with prostate cancer with two or more relatives with prostate cancer were tested for mutations in 22 tumour suppressor genes. Fourteen men (7.3%) were found to carry a mutation in one of eight genes⁵⁹. Nicolosi et al. reported on a large series of 3607

Table 1. Germline Mutation frequency in Men with Prostate Cancer and Population Controls.								
	Exome Aggregation Cohort General Population ī = 53,105 (%)	Lee et al. ⁵⁸ Localized Prostate Cancer *ī = 1174 European population (%)	TGCA cohort Localized Prostate Cancer $\bar{\imath}=499~(\%)$	Leongamornlert et al. ⁵⁹ Familial Prostate Cancer $\bar{i} = 191$ (%)	Prichard et al. ⁵⁶ Metastatic Prostate Cancer $\bar{i} = 692$ (%)	Castro et al. Prorepair-B ⁴⁶ Metastatic CR Prostate Cancer $\bar{\imath}=419~(\%)$	Nicolosi et al. ⁶² Invitae Prostate Cancer** ī= 3607 (%)	
All Genes: % +ve	2.7%	4.0%	4.6%	7.3%	11.8%	16.2%	17.2%	
BRCA1	0.22	0.77	0.60	0.52	0.87	1.0	1.25	
BRCA2	0.29	1.0	0.20	2.10	5.35	6.2	4.74	
ATM	0.25	0.51	1.0	1.04	1.59	1.9	2.03	
PALB2	0.12	0.17	0.40	0.52	0.43	_	0.56	
CHEK2	0.61	0.34	0.40	1.04	1.87	0.24	2.88	
MLH1	0.02	0.0	0	_	0	_	0.06	
MSH2	0.04	0.17	0.20	-	0.14	0.24	0.69	
PMS2	0.11	0.09	0.20	0.52	0.29	_	0.54	
MSH6	0.08	0.09	0.20	_	0.14	_	0.45	
HOXB13	_	_	_	-	_	_	1.12	
RAD51D	0.08	0.0	0.20	-	0.43	_	0.15	

^{*}Men who subsequently developed metastatic disease were excluded

^{**}Referral-based genetic testing for personal history of prostate cancer and includes men with additional risk factors for hereditary cancer.

men with prostate cancer with varying family history and ethnicity tested by Invitae⁶². The overall mutation detection rate reported was considerably higher than for previous studies (17.2%), but these patients were selected and referred for testing. Many had a family history of cancer. The highest mutation frequencies were found in men with prostate cancer and a family history of ovarian cancer (22.8%), a family history of pancreatic cancer (19.4%) and Ashkenazi Jewish ancestry (22.2%).

In the next section we consider prostate cancer susceptibility genes individually.

BRCA1

BRCA1 predisposes women to breast, ovarian, fallopian tube and peritoneal cancer $^{52.63}$. An early study of prostate cancer in *BRCA1* carriers found an of elevated risk for prostate cancer among those under age 65 with a relative risk of 1.82 (95% CI: 1.01–3.29, $p = 0.05)^{64}$. This was confirmed in a second series of 913 men with clinically evident prostate cancer (RR = 3.75 (95% CI: 1.02–9.6)) translating to approximately 8.6% risk to age 65 65 . A recent prospective study of prostate cancer risk found men who carry a *BRCA1* mutation have an absolute risk for prostate cancer of 21% to age 75 and 29% to age 85 66 . Among men with metastatic prostate cancer, approximately 1% carry a *BRCA1* mutation 46,56 . It is not clear that *BRCA1* predisposes to a more aggressive form of prostate cancer.

BRCA2

BRCA2 is the best-established prostate cancer susceptibility gene. *BRCA2* also predisposes women to breast and ovarian cancer (including fallopian tube and peritoneal cancer)⁶⁷. The risks for pancreatic cancer, melanoma, and for men, prostate and breast cancer, are also higher than general population risk^{64,68}. Kote-Jarai et al. estimated that *BRCA2* confers an approximately 8.6 fold increase in prostate cancer risk by age 65, with a 15% absolute risk by age 65⁶⁹. In the prospective study by Nyberg et al., *BRCA2* carriers had an absolute risk of prostate cancer of 27% to age 75 and the risk estimate reached 60% by age 85. To some extent these risk estimates depend on the intensity of surveillance.

Among men with *BRCA* mutations, the family history impacts upon risk. For *BRCA1* carriers, risk for prostate cancer was higher for those with a positive family history of prostate cancer (SIR = 3.2) than for those without (SIR = 2.3). For *BRCA2* carriers the SIR was 7.3 for those a positive family history and was 3.9 for those without⁶⁶.

The *BRCA2* mutation frequency is similar for men with earlyonset prostate cancer (<age 55) (2.3%) and for men with familial prostate cancer (3 or more cases of prostate cancer) (2.1%)⁵⁹. The frequency of *BRCA2* mutations was 4.5% in a large series of men with prostate cancer referred for testing with varying family histories, ethnicities and stages of disease⁶².

Multiple studies have found poor survival for *BRCA2* carriers compared with noncarriers ^{28,29,61,70–72} *BRCA2* carriers have a stronger association with Gleason score ≥ 7 cancer than with Gleason score ≤ 6, cancer and a higher risk of death from prostate cancer⁶⁶. Castro et al. examined 2019 men with prostate cancer (18 BRCA1 carriers, 61 BRCA2 carriers and 1940 noncarriers) with the aims of determining the independent prognostic value of BRCA1 and BRCA2 mutations on prostate cancer characteristics and survival²⁹. They found that carriers were more likely to have poorly differentiated cancer (Gleason score ≥ 8) (35% vs 15%; *p* = 0.00003), advanced stage (T3-T4) (37% vs 28%; p = 0.003) and metastatic spread (M1:18% vs. 9%; p = 0.005). Median causespecific overall survival was shorter in carriers than noncarriers (8.6 vs. 15.7 years; $p = 7 \times 10^{-8}$). A trend toward better median causespecific survival was observed in BRCA1 carriers compared with BRCA2 but it was not significant (10.5 and 8.6 years, p = 0.37). The 5-year metastases-free survival was also lower in carriers than in noncarriers (77% vs. 93%; p=0.0001). In the Prorepair-B study of men with Spanish ancestry and metastatic castrate-resistant prostate cancer 3.3% carried a *BRCA2* mutation, slightly lower than the 5.4% among the 692 men with metastatic prostate cancer in the study by Pritchard et al^{46,56}. *BRCA2* accounted for 44% of all pathogenic variants detected among the men with metastatic cancer in the Pritchard study.

ATM

The ATM gene (Ataxia Telangiectasia Mutated) is required for cell response to DNA damage and for genome stability. Women who carry an ATM mutation have a 20-30% lifetime risk to develop breast cancer, and carriers also have a possible but as yet undefined risk for other cancers, such as prostate, pancreatic and colon cancer^{73–75}. The frequency of ATM mutations ranges from approximately 0.5% in the general population (Exome Aggregation Consortium), to 1% in men with localized prostate cancer to 1.6% in men with metastatic prostate cancer^{56,60}. In addition, ATM mutations have been associated with aggressive disease, and with metastatic disease^{28,56}. Darst et al. performed a case-control study of aggressive vs. nonaggressive prostate cancers in men of European ancestry⁷⁶. They found ATM was associated with aggressive disease, with 1.6% of aggressive and 0.8% of nonaggressive cases carrying a pathogenic or likely pathogenic variant (OR = 1.88, 95% CI: 1.10–3.22, p = 0.02). In Poland, 50% of cancers diagnosed among ATM carriers were Gleason score 8-10, compared to 22.7% in noncarriers $(p = 0.03)^{77}$.

CHEK2

CHEK2, also called cell cycle checkpoint kinase 2 is a cell cycle check point regulator. The variant c.1100delC accounts for the majority of pathogenic variants in this gene⁷⁸. CHEK2 mutations confer an increased risk for breast (in both males and females), thyroid, colon, and prostate cancer⁷⁸⁻⁸². Hale et al. pooled five studies and examined prostate cancer risk associated with the 1100delC variant. They found the odds ratios were 1.98 (95% CI: 1.23-3.18) for unselected cases and 3.39 (95% CI: 1.78-6.47) for familial cases⁸³. Cybulski et al. examined prostate cancer risk associated with the truncating variant 1100delC in addition to two other truncating variants (IVS2 + 1 G > A (c.444 + 1 G.A), del5395) and one missense variant (p.1157T). The p.1157T missense variant was associated with an odds ratio of 1.8 (95% CI: 1.5-2.2), p < 0.0001, as compared with the truncating variants combined (OR = 2.1 (95% CI: 1.4–3.0), p = 0.0001. For the 1100delC variant alone the odds ratio was 3.2 (95% CI: 1.4–7.5), $p = 0.009^{84}$.

CHEK2 variants are found in fewer than one percent of the general population (0.61% in the Exome Aggregation Consortium) and in two percent of men with metastatic prostate cancer, giving a relative risk for metastatic prostate cancer of 3.1 (95% Cl: 1.5–5.6), $p=0.0002^{56}$. CHEK2 mutations were among the most commonly found variants in men with metastatic cancer (12% of 84 variants detected among 692 men with metastatic disease)⁵⁶.

PALB2

PALB2 is a known breast cancer susceptibility gene with risk of breast cancer estimated to be approximately 14% to age 50 and 53% to age 80⁸⁵. It is also associated with a moderate risk of ovarian cancer (approximately 5% to age 80)⁸⁶. Risks may also be increased for pancreatic cancer (2–3%) and male breast cancer (1%). An association with gastric cancer has also been reported^{87,88}.

Studies of families with mutations in *PALB2* have not demonstrated an increased risk for prostate cancer, but *PALB2* variants have been associated with aggressive prostate cancer and with metastatic prostate cancer^{24,25,86}. Pritchard et al. found an increased frequency of *PALB2* mutations in the series of 692



men with metastatic prostate cancer when compared with a large population of people unaffected with cancer (RR 3.5, 95% Cl: 0.7-10.3, $p=0.05)^{56}$. Wokolorczyk et al tested 5472 unselected Polish men with prostate cancer and 8016 Polish controls for two founder *PALB2* variants. They found that *PALB2* carriers were not more common in prostate cancer patients overall, but were common in those diagnosed with aggressive prostate cancers of high grade (Gleason score 8–10) than noncarriers (64.3% vs. 18.1%, p < 0.0001). The five-year survival for *PALB2* carriers was 42% compared with 72% for noncarriers (p=0.006)²⁴.

MMR genes

The mismatch repair (MMR) genes (*MLH1*, *MSH2*, *MSH6*, *PMS2*) play a key role in genomic stability and are associated with significantly increased risks varying by gene for colon cancer (15–46%), endometrial cancers (43 to 57%) and ovarian cancer (10–17%)^{89–91}. Mutations in MMR genes are associated with additional cancers such as urothelial, gastric, small bowel, pancreatic, hepatobiliary, brain, and sebaceous carcinomas^{89,92}. Germline mutations in these genes are collectively known to cause Lynch syndrome.

In a study of Lynch Syndrome families, the cumulative lifetime risk (to age 80) of prostate cancer was estimated to be 30% in MMR mutation carriers^{57,93}. MSH2 was the most strongly associated with prostate cancer of the MMR genes, with a risk of 23.8% risk to age 75 in the Prospective Lynch Syndrome Database^{94,95}. Similarly, Barrow et al found a 10-fold increased risk of prostate cancer associated with MSH2 in study of male Lynch syndrome mutation carriers and their first-degree male relatives who were part of the Manchester Regional Lynch Syndrome database (RR 10.41; 95% CI: 2.80-26.65)⁸⁹. In the series of 3607 men with prostate cancer, 2% carried a mutation in one of four DNA mismatch repair genes (MLH1, MSH2, MSH6 and PMS2). MSH2 accounted for the greatest proportion of these mutations followed by PMS2 and MSH6 with only 2 of 58 pathogenic variants in these 4 genes identified in MLH162. In studies to date, the age at onset and aggressiveness associated with MMR-related prostate cancers do not appear to be different than observed in sporadic cases⁹⁶.

RAD51D

RAD51D is involved in homologous recombination repair and is primarily associated with hereditary ovarian cancer, and to a lesser extent, breast cancer^{78,79}. Germline *RAD51D* mutations are relatively rare in men with prostate cancer, (3/692 = 0.43%). Although the mutation frequency is low, the relative risk of metastatic disease was 5.7 (95% Cl: 1.2–16.7, p = 0.02)⁵⁶. Further studies are needed to better define the role of *RAD51D* in the development of aggressive and metastatic prostate cancer.

HOXB13

HOXB13 is a homeobox transcription factor gene and a single rare variant G84E confers a significantly increased risk for prostate cancer^{97,98} The odds ratios for development of associated with this variant range from 2.9 (95% CI: 1.9–4.6) in the UK Genetic Prostate Cancer Study (UKGPCS) to 8.8 (95% CI: 4.9–15.7) among men with a family history in a Finnish study by Laitinen et al.^{99,100}.

Nyberg et al estimated the absolute risk of prostate cancer in a large UK-based kin-cohort study to be 60% to age 85 among men with the G84E variant with no family history and 98% for men with the variant and two relatives with prostate cancer at young ages, compared to 15% for noncarriers¹⁰¹. In Poland the G84E allele was associated with a five-fold increase in the risk of unselected prostate cancer¹⁰².

In the Finnish study, 8.4% (16 of 190) men with prostate cancer and a positive family history were positive for the *HOXB13* variant, higher than observed in other cohorts of familial prostate cancer but similar to that in Sweden, suggesting a founder effect for the variant ¹⁰⁰. The International Consortium for Prostate Cancer Genetics (ICPCG) genotyped a large sample of prostate cancer families and found the variant was present in approximately 5% of prostate cancer families of European descent ⁹⁹. The *HOXB13* variant has been found to be associated with younger age at diagnosis and higher PSA values at diagnosis ^{100,103,104}. There is little data to date on the presence of *HOXB13* mutations in non-European populations.

Prostate cancer: who to test

Current guidelines for germline genetic testing tend to be complex and focus on men with prostate cancer. The NCCN Guidelines (V 2.2021 for Prostate Cancer and the NCCN Guidelines V2.2021 for High-Risk Assessment for Breast, Ovarian and Pancreatic Cancer) recommend germline genetic testing for patients with prostate cancer who satisfy additional criteria such as Ashkenazi Jewish ancestry, family history of cancer and/or based on characteristics of the prostate cancer, such as grade, stage, histology and the presence of metastatic disease. A man with a likelihood 2.5–5% of carrying a pathogenic BRCA1/2 variant (in prior probability models such as Tyrer Cuzick, BRCAPro, CanRisk) is also eligible for testing 11. The likelihood of detecting a germline pathogenic variant in men with prostate cancer are presented by gene and by study in Table 1. All genes listed in Table 1 are considered to be actionable hereditary cancer predisposition genes, but not all genes pose similar risk for prostate cancer or for more aggressive disease. The genes which are believed to have the greatest impact in terms of risk are highlighted in Table 2 (BRCA1, BRCA2, ATM and PALB2). HOXB13 is not included in many of these studies but is associated with prostate cancer in European populations.

The prevalence of mutations according to risk group is also presented in Table 2. The prevalence is highest for men with a female relative with ovarian cancer and for men with metastatic prostate cancer. The majority of mutations are in *BRCA1* and *BRCA2*, although the number of genes tested varies from study to study.

The 2019 Philadelphia Prostate Cancer Consensus Conference on the implementation of Germline Genetic Testing for Prostate Cancer identified challenges in the implementation of testing in three main areas: determining which genes should be included in panels, variability in guidelines regarding who should have genetic testing and the availability of genetic services³⁰.

Population testing of all men will identify the greatest number of unaffected men at risk for prostate cancer and allow for targeted screening and possibly prevention; however, this is not

Table 2. The prevalence of mutations by various at risk groups.						
Group	Prevalence of LP/P variant	Prevalence of LP/P variant in: BRCA1/BRCA2, ATM & PALB2				
Men diagnosed with localized prostate cancer	4.0% ⁵⁸ , 4.6% ⁶⁰	2.5% ⁵⁸ , 2.2% ⁶⁰				
Men diagnosed with metastatic prostate cancer	11.8% ⁵⁶	8.2% ⁵⁶				
Affected men with 2 relatives with prostate cancer	7.3% ⁵⁹	4.2% ⁵⁹				
Men with first degree relative with epithelial ovarian cancer	10% (for BRCA only ¹⁰⁹)	10% (for BRCA only ¹⁰⁹)				



currently feasible. A focused approach is required. Genetic testing of patients allows an opportunity to identify those who are more likely to progress to advanced or metastatic disease and provides and opportunity for intervention. Testing for underlying hereditary predisposition has become standard of care for patients with other cancers, such as colon cancer and epithelial ovarian cancer^{11,105}. Goals for genetic testing for prostate cancer should include prevention of progression of the disease in men at highest risk and improvement in treatment of hereditary prostate cancer. Currently testing is not recommended for all men with prostate cancer, rather is based on the probability that a mutation is likely to be found. In the event that personalised mutation-specific treatments for prostate cancer are developed, there will be an impetus to expand testing to more prostate cancer patients. This will also allow the identification of other family members at risk for a range of cancers (cascade testing).

Models of genetic testing. Prostate cancer is very common in Europe and North America and if genetic testing is to be expanded to more men with prostate cancer, it is important to consider how to do it effectively and efficiently. The current model of genetic services includes pre- and post-testing genetic counselling and is time consuming and expensive. There are too few genetic counsellors to expand this model to testing all prostate cancer patients and their specialized skills and training are most needed to support patients who receive a positive genetic test result. To expand genetic testing, other options should be considered. Many models are being explored and implemented including direct-to-consumer genetic testing, sponsored genetic testing, and physician-mediated genetic testing (mainstream testing). Each model has unique challenges. McCuaig et al. provide an informative summary of this in regard to testing all women with epithelial ovarian cancer; these considerations also apply to men with prostate cancer¹⁰⁶.

TheScreenProject is an example of a guided direct-to-consumer genetic testing model for BRCA1 and BRCA2, with the patient given the option of expanded panel testing¹⁰⁷. All men and women over age 18 are eligible to self-refer to the study including those with and without cancer. The test is available to all men with prostate cancer, regardless of clinical presentation or family history. They register online, are invited to watch a pre-test counselling video, provide informed consent online, pay a fee for testing and are sent a saliva kit to submit their DNA for testing. A genetic counsellor is available throughout the process to all participants. Those participants who receive a positive test result are contacted by the study genetic counsellor and referred to a local clinic for counselling and follow-up. Where available, men with a positive result are referred to a high-risk clinic that provides both clinical screening and opportunities for further research studies¹⁰⁸. In the first phase of the study the overall satisfaction with the process was very high. Only 5% of participants with a negative result requested genetic counselling. The principal difference between TheScreen-**Project** and other Canadian testing programs is that the patient is required to pay 250 USD to be tested whereas provincial programs are without cost to the patient (if eligible).

CONCLUSIONS

The genetic basis of prostate cancer has been well established, the most relevant genes identified and we are now in the position to translate this information into better cancer care. Challenges include identifying more men who carry a cancer predisposing mutation, in particular prior to a cancer diagnosis or at the time of early diagnosis. Genetic testing should be done according to the presentation of prostate cancer and the family history of prostate and ovarian cancer or where there is a known mutation in the family. The majority of the genes involved in homologous recombination repair which predispose women to breast cancer

are also associated with high Gleason grade prostate cancer and with a propensity to metastasize. It is probably unwise to consider these men as candidates for active surveillance.

There are several priority areas for further research:

To identify means to prevent hereditary prostate cancer.

To estimate the mortality rates for men with localised prostate cancer and a susceptibility gene.

To determine the impact of screening with PSA and MRI on mortality.

To compare the outcomes of radical prostatectomy and radiotherapy in men with inherited susceptibility mutations.

To develop personalised treatment options. There is evidence that specific treatment (i.e. olaparib) may be warranted for *BRCA2* carriers but this is not yet the case for men with mutations in other genes.

It is important that collaborative observational studies are conducted which follow men with mutations in order to evaluate various prevention and treatment options. Translational research is needed so that the benefits of genetic testing that have been realized for women are realized for men. The goal should be to expand our current focus from the treatment of men with metastatic disease towards prevention, screening and the treatment of early-stage disease.

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REFERENCES

- 1. National Cancer Institute, Surveillance Epidemiology and End Results Program: Cancer Stat Facts: Prostate Cancer. (2020).
- 2. Tsodikov, A. et al. Is prostate. Cancer Differ. black men.? Answ. 3 Nat. Hist. models Cancer 123, 2312–2319 (2017).
- Bruner, D. W., Moore, D., Parlanti, A., Dorgan, J. & Engstrom, P. Relative risk of prostate cancer for men with affected relatives: systematic review and metaanalysis. Int J. Cancer 107, 797–803 (2003).
- Goldgar, D. E., Easton, D. F., Cannon-Albright, L. A. & Skolnick, M. H. Systematic population-based assessment of cancer risk in first-degree relatives of cancer probands. J. Natl Cancer Inst. 86, 1600–1608 (1994).
- Schaid, D. J. The complex genetic epidemiology of prostate cancer. Hum. Mol. Genet 13, R103–R121 (2004).
- Schumacher, F. R. et al. Association analyses of more than 140,000 men identify 63 new prostate cancer susceptibility loci. *Nat. Genet* 50, 928–936 (2018).
- Lecarpentier, J. et al. Prediction of Breast and Prostate Cancer Risks in Male BRCA1 and BRCA2 Mutation Carriers Using Polygenic Risk Scores. J. Clin. Oncol. 35, 2240–2250 (2017).
- Barnes, D. R. et al. Breast and Prostate Cancer Risks for Male BRCA1 and BRCA2 Pathogenic Variant Carriers Using Polygenic Risk Scores. J. Natl. Cancer Inst. 114, diab147 (2021)
- Amin Al Olama, A. et al. Risk Analysis of Prostate Cancer in PRACTICAL, a Multinational Consortium, Using 25 Known Prostate Cancer Susceptibility Loci. Cancer Epidemiol. Biomark. Prev. 24, 1121–1129 (2015).
- Liede, A. et al. Evaluation of the needs of male carriers of mutations in BRCA1 or BRCA2 who have undergone genetic counseling. Am. J. Hum. Genet 67, 1494–1504 (2000).
- 11. NCCN. Clinical Practice Guidelines in Oncology, Genetic/Familial High-Risk Assessment: Breast, Ovarian, and Pancreatic V2.2021.
- van Adelsberg, J. et al. The VIOXX in Prostate Cancer Prevention study: cardiovascular events observed in the rofecoxib 25 mg and placebo treatment groups. Curr. Med Res Opin. 23, 2063–2070 (2007).
- Klein, E. A. et al. Vitamin E and the risk of prostate cancer: the Selenium and Vitamin E Cancer Prevention Trial (SELECT). JAMA 306, 1549–1556 (2011).
- Thompson, I. M. et al. Long-term survival of participants in the prostate cancer prevention trial. N. Engl. J. Med 369, 603–610 (2013).
- Andriole, G. L. et al. Effect of dutasteride on the risk of prostate cancer. N. Engl. J. Med 362, 1192–1202 (2010).
- Schröder, F. H. et al. Screening and prostate-cancer mortality in a randomized European study. N. Engl. J. Med 360, 1320–1328 (2009).
- Hugosson, J. et al. Eighteen-year follow-up of the Göteborg Randomized Population-based Prostate Cancer Screening Trial: effect of sociodemographic variables on participation, prostate cancer incidence and mortality. Scand. J. Urol. 52, 27–37 (2018).



- Pinsky, P. F. et al. Extended mortality results for prostate cancer screening in the PLCO trial with median follow-up of 15 years. Cancer 123, 592–599 (2017).
- USPSTF. U. S. Preventive Services Task Force: Final Recomendation Statement: Prostate Cancer Screening. (2018).
- NCCN. Clinical Practice Guidelines in Oncology, Prostate Cancer Early Detection V1.2021.
- ACS. American Cancer Society Recommendations for Prostate Cancer Early Detection, https://tinyurl.com/374wh3h4 (2021).
- Page, E. C. et al. Interim Results from the IMPACT Study: Evidence for Prostatespecific Antigen Screening in BRCA2 Mutation Carriers. Eur. Urol. 76, 831–842 (2019)
- Segal, N. et al. Imaging-based prostate cancer screening among BRCA mutation carriers-results from the first round of screening. *Ann. Oncol.* 31, 1545–1552 (2020).
- 24. Wokołorczyk, D. et al. PALB2 mutations and prostate cancer risk and survival. *Br. J. Cancer* **125**, 569–575 (2021).
- Darst, B. F. et al. Germline Sequencing DNA Repair Genes in 5545 Men With Aggressive and Nonaggressive Prostate Cancer. J. Natl Cancer Inst. 113, 616–625 (2021).
- CCS. Canadian Cancer Statistics Advisory Committee. Canadian Cancer Statistics 2018. Toronto, ON: Canadian Cancer Society; 2018. Available at: cancer.ca/ Canadian-Cancer-Statistics-2018-EN (accessed [date]).
- Siegel, D. A., O'Neil, M. E., Richards, T. B., Dowling, N. F. & Weir, H. K. Prostate Cancer Incidence and Survival, by Stage and Race/Ethnicity - United States, 2001-2017. MMWR Morb. Mortal. Wkly Rep. 69, 1473–1480 (2020).
- Carter, H. B. et al. Germline Mutations in ATM and BRCA1/2 Are Associated with Grade Reclassification in Men on Active Surveillance for Prostate Cancer. Eur. Urol. 75, 743–749 (2019).
- Castro, E. et al. Germline BRCA mutations are associated with higher risk of nodal involvement, distant metastasis, and poor survival outcomes in prostate cancer. J. Clin. Oncol. 31, 1748–1757 (2013).
- Giri, V. N. et al. Implementation of Germline Testing for Prostate Cancer: Philadelphia Prostate Cancer Consensus Conference 2019. J. Clin. Oncol. 38, 2798–2811 (2020).
- Castro, E. et al. Effect of BRCA Mutations on Metastatic Relapse and Causespecific Survival After Radical Treatment for Localised Prostate Cancer. Eur. Urol. 68, 186–193 (2015).
- Iannuzzi, C. M., Atencio, D. P., Green, S., Stock, R. G. & Rosenstein, B. S. ATM mutations in female breast cancer patients predict for an increase in radiationinduced late effects. *Int J. Radiat. Oncol. Biol. Phys.* 52, 606–613 (2002).
- Cesaretti, J. A. et al. ATM sequence variants are predictive of adverse radiotherapy response among patients treated for prostate cancer. Int J. Radiat. Oncol. Biol. Phys. 61, 196–202 (2005).
- Bergom, C. et al. The Implications of Genetic Testing on Radiation Therapy Decisions: A Guide for Radiation Oncologists. Int J. Radiat. Oncol. Biol. Phys. 105, 698–712 (2019).
- 35. Morash, C. et al. Active surveillance for the management of localized prostate cancer: Guideline recommendations. *Can. Urol. Assoc. J.* **9**, 171–178 (2015).
- Klotz, L. et al. Long-term follow-up of a large active surveillance cohort of patients with prostate cancer. J. Clin. Oncol. 33, 272–277 (2015).
- Bokhorst, L. P. et al. A Decade of Active Surveillance in the PRIAS Study: An Update and Evaluation of the Criteria Used to Recommend a Switch to Active Treatment. Eur. Urol. 70, 954–960 (2016).
- Han, M., Partin, A. W., Pound, C. R., Epstein, J. I. & Walsh, P. C. Long-term biochemical disease-free and cancer-specific survival following anatomic radical retropubic prostatectomy. The 15-year Johns Hopkins experience. *Urol. Clin. North Am.* 28, 555–565 (2001).
- Gonzalez, C. M. et al. Preoperative PSA level significantly associated with interval to biochemical progression after radical retropubic prostatectomy. *Urology* 64, 723–728 (2004).
- Hull, G. W. et al. Cancer control with radical prostatectomy alone in 1,000 consecutive patients. J. Urol. 167, 528–534 (2002).
- Amling, C. L. et al. Long-term hazard of progression after radical prostatectomy for clinically localized prostate cancer: continued risk of biochemical failure after 5 years. J. Urol. 164, 101–105 (2000).
- Pound, C. R. et al. Natural history of progression after PSA elevation following radical prostatectomy. JAMA 281, 1591–1597 (1999).
- Tutt, A. N. J. et al. Adjuvant Olaparib for Patients with BRCA1- or BRCA2-Mutated Breast Cancer. N. Engl. J. Med. 384, 2394–2405 (2021).
- Kirby, M., Hirst, C. & Crawford, E. D. Characterising the castration-resistant prostate cancer population: a systematic review. *Int J. Clin. Pr.* 65, 1180–1192 (2011).
- Annala, M. et al. Circulating Tumor DNA Genomics Correlate with Resistance to Abiraterone and Enzalutamide in Prostate Cancer. Cancer Disco. 8, 444–457 (2018).

- Castro, E. et al. PROREPAIR-B: A Prospective Cohort Study of the Impact of Germline DNA Repair Mutations on the Outcomes of Patients With Metastatic Castration-Resistant Prostate Cancer. J. Clin. Oncol. 37, 490–503 (2019).
- Vandekerkhove, G. et al. Circulating Tumor DNA Abundance and Potential Utility in De Novo Metastatic Prostate Cancer. Eur. Urol. 75, 667–675 (2019).
- 48. Lozano, R. et al. Genetic aberrations in DNA repair pathways: a cornerstone of precision oncology in prostate cancer. *Br. J. Cancer* **124**, 552–563 (2021).
- de Bono, J. et al. Olaparib for Metastatic Castration-Resistant Prostate Cancer. N. Engl. J. Med. 382, 2091–2102 (2020).
- Abida, W. et al. Rucaparib in Men With Metastatic Castration-Resistant Prostate Cancer Harboring a. J. Clin. Oncol. 38, 3763–3772 (2020).
- Abida, W. et al. Analysis of the Prevalence of Microsatellite Instability in Prostate Cancer and Response to Immune Checkpoint Blockade. *JAMA Oncol.* 5, 471–478 (2019).
- Ford, D., Easton, D. F., Bishop, D. T., Narod, S. A. & Goldgar, D. E. Risks of cancer in BRCA1-mutation carriers. Breast Cancer Linkage Consortium. *Lancet* 343, 692–695 (1994).
- Dong, X. et al. Mutations in CHEK2 associated with prostate cancer risk. Am. J. Hum. Genet. 72, 270–280 (2003).
- Seppälä, E. H. et al. CHEK2 variants associate with hereditary prostate cancer. Br. J. Cancer 89, 1966–1970 (2003).
- Angèle, S. et al. ATM polymorphisms as risk factors for prostate cancer development. Br. J. Cancer 91, 783–787 (2004).
- Pritchard, C. C. et al. Inherited DNA-Repair Gene Mutations in Men with Metastatic Prostate Cancer. N. Engl. J. Med 375, 443–453 (2016).
- Grindedal, E. M. et al. Germ-line mutations in mismatch repair genes associated with prostate cancer. Cancer Epidemiol. Biomark. Prev. 18, 2460–2467 (2009).
- Lee, D. J. et al. Association of Inherited Mutations in DNA Repair Genes with Localized Prostate Cancer. Eur Urol. 18, 559–567 (2021).
- Leongamornlert, D. et al. Frequent germline deleterious mutations in DNA repair genes in familial prostate cancer cases are associated with advanced disease. Br. J. Cancer 110, 1663–1672 (2014).
- Network, C. G. A. R. The Molecular Taxonomy of Primary Prostate Cancer. Cell 163, 1011–1025 (2015).
- Na, R. et al. Germline Mutations in ATM and BRCA1/2 Distinguish Risk for Lethal and Indolent Prostate Cancer and are Associated with Early Age at Death. Eur. Urol. 71, 740–747 (2017).
- Nicolosi, P. et al. Prevalence of Germline Variants in Prostate Cancer and Implications for Current Genetic Testing Guidelines. *JAMA Oncol.* 5, 523–528 (2019).
- Liede, A., Karlan, B. Y. & Narod, S. A. Cancer risks for male carriers of germline mutations in BRCA1 or BRCA2: a review of the literature. *J. Clin. Oncol.* 22, 735–742 (2004).
- 64. Thompson, D. & Easton, D. F., Breast Cancer Linkage, C. Cancer Incidence in BRCA1 mutation carriers. *J. Natl Cancer Inst.* **94**, 1358–1365 (2002).
- Leongamornlert, D. et al. Germline BRCA1 mutations increase prostate cancer risk. Br. J. Cancer 106, 1697–1701 (2012).
- Nyberg, T. et al. Prostate Cancer Risks for Male BRCA1 and BRCA2 Mutation Carriers: A Prospective Cohort Study. Eur. Urol. 77, 24–35 (2020).
- Kuchenbaecker, K. B. et al. Risks of Breast, Ovarian, and Contralateral Breast Cancer for BRCA1 and BRCA2 Mutation Carriers. JAMA 317, 2402–2416 (2017).
- Consortium, B. C. L. Cancer risks in BRCA2 mutation carriers. J. Natl Cancer Inst. 91, 1310–1316 (1999).
- Kote-Jarai, Z. et al. BRCA2 is a moderate penetrance gene contributing to young-onset prostate cancer: implications for genetic testing in prostate cancer patients. Br. J. Cancer 105, 1230–1234 (2011).
- Mitra, A. et al. Prostate cancer in male BRCA1 and BRCA2 mutation carriers has a more aggressive phenotype. Br. J. Cancer 98, 502–507 (2008).
- Tryggvadóttir, L. et al. Prostate cancer progression and survival in BRCA2 mutation carriers. J. Natl Cancer Inst. 99, 929–935 (2007).
- Akbari, M. R. et al. The impact of a BRCA2 mutation on mortality from screendetected prostate cancer. Br. J. Cancer 111, 1238–1240 (2014).
- Swift, M., Reitnauer, P. J., Morrell, D. & Chase, C. L. Breast and other cancers in families with ataxia-telangiectasia. N. Engl. J. Med 316, 1289–1294 (1987).
- 74. Jerzak, K. J., Mancuso, T. & Eisen, A. Ataxia-telangiectasia gene. *Curr. Oncol.* 25, e176–e180 (2018).
- van Os, N. J. et al. Health risks for ataxia-telangiectasia mutated heterozygotes: a systematic review, meta-analysis and evidence-based guideline. Clin. Genet 90, 105–117 (2016).
- Darst, B. F. Findings from a Genetic Sequencing Investigation of Men with Familial and Aggressive Prostate Cancer. Eur. Urol. 79, 362–363 (2021).
- Wokołorczyk, D. et al. Mutations in ATM, NBN and BRCA2 predispose to aggressive prostate cancer in Poland. Int J. Cancer 147, 2793–2800 (2020).
- Dorling, L. et al. Breast Cancer Risk Genes Association Analysis in More than 113,000 Women. N. Engl. J. Med 384, 428–439 (2021).

- 79. Hu, C. et al. A Population-Based Study of Genes Previously Implicated in Breast Cancer, N. Engl. J. Med 384, 440-451 (2021).
- 80. Cybulski, C. et al. Risk of breast cancer in women with a CHEK2 mutation with and without a family history of breast cancer. J. Clin. Oncol. 29, 3747-3752 (2011).
- 81. Siołek, M. et al. CHEK2 mutations and the risk of papillary thyroid cancer. Int J. Cancer 137, 548-552 (2015).
- 82. Wasielewski, M. et al. CHEK2 1100delC and male breast cancer in the Netherlands. Breast Cancer Res Treat. 116, 397-400 (2009).
- 83. Hale, V., Weischer, M. & Park, J. Y. CHEK2 (*) 1100delC Mutation and Risk of Prostate Cancer. Prostate Cancer 2014, 294575 (2014).
- 84. Cybulski, C. et al. An inherited NBN mutation is associated with poor prognosis prostate cancer. Br. J. Cancer. 108, 461-468 (2013).
- 85. Antoniou, A. C. et al. Breast-cancer risk in families with mutations in PALB2. N. Engl. J. Med 371, 497-506 (2014).
- 86. Yang, X. et al. Cancer Risks Associated With Germline PALB2 Pathogenic Variants: An International Study of 524 Families. J. Clin. Oncol. 38, 674–685
- 87. Fewings, E. et al. Germline pathogenic variants in PALB2 and other cancerpredisposing genes in families with hereditary diffuse gastric cancer without CDH1 mutation: a whole-exome sequencing study. Lancet Gastroenterol. Hepatol. 3, 489-498 (2018).
- 88. Huang, K. L. et al. Pathogenic Germline Variants in 10,389 Adult Cancers. Cell 173, 355-370.e314 (2018).
- 89. Barrow, E. et al. Cumulative lifetime incidence of extracolonic cancers in Lynch syndrome: a report of 121 families with proven mutations. Clin. Genet 75,
- 90. Ten Broeke, S. W. et al. Cancer Risks for PMS2-Associated Lynch Syndrome. J. Clin. Oncol. 36, 2961-2968 (2018).
- 91. Møller, P. et al. Cancer risk and survival in path_MMR carriers by gene and gender up to 75 years of age: a report from the Prospective Lynch Syndrome Database, Gut 67, 1306-1316 (2018).
- 92. Bonadona, V. et al. Cancer risks associated with germline mutations in MLH1. MSH2, and MSH6 genes in Lynch syndrome. JAMA 305, 2304-2310 (2011).
- 93. Raymond, V. M. et al. Elevated risk of prostate cancer among men with Lynch syndrome. J. Clin. Oncol. 31, 1713-1718 (2013).
- 94. Dominguez-Valentin, M. et al. Cancer risks by gene, age, and gender in 6350 carriers of pathogenic mismatch repair variants; findings from the Prospective Lynch Syndrome Database. Genet Med 22, 15-25 (2020).
- 95. Engel, C. et al. Risks of less common cancers in proven mutation carriers with lynch syndrome. J. Clin. Oncol. 30, 4409-4415 (2012).
- 96. Ryan, S., Jenkins, M. A. & Win, A. K. Risk of prostate cancer in Lynch syndrome: a systematic review and meta-analysis. Cancer Epidemiol. Biomark. Prev. 23, 437-449 (2014).
- 97. Kote-Jarai, Z. et al. Prevalence of the HOXB13 G84E germline mutation in British men and correlation with prostate cancer risk, tumour characteristics and clinical outcomes. Ann. Oncol. 26, 756-761 (2015).
- 98. Akbari, M. R. et al. Association between germline HOXB13 G84E mutation and risk of prostate cancer. J. Natl Cancer Inst. 104, 1260-1262 (2012).
- 99. Xu, J. et al. HOXB13 is a susceptibility gene for prostate cancer: results from the International Consortium for Prostate Cancer Genetics (ICPCG). Hum. Genet 132, -14 (2013).
- 100. Laitinen, V. H. et al. HOXB13 G84E mutation in Finland: population-based analysis of prostate, breast, and colorectal cancer risk. Cancer Epidemiol. Biomark. Prev. 22, 452-460 (2013).
- 101. Nyberg, T. et al. Homeobox B13 G84E Mutation and Prostate Cancer Risk. Eur. Urol. 75, 834-845 (2019).
- 102. Kluźniak. W. et al. The G84E mutation in the HOXB13 gene is associated with an increased risk of prostate cancer in Poland. Prostate 73, 542-548 (2013).
- 103. Ewing, C. M. et al. Germline mutations in HOXB13 and prostate-cancer risk, N. Engl. J. Med 366, 141-149 (2012).
- 104. Chen, Z. et al. The G84E mutation of HOXB13 is associated with increased risk for prostate cancer: results from the REDUCE trial. Carcinogenesis 34, 1260-1264 (2013).

- 105. NCCN. Clinical Practice Guidelines in Oncology, Genetic/Familial High-Risk Assessment: Colorectal V1, 2021.
- 106. McCuaig, J. M. et al. Evolution of genetic assessment for BRCA-associated gynaecologic malignancies: a Canadian multisociety roadmap. J. Med Genet 55, 571-577 (2018).
- 107. Narod, S. A. et al. The Screen Project: Guided Direct-To-Consumer Genetic Testing for Breast Cancer Susceptibility in Canada, Cancers (Basel) 13, 1894 (2021).
- 108. Lorentz, J., Liu, S. K. & Vesprini, D. Male Oncology Research and Education program for men at high risk for prostate cancer. Curr. Oncol. 25, 170–175 (2018).
- 109. Harter, P. et al. Prevalence of deleterious germline variants in risk genes including BRCA1/2 in consecutive ovarian cancer patients (AGO-TR-1). PLoS One 12. e0186043 (2017).

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AUTHOR CONTRIBUTIONS

A.F., R.C. and S.N. performed the literature review and wrote the first draft of the paper. All other authors reviewed and edited the paper to create the final draft.

COMPETING INTERESTS

Dr. Neil Fleshner has received consulting fees from Amgen, Janssen, Astellas, Bayer, Sanofi, Abbvie, Ferring, has stock holdings in POINT Biopharma and Verity pharmaceuticals and is the Chief Medical Officer for Verity pharmaceuticals and POINT Biopharma. Dr. Fleshner holds research grants (paid to the institution) from Janssen, Astellas, Bayer, Sanofi, Nucleix, Progenix, SpectraCure AB, Bavarian Nordic. Dr. Mohammad Akbari holds stock and other ownership interests in Genewise Inc. All other authors have no competing interests to report.

ADDITIONAL INFORMATION

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