



BRCA genetic testing and counseling in breast cancer: how do we meet our patients' needs?



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BRCA1 and *BRCA2* are tumor suppressor genes that have been linked to inherited susceptibility of breast cancer. Germline *BRCA1/2* pathogenic or likely pathogenic variants (gBRCAm) are clinically relevant for treatment selection in breast cancer because they confer sensitivity to poly(ADP-ribose) polymerase (PARP) inhibitors. *BRCA1/2* mutation status may also impact decisions on other systemic therapies, risk-reducing measures, and choice of surgery. Consequently, demand for gBRCAm testing has increased. Several barriers to genetic testing exist, including limited access to testing facilities, trained counselors, and psychosocial support, as well as the financial burden of testing. Here, we describe current implications of gBRCAm testing for patients with breast cancer, summarize current approaches to gBRCAm testing, provide potential solutions to support wider adoption of mainstreaming testing practices, and consider future directions of testing.

BRCA1 and *BRCA2* were identified in the 1990s as genes linked to inherited susceptibility to breast cancer^{1,2}. As tumor suppressor genes, they encode proteins that are crucial for the repair of complex DNA damage (such as double-strand breaks) by homologous recombination³. Germline mutations (i.e., pathogenic or likely pathogenic variants) in *BRCA1/2* (gBRCAm) affecting this vital DNA repair pathway predispose individuals to developing breast cancer by impairing homologous recombination and causing genomic instability³.

The advent of poly(ADP-ribose) polymerase (PARP) inhibitors has revolutionized the therapeutic landscape for cancers associated with gBRCAm, including breast, ovarian, prostate, and pancreatic cancer⁴. For breast cancer, the focus of this article, PARP inhibitors are approved for early and advanced disease harboring gBRCAm based on the results of major clinical trials: for olaparib, OlympiAD and OlympiA; and for talazoparib, EMBRACA⁵⁻⁷. Given the opportunity for therapeutic targeting of gBRCAm, timely determination of gBRCAm status is critical to guide treatment decisions, and demand for gBRCAm testing has rapidly increased in recent years⁸. High-throughput sequencing technologies have made analysis of cancer-susceptibility genes rapid and affordable⁸. However, there is concern that the demand for gBRCAm testing may overwhelm current

genetic services⁹. Furthermore, barriers at the individual-, provider-, systems-, and policy-levels exist, which restrict access to genetic testing resources and genetic counseling¹⁰. Innovative methods of mainstreaming genetic services may help overcome some of these challenges. Education and resources to support appropriate counseling for gBRCAm testing, as well as information on the implications of testing, and models for genetic test consent, are urgently needed to support the evolving clinical space.

In this review, we describe the implications of gBRCAm testing for potential surgical approaches and treatment in patients with breast cancer, summarize the various approaches to gBRCAm testing (including traditional and alternative models), provide practical resources to support mainstreaming of the gBRCAm testing pathway, and consider the relevance of genetic testing in breast cancer in the future.

Biology of BRCAm in breast cancer

Hereditary breast and ovarian cancer (HBOC) syndrome accounts for approximately 10% of breast cancer cases¹¹. *BRCA1* and *BRCA2* are the main genes involved in genetic susceptibility to breast cancer¹². HBOC is associated with early-onset breast cancer, and an increased risk of other cancers, including ovarian, pancreatic, fallopian tube, and prostate³. The

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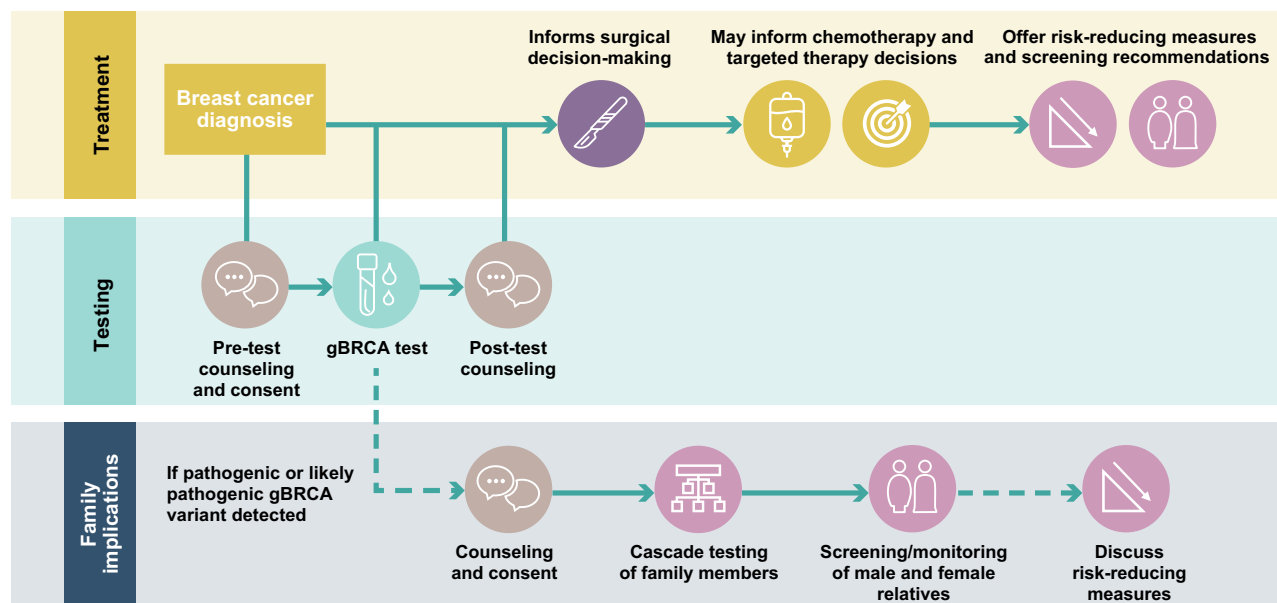


Fig. 1 | The pathway from gBRCAm testing to decisions relating to risk-reducing measures, choice of surgery, and systemic therapies.

cumulative lifetime risk of developing breast cancer by age 80 years is high at 72 and 69% for *BRCA1* and *BRCA2* mutation carriers, respectively¹³. Female gBRCAm carriers also have a 44% (*BRCA1*) and 17% (*BRCA2*) cumulative risk of developing ovarian cancer¹³.

Patients harboring gBRCAm are more likely to develop breast cancer at a younger age, with approximately 12% of the cases arising in women ≤ 40 years of age attributed to pathogenic or likely pathogenic variants in *BRCA1* or *BRCA2*¹⁴. These breast cancers have distinct biological features: among individuals with gBRCA1m, breast cancers are typically hormone receptor-negative (~76%) and human epidermal growth factor receptor 2 (HER2)-negative (94%), while breast cancers developing in individuals with gBRCA2m are more frequently hormone receptor-positive (83%) and HER2-negative (89%)¹⁴.

Goals of gBRCAm testing in breast cancer

Available evidence regarding surgical and systemic treatment outcomes in patients with gBRCAm breast cancer highlights the importance of determining gBRCAm status prior to finalizing treatment decisions. Clinical practice guidelines further reinforce the role of gBRCAm testing in the context of treatment decision-making, beyond its importance for risk management and cascade testing^{11,15}. The presence of gBRCAm may impact decisions about risk-reducing measures, choice of surgery, and systemic therapies (Fig. 1).

Surgical decision-making

Breast-conserving surgery (BCS). BCS aims to remove the breast tumor, with clear margins, in a manner that is cosmetically acceptable to the patient¹⁶. Although BCS is recommended for most patients with early-stage operable breast cancer¹⁵, the best approach for patients harboring gBRCAm is unclear. Practice guidelines recommend that gBRCAm status should not preclude the use of BCS as a surgical option for breast cancer¹⁷. However, these patients should be counseled regarding the risk of ipsilateral breast cancer recurrence, new primary breast cancer in the treated breast, and contralateral breast cancer, noting that intensified surveillance is a reasonable treatment strategy for breast cancer^{11,17}.

Contralateral risk-reducing mastectomy (CRRM). Some women with a confirmed gBRCAm opt for CRRM over BCS, which is removal of the unaffected breast to reduce the risk of contralateral breast cancer, with or without the option of breast reconstruction¹⁸. A meta-analysis of

outcomes in patients with gBRCAm found that CRRM reduced the relative risk of contralateral breast cancer by 93% versus surveillance and significantly increased overall survival (OS) versus surveillance¹⁹. It should be noted that benefit from CRRM was not maintained in all studies after adjusting for confounding factors²⁰, and the absolute survival benefits of mastectomy (both ipsilateral and contralateral) are heavily dependent on patient prognosis; patients with aggressive types of disease, and especially those with little response from neoadjuvant systemic therapy regimens, are at higher risk from distant metastasis than local recurrence or a new primary in the contralateral breast.

Risk-reducing salpingo-oophorectomy (RRSO). While RRSO is indicated in gBRCAm carriers, its effect on breast cancer risk reduction is not clear²¹. A recent systematic review and meta-analysis of 21,022 patients demonstrated a 37 and 49% reduction in the risk of developing breast cancer following RRSO compared with no RRSO in patients with gBRCA1m and gBRCA2m, respectively, with the effect particularly pronounced in younger women with gBRCAm²². A retrospective analysis in 676 women harboring gBRCAm showed that oophorectomy decreased mortality in patients with gBRCA1m and decreased breast cancer-specific mortality in patients with estrogen receptor (ER)-negative gBRCAm breast cancer²³. Other studies have failed to demonstrate a benefit of RRSO on breast cancer risk^{24,25}.

Systemic treatment decision-making

Chemotherapy. gBRCAm advanced breast cancers are sensitive to platinum-based and non-platinum-based chemotherapy regimens^{26–29}. For early breast cancer, patients with gBRCAm are treated with anthracycline/taxane-based regimens, similar to those individuals with sporadic breast cancers³⁰. The clinical value of adding platinum therapy to neoadjuvant chemotherapy for patients with gBRCAm tumors is inconclusive. The phase 3 BrightNess trial concluded that the addition of carboplatin, with or without veliparib, to neoadjuvant chemotherapy significantly improved pathological complete response (pCR) rates among patients with triple-negative breast cancer (TNBC), regardless of gBRCA status³¹. Furthermore, a meta-analysis of neoadjuvant regimens in patients with gBRCAm TNBC reported improved pCR rates when platin derivatives were combined with anthracyclines and taxanes, although it was unclear if this combination offered a clinically meaningful benefit over standard chemotherapy alone³². However, GeparSixto and INFORM did not show a benefit to adding carboplatin or cisplatin,

respectively, to neoadjuvant chemotherapy for patients with gBRCAm early breast cancer^{26,33}. Exploratory translational analyses of BrighTNess sought to elucidate the differences in benefit observed for patients with breast cancer and gBRCAm³⁴. Higher PAM50 proliferation score, CIN70 score, and GeparSixto immune signature were associated with higher odds of pCR for both patients with and without gBRCAm, and thus have been proposed as potentially useful biomarkers for determining addition of carboplatin to neoadjuvant chemotherapy³⁴, but have yet to be validated for clinical practice.

PARP inhibition. PARP inhibitors block the enzyme that has a vital role in repairing DNA single-strand breaks. They exploit the double-strand break repair deficiency of BRCAm cells, which accumulate unrepaired, toxic DNA double-strand breaks, thus resulting in tumor cell death (i.e., synthetic lethality). Olaparib is licensed for the adjuvant treatment of gBRCAm, HER2-negative high-risk early breast cancer, and for gBRCAm (tumor BRCAm in Japan), HER2-negative locally advanced (EU) or metastatic (EU and US) breast cancer. Talazoparib is approved for the treatment of gBRCAm, HER2-negative locally advanced or metastatic breast cancer in the US, Europe, and several other countries worldwide.

For advanced gBRCAm HER2-negative breast cancer, PARP inhibitors were approved based on the results of the OlympiAD (olaparib) and EMBRACA (talazoparib) clinical trials^{5,6,35,36}. In OlympiAD, olaparib had significantly improved median progression-free survival (PFS) versus standard chemotherapy treatment of physician's choice (7.0 months vs 4.2 months; HR 0.58 [95% CI 0.43–0.80]; $P < 0.001$) in patients with gBRCAm HER2-negative metastatic breast cancer⁵. Median OS was 19.3 months for olaparib and 17.1 months for standard chemotherapy (HR 0.89 [95% CI 0.67–1.18])³⁵. In subanalyses, a potential OS benefit with first-line olaparib versus chemotherapy was observed (median 22.6 vs 14.7 months; HR 0.55 [95% CI 0.33–0.95]), with 3-year survival at 40.8% with olaparib and 12.8% with treatment of physician's choice, which, notably, did not include a platinum regimen^{5,35}. In EMBRACA, talazoparib significantly improved median PFS versus standard chemotherapy (8.6 vs 5.6 months; HR 0.54 [95% CI 0.41–0.71]; $P < 0.001$) in patients with gBRCAm advanced breast cancer⁶, with no observed improvements in OS³⁷.

For early breast cancer, olaparib was approved based on the results of the phase 3 OlympiA study in patients with high-risk early gBRCAm HER2-negative breast cancer who had completed local treatment and neoadjuvant or adjuvant chemotherapy^{7,38}. In the second prespecified analysis of OlympiA, adjuvant olaparib was associated with significantly improved OS versus placebo, with a 32% reduced risk of death (HR 0.68; 98.5% CI 0.47–0.97; $P = 0.009$)⁷. Significantly improved invasive disease-free survival (IDFS; HR 0.63; 95% CI 0.50–0.78) was also shown, consistent with the significantly improved IDFS reported at the first prespecified analysis (HR 0.58; 99.5% CI 0.41–0.82; $P = 0.001$)⁷.

These positive results in the adjuvant setting raised the question of whether PARP inhibitors may also have a place for neoadjuvant treatment of HER2-negative early breast cancer; however, trials have reported mixed results. In the BrighTNess trial, described above, addition of veliparib did not add benefit over neoadjuvant carboplatin/paclitaxel alone³¹. The phase 2 GeparOLA study comparing neoadjuvant paclitaxel plus olaparib to paclitaxel/carboplatinum in patients with HER2-negative breast cancer and homologous recombinant deficiency did not meet its primary endpoint (exclusion of a pCR rate of $\leq 55\%$)³⁹, but did report a numerically improved pCR rate with paclitaxel/olaparib followed by epirubicin/cyclophosphamide (55.1%) versus paclitaxel/carboplatinum (48.6%) followed by epirubicin/cyclophosphamide, and a more favorable tolerability profile for paclitaxel/olaparib³⁹. In the single-arm neoTALA trial, patients with gBRCAm, early-stage TNBC were treated with talazoparib followed by definitive surgery⁴⁰. Although neoadjuvant talazoparib was active, the pCR rates did not meet the prespecified threshold of efficacy⁴⁰. Other neoadjuvant trials are ongoing to enhance our understanding of the potential use of PARP inhibitors in early breast cancer. Of potential interest is the opportunity to evaluate

alternative PARP inhibitor combinations (e.g., with immunotherapy), and tailor therapy according to the patient. For example, in the ongoing OlympiAN trial (NCT05498155) patients with deleterious/suspected deleterious BRCAm and operable, early-stage, HER2-negative, ER-negative/ER-low breast cancer are assigned olaparib (lower-risk cohort) or olaparib plus durvalumab (higher-risk cohort), and assessed for pCR⁴¹.

PARP inhibitors are an important treatment strategy for gBRCAm breast cancer and rely on timely access to genetic testing to guide the most appropriate treatment selection, particularly in the early breast cancer setting.

Cyclin-dependent kinase 4/6 (CDK4/6) inhibitors. A CDK4/6 inhibitor in combination with endocrine therapy is a recommended option for first-line treatment for certain patients with hormone receptor-positive/HER2-negative advanced or metastatic breast cancer^{15,42}. Use of CDK4/6 inhibitors has also extended into earlier lines of treatment, with abemaciclib plus endocrine therapy a treatment option in the adjuvant setting for patients with hormone receptor-positive/HER2-negative, high-risk breast cancer¹⁵, and positive results having been reported for ribociclib (NATALEE)⁴³. While the optimal sequence is not known, recent guideline updates note that when patients are eligible for both adjuvant olaparib and abemaciclib then olaparib should be given first^{30,44}. Real-world evidence has suggested that patients with hormone receptor-positive advanced breast cancer and gBRCAm may have inferior outcomes with CDK4/6 inhibition or endocrine therapy versus those without gBRCAm^{45–49}. This emerging finding highlights the potential importance of early detection of gBRCAm in patients with hormone receptor-positive breast cancer ahead of treatment selection, especially in light of recent CDK4/6 inhibitor approval in the early breast cancer setting.

Immunotherapy. There is limited evidence on the effectiveness of immunotherapy in patients with gBRCAm breast cancer. A recent sub-study from the phase 3 IMpassion130 trial of the anti-programmed death-ligand 1 (PD-L1) antibody atezolizumab showed that, in combination with nab-paclitaxel, patients with PD-L1-positive advanced TNBC had an OS and PFS benefit regardless of *BRCA1/2* mutation status (germline or somatic)⁵⁰. The efficacy of neoadjuvant PARP inhibition in combination with immunotherapy is under investigation; for example, olaparib in combination with durvalumab is being investigated in the aforementioned OlympiAN study⁴¹.

Screening and counseling for family members

The burden of gBRCAm in breast cancer extends beyond the affected individual, with other family members facing decisions regarding gBRCAm testing, as well as considerations of family planning. In case of a familial association, genetic testing is recommended by the NCCN Clinical Practice Guidelines in Oncology (NCCN Guidelines[®]) for unaffected family members²¹. If a pre-symptomatic individual is identified as a carrier of gBRCAm, intensified surveillance for breast cancer is recommended, which differs per guideline but may include regular magnetic resonance imaging (MRI), ultrasound, mammography, and/or clinical breast exam, with guidance provided based on age^{11,21}. For patients harboring gBRCAm with a diagnosis of breast cancer who have not undergone bilateral mastectomy, National Comprehensive Cancer Network[®] (NCCN[®]) recommends that breast MRI and mammography should continue as recommended, based on age²¹.

For individuals undergoing pre-symptomatic testing (known gBRCAm in a family member), it is recommended that pre-test counseling topics include options for screening and early detection, the benefits and disadvantages of risk-reducing surgery (including the extent of cancer risk reduction, risks associated with surgery, management of menopausal symptoms with RRSO, psychosocial and quality-of-life impacts, and life expectancy), the benefits and limitations of reconstructive surgery and reproductive options, and the psychological implications of pre-

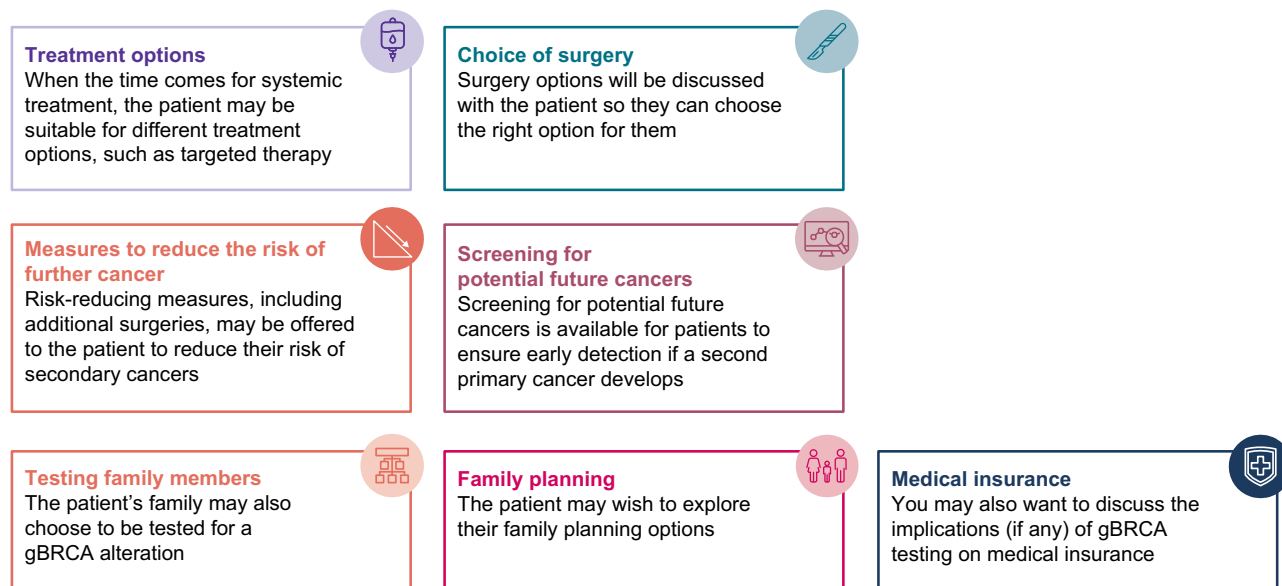


Fig. 2 | gBRCAm counseling and testing in clinical practice.

symptomatic diagnosis^{11,21}. Consideration is required with regard to reproductive concerns and the psychosocial impact of undergoing RRSO in gBRCAm carriers²¹.

gBRCAm counseling and testing in clinical practice

Implementation of guideline recommendations for gBRCAm counseling and testing

Practice guidelines for genetic counseling and gBRCAm testing are predominantly based on personal and family history of breast, ovarian, pancreatic, and/or prostate cancer; young age at diagnosis; male breast cancer; and multiple tumors (breast and ovarian) in the same patient²¹. More than 32 guidelines for gBRCAm testing relevant to breast cancer exist worldwide^{11,21,51,52}, and the recommendations are often inconsistent. Many guidelines do not include recommendations for genetic counseling, or only provide counseling recommendations for patients who have been identified as carriers of gBRCAm⁵¹. Some guidelines recommend gBRCAm testing after genetic counseling and personalized risk assessment, and/or if the result is likely to influence the individual's choice of primary treatment⁵¹. Some guidelines recommend testing based upon percentage risk of harboring a BRCA mutation, but there is a lack of consensus on the threshold used to determine whether an individual is eligible for genetics clinic referral/testing (10% vs 5%)^{21,53}, and some guidelines propose testing all patients under certain circumstances (e.g., with ER-positive advanced breast cancer and resistance to endocrine therapy), considering that PARP inhibitors have a greater risk-benefit ratio than chemotherapy⁵⁴. There are limited treatment recommendations and algorithms for women with gBRCAm-associated advanced breast cancer⁵¹. Greater consensus and cohesion of guidelines would be useful for patients and the medical community covering the topics highlighted in Fig. 2.

Disparities in gBRCAm testing in clinical practice

There has been a systemic underuse of gBRCAm testing over the past two decades, which has led to inappropriate and inconsistent testing and, consequently, missed opportunities for cancer prevention and management⁵⁵. Historically, NCCN criteria have been seen to be the least restrictive of the models, identifying a larger percentage of carriers compared with other models. However, the complex nature of the NCCN criteria render them difficult to implement in real-world clinical practice⁵¹, with low adherence rates reported⁵⁶. Expansion of NCCN criteria to include all women diagnosed at ≤ 65 years of age was shown to improve sensitivity of the selection criteria, without requiring testing of all women with breast cancer⁵⁷.

Although recent data from some centers and countries suggest widespread routine gBRCAm testing⁵⁸, a number of reports highlight the need for broader eligibility criteria for gBRCAm testing to ensure that more individuals can have access^{55,57,59,60}. Notably, patient eligibility for gBRCAm testing has been shown to vary depending on different testing criteria and recommendations, ranging from over 98% using recent guidelines published by the American Society of Breast Surgeons (ASBrS) to only around 30% eligibility using the Breast and Ovarian Analysis of Disease Incidence and Cancer Estimation Algorithm (BOADICEA) criteria^{55,57} (Fig. 3). Simplified, cost-effective eligibility criteria for gBRCAm testing, based on individual rather than family history criteria, have been proposed by the Mainstreaming Cancer Genetics (MCG) group. The five eligibility criteria include: (1) ovarian cancer diagnosis, (2) breast cancer diagnosed ≤ 45 years of age, (3) two primary breast cancers, both diagnosed ≤ 60 years of age, (4) TNBC diagnosis, and (5) male breast cancer diagnosis⁵⁵. In an analysis of different guidelines, using these criteria would have tested 92% of people and detected 100% of gBRCAm carriers⁵⁵. An additional sixth criteria (breast cancer, plus a parent, sibling, or child meeting any of the other criteria) further improved the eligibility rate to 97% (MCGplus)⁵⁵, while expansion of NCCN criteria (v1.2020) to include individuals diagnosed at ≤ 65 years of age, as recommended by ASBrS, increased testing eligibility to include over 98% of BRCAm carriers⁵⁷ (Fig. 3). Both the MCG and MCGplus criteria were deemed cost-effective, with cost-effective ratios of \$1330 and \$1225 per discounted quality-adjusted life year for the MCG and MCGplus criteria, respectively⁵⁵. Additional studies have sought to investigate the cost-effectiveness of BRCA testing in all patients with breast cancer, with several studies conducted in countries such as Australia, China, Norway, Malaysia, the UK, and the US finding this to be a potentially cost-effective strategy⁶¹⁻⁶⁵.

Traditional genetic counseling and testing pathway

The traditional pathway of genetic testing involves individualized patient referral to the genetics department for the management of pre-test genetic counseling, consenting, sample acquisition, and return of results (Fig. 4). Pre-test counseling, and the process of informed consent, focuses on giving patients sufficient information about the test, its limitations, and the consequences (including psychological) of a positive result, to enable an informed decision as to whether or not to proceed⁹. Patients who test positive for gBRCAm receive post-test support from a geneticist/genetic counselor/expert^{9,66}.

Genetic professionals offering counseling include both medical genetic physicians (professionals with advanced training, such as an

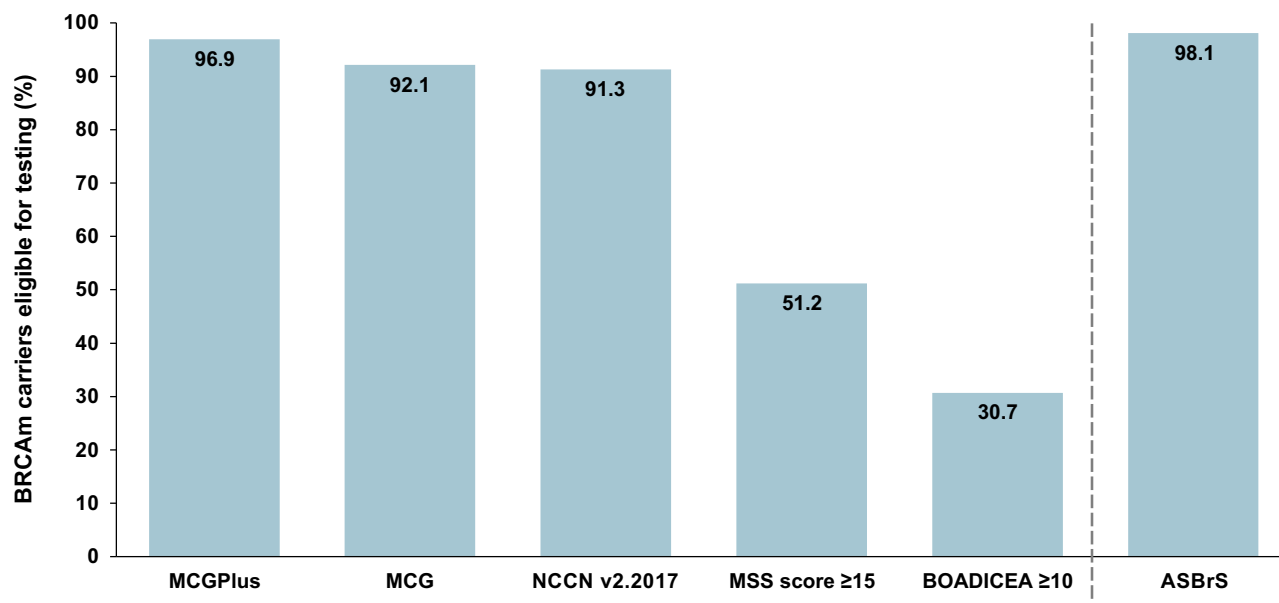


Fig. 3 | Eligibility for BRCAm testing using different testing criteria. The graph shows estimates of patient eligibility for BRCA testing among BRCAm carriers. Data to the left of the dashed line is reproduced from a report in 2019 by the MCG group assessing rates of testing eligibility by different criteria⁵⁵, while the bar to the right of the dashed line illustrates the result of an analysis by ASBrS published in 2020, examining the effect of including all individuals meeting NCCN criteria v1.2020 plus

those diagnosed with breast cancer at ≤ 65 years⁵⁷. ASBrS, American Society of Breast Surgeons; BOADICEA, Breast and Ovarian Analysis of Disease Incidence and Cancer Estimation Algorithm (≥ 10 refers to a 10% or greater probability that a BRCA1 or BRCA2 mutation is present); MCG, Mainstreaming Cancer Genetics; MSS, Manchester Scoring System; NCCN, National Comprehensive Cancer Network* (NCCN*).

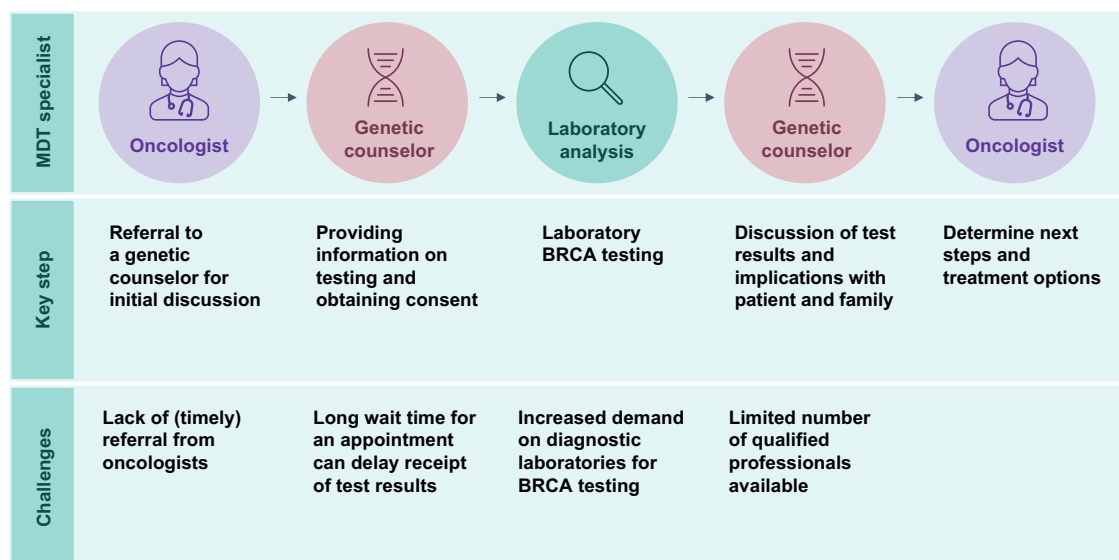


Fig. 4 | Key steps and challenge points in traditional gBRCAm testing pathways. MDT, multidisciplinary team.

MD with a specialization in genetic medicine) and genetic counselors (professionals with a specialized Masters degree in genetic counseling)^{67,68}. Genetic counseling by a trained genetics clinician has been shown to improve patient knowledge, understanding, and satisfaction among patients⁶⁹, and is recommended in multiple guidelines^{11,21}. While advantages of this type of care are clear, disadvantages include that it can be time-consuming, and a limited number of professionals are appropriately trained. When rapid access to test results is required to inform treatment decisions in a time-sensitive manner, especially for those undergoing upfront surgery, it may not be possible to maintain this workflow, and innovative alternatives may be required⁷⁰.

Although genetic counseling is recommended, a dearth of adequately trained professionals in this field may limit access⁷¹, with some countries imposing legal requirements for practicing genetic counseling⁷². Where possible, non-geneticist physicians might feel the need to counsel and test patients themselves without support, despite increasing demands on their time and shorter appointment times^{69,71}. Across Canada and the US, there are approximately 1.5 genetic counselors per 100,000 individuals, and it is estimated that double the workforce will be needed to meet future demands⁷³. There has been an increase in genetic counselors reporting the use of multiple types of delivery models, including telephone and tele-genetics, with an aim of improving access and efficiency of genetic counseling; however, barriers remain that can hinder implementation of these

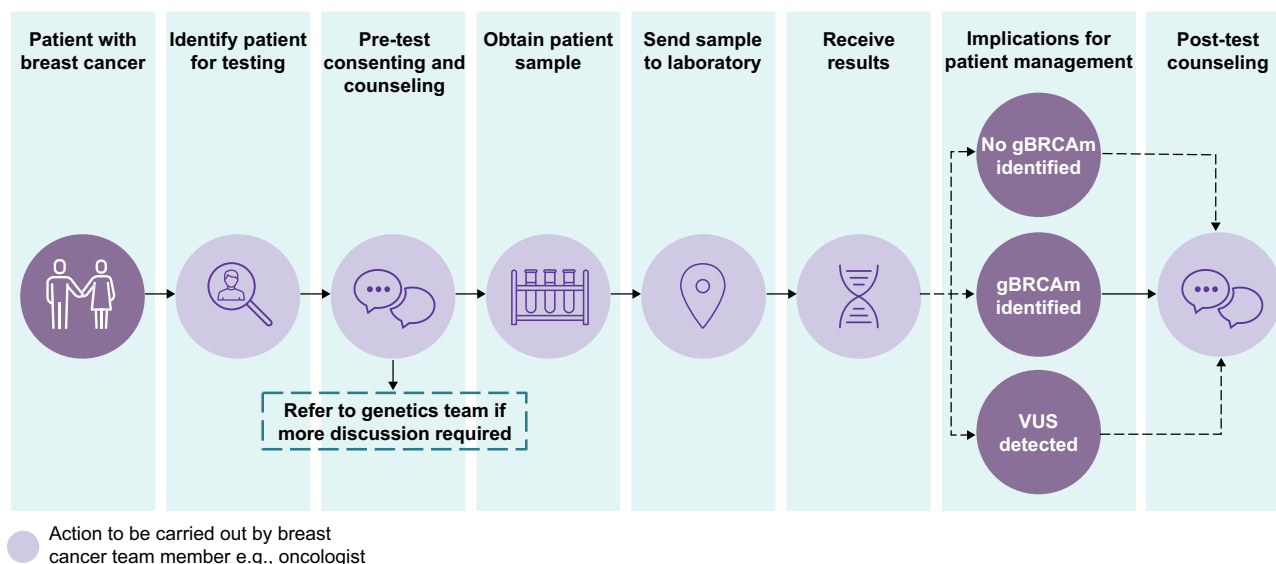


Fig. 5 | Example gBRCAm testing pathway to illustrate the mainstream genetic testing pathway. VUS, variant of uncertain significance.

models⁷⁴. In a large, US population-based study, only 62% of high-risk patients with newly diagnosed breast cancer who were tested had a genetic counselor session⁷⁵. Furthermore, 66% of all patients, and 81% of high-risk patients, wanted testing but only 29 and 53% received it, respectively⁷⁵. The most common reason for high-risk patients not being tested was “my doctor didn’t recommend it”⁷⁵. Wait times to see genetic specialists can also be substantial. In the UK, the Nottingham University Hospitals National Health Service (NHS) Trust reports wait times of 12–14 weeks for an initial appointment and 4–6 months to receive results⁷⁶. This highlights the need for alternative models of counseling and consenting of patients to ensure all eligible patients receive testing in a timely manner.

Systemic and societal barriers can impede equitable access to the benefits of genetic testing. Suboptimal testing rates among individuals of low socioeconomic status have been largely attributed to perceived/actual financial costs of genetic testing, with patients and healthcare providers often unclear as to whether genetic counseling services and follow-up care are covered by health insurance^{10,77}. Strategies to improve testing rates in this patient demographic include the integration of genetic counselors into primary care settings to reduce travel time and costs to the patient⁷⁸, and lobbying for expanded health insurance coverage for genetic counseling and testing services⁷⁹.

Reports from US ovarian and breast cancer centers have consistently found racial/ethnic disparity in access to genetic testing, with referral rates being higher for non-Hispanic White women than for women of other races^{80–82}. Lower awareness of the genetic basis of risk, incomplete family history, and mistrust of medical confidentiality may contribute to racial/ethnic disparities in referrals for genetic testing^{79,83}. In addition, the detection of pathogenic variants may be decreased, and variants of uncertain significance increased, in non-White individuals^{84–86}, as genomic reference databases provide poor genetic representation of non-White populations^{87,88}. Whilst initiatives have been established to address gaps in the diversity of genomic data⁸⁹, additional strategies are required to increase genetic testing rates among non-White populations. These include the development of culturally and linguistically tailored educational material, extended appointment availability, increased training of primary care-based specialists to mitigate unconscious or implicit biases, and a drive to recruit and train more healthcare providers from minority backgrounds^{79,80,90}.

Mainstream genetic counseling and testing pathways

In mainstream genetic testing pathways, medical oncology teams are responsible for pre-test genetic counseling, obtaining consent, scheduling the genetic test, and using the results to guide treatment decisions

(Fig. 5)^{55,91,92}. Implementation of mainstream models has enabled more efficient testing of patients with ovarian cancer and has significantly increased the proportion of patients being offered genetic testing^{93–95}.

Mainstream genetic testing models for patients with breast cancer have also proven effective, with high pathogenic variant detection rates and a reduced burden on genetic services observed^{55,66,76}. A Canadian study reported a significant decrease in wait time from referral to the return of genetic test results using an oncology clinic-based model compared with a traditional model in patients with breast or ovarian cancer (403 vs 191 days; $P < 0.001$)⁹⁶. Other studies support that oncologist-led mainstreaming results in increased testing uptake and shorter test-turnaround times^{87,98}. A systematic review of 15 studies in patients with breast, ovarian, endometrial, or prostate cancer showed that turnaround times with the mainstream approach are lower than those with the traditional pathway, with results typically obtained 3–6 weeks after discussing and ordering the genetic test⁹². Another study in patients with breast cancer measured an 85% reduction in time to test result using the mainstream model compared with the traditional model (4 vs 25 weeks)⁵⁵. A mainstreaming program in Australia had successful uptake with a notable gBRCAm detection rate and a reduced burden on the center, enabling reallocation of resources to streamline the genetic testing process⁶⁶. Mainstream models also reduce genetic consultation requirements versus traditional models^{55,66}.

Perspectives of the multidisciplinary team. Oncogenetic partnership models, in which clinical teams order genetic testing in collaboration with geneticists and implement counseling at both an individual and group level, have been shown to improve access to counseling and reduce turnaround times for genetic testing⁹⁹. However, the feasibility of implementing new testing strategies may vary by region.

As part of the MCG program in breast cancer in the UK and Malaysia, 100% of team members (12 oncologists, 8 surgeons, and 3 nurse specialists) reported feeling confident to approve patients for genetic testing, and believed that the process worked well⁵⁵. Similar experiences have been reported among ovarian cancer teams^{9,91}. However, another study surveying oncologists, clinical geneticists, and surgeons found that while oncologists and clinical geneticists were mainly positive about the introduction of mainstream approaches, surgeons were not keen to implement mainstream services in their breast clinic, feeling that they did not have the expertise, time, or capacity to undertake the extra responsibility, and that genetic testing did not have much relevance for their treatment decision-making¹⁰⁰.

Nurses play an integral role within the oncology team, with clinical nurse specialists often being the key point of contact for patients throughout

the cancer pathway and thus ideally placed to deliver information on gBRCAm testing. A single-center UK study assessing the use of clinical nurse specialists in consenting patients with ovarian cancer for gBRCAm testing showed that there was no difference in patient-reported satisfaction compared with oncologist-led consenting, and nurses felt confident in counseling, consenting, and returning results⁹. A specialist, nurse-led breast cancer MCG service established at the Nottingham Breast Institute, UK, has reduced wait time from the date of testing to the date of results to 36 days compared with an historical wait time of 4–6 months, while also delivering continuity of care for patients, releasing oncologists' time, and allowing oncologists and patients to consider treatment options at an earlier time point⁷⁶. The potential for nurses to play a role in decision coaching in healthy individuals who are carriers of gBRCAm is being explored, with preliminary results suggesting the approach is feasible^{101,102}. We provide an educational guide for nurses to outline the role that nurses can play in the gBRCA testing pathway and support conversations around nurse-obtained consent (supplementary information: *Nurse consenting guide for germline BRCA testing*).

The patients' perspective. Genetic testing in mainstream oncology units is widely accepted by patients^{55,66,91,103–106}. In the MCG breast cancer program, 96% of patients were happy that genetic testing was performed by their cancer team⁵⁵. Some patients reported a preference for their medical oncologist or their oncology nurse to deliver pre-test counseling, because medical oncologists could use the information gained through genetic testing for treatment decisions, and because nurses are more familiar with, and better understand, the individual patient experience¹⁰⁵.

Educational needs for non-genetic specialists. Ensuring appropriate training on an ongoing basis for those involved in consenting and arranging genetic testing is paramount to the success of mainstream gBRCAm testing⁶⁶. An early study evaluating patient experiences of gBRCAm testing in the US (all tumor types) found that the quality of information given to patients by non-certified genetic healthcare professionals (HCPs) was not as consistent as that given by certified genetic HCPs, with far fewer patients in mainstream testing versus traditional counseling recalling having had a pre-test discussion, and what that included¹⁰⁷. The types of training required by non-genetic specialists include generic consent training, plus specific genetics training, which involves learning how to identify eligible patients for gBRCAm testing, the relevance of gBRCAm testing, the significance for patients with a positive or negative result, the significance of a gBRCAm variant requiring evaluation, and implications of a positive test for family members⁹. Workshops designed to improve HCP knowledge and self-confidence have been shown to significantly enhance ability to overcome communication difficulties in relation to genetic testing and counseling¹⁰⁸. We provide educational guides to support healthcare providers in their understanding of the gBRCA testing and consenting pathway (supplementary information: *HCP guide to genetic counseling: Understanding germline BRCA testing and its clinical implications in breast cancer* and *Germline BRCA testing pathway infographic*), as well as useful language to help explain the process to patients (supplementary information: *Patient-HCP flipbook: What you need to know about BRCA testing*).

Use of digital tools

Digital tools are being increasingly used across the genetic testing pathway for clinical assessment, family history taking, education, post-test counseling, and follow-up, and include web-based tools, mobile applications, chatbots, videos, and games^{73,109,110}. They have been shown to improve access to genetic testing (particularly for patients in under-served areas), reduce waiting times, enhance continuity of care, increase patient engagement, and free up time for other patient-centered consultations^{73,110}. Digital tools are associated with positive patient outcomes, including increased knowledge and reduced decision conflict, and achieve similar patient outcomes to in-person consultations¹⁰⁹.

There are no digital tools that offer a comprehensive solution across the entire genetic counseling and testing pathway, with most tools developed for

use in the pre-test counseling phase only¹⁰⁹. The Genetics Navigator is currently in development and aims to supplement in-person consultations and support the full genetic testing pathway, including pre-test counseling, education, decision support, laboratory reporting, personalized return of results, and post-test counseling¹¹⁰. A digital pathway has also been integrated into the UK NHS clinical, laboratory, and informatics systems for delivery of gBRCAm testing to cancer patients and has been piloted as part of the BRCA-DIRECT study¹¹¹. Results demonstrated that uptake of genetic testing using the digital pathway was non-inferior to those receiving pre-test information via telephone, with similarly good patient satisfaction and knowledge and low anxiety scores¹¹¹.

The future of genetic testing in breast cancer

Germline versus somatic mutation testing

Genetic testing of tumor tissue has the potential to identify both germline and somatic pathogenic (or likely pathogenic) variants, and thus identify more people who might benefit from targeted therapies. Indeed, several studies have demonstrated clinical benefit with PARP inhibitor treatment for somatic BRCAm (sBRCAm) breast cancers^{112–114}. High concordance between germline and tumor BRCAm testing in breast and ovarian cancer has been observed^{115–120}; however, while sBRCAm and gBRCAm can be mutually exclusive in breast cancer, and not all mutation types can be detected by current clinical testing methods, it is possible that patients with metastatic breast cancer could benefit more from tumor testing than germline testing, as other abnormalities and targets could be identified^{121–123}. For example, approvals of alpelisib plus fulvestrant for the treatment of *PIK3CAm* advanced or metastatic breast cancer¹²⁴, capivasertib plus fulvestrant for advanced or metastatic breast cancer with *PIK3CAm*, *AKT1m*, or *PTENm*¹²⁵, and pembrolizumab for the treatment of unresectable or metastatic solid tumors of any type with high tumor mutational burden¹²⁶ may have led to an increase in patient referrals for tumor testing using a gene panel assay. An increasing number of patients with BRCAm breast cancer could, therefore, receive an incidental positive result for BRCAm and be subsequently offered a gBRCA test to confirm the somatic or germline status, in accordance with NCCN Guidelines^{*21}.

Parallel testing of normal and tumor material offers an alternative approach that allows direct differentiation of somatic versus germline pathogenic (or likely pathogenic) variants, leading to timely treatment selection and genetic counseling that may otherwise be delayed with germline- or tumor-only testing¹²⁷. Somatic testing alone would not distinguish between germline and somatic pathogenic (or likely pathogenic) variants, and thus may not be useful for determining future surveillance/surgery options for the patient, and may not benefit family members¹²⁸. Therefore, an increasing number of centers are moving toward parallel testing for patients with a breast cancer diagnosis¹²⁷. Analysis of circulating tumor DNA has the potential to identify both somatic and germline variants, and may offer a non-invasive alternative to tissue testing¹²⁹.

Genetic testing beyond BRCA

Panel testing allows for the screening of multiple genes beyond *BRCA1* and *BRCA2* that may be associated with tumor development and/or treatment response¹³⁰. For example, several other factors in the homologous recombination pathway have emerged as clinically relevant in surgical and treatment decision-making^{131,132}. Pathogenic variants in breast cancer susceptibility genes beyond *BRCA1* and *BRCA2* are increasingly being considered in clinical trials with targeted therapies^{113,133,134} and further recommendations for risk reduction, screening, and treatment strategies for carriers of these variants are being incorporated into clinical practice guideline updates and risk assessment tools^{11,21,52}. For example, current NCCN Guidelines recommend discussion of risk-reducing mastectomy with patients found to harbor pathogenic or likely pathogenic variants in *CDH1*, *PALB2*, *PTEN*, *STK11*, or *TP53*, and consideration of RRSO at 45–50 years of age in patients with pathogenic or likely pathogenic variants in *PALB2*, *RAD51C*, or *RAD51D*²¹. The web-based CanRisk tool, which integrates the presence of pathogenic variants in eight breast cancer susceptibility genes with several other risk

factors to estimate the personal risk of breast cancer, is currently endorsed by multiple clinical guidelines^{135,136}.

To summarize, advancements in patient information and care, in particular the introduction of PARP inhibitors for the treatment of breast and other cancers, have resulted in a substantial increase in demand for genetic testing. This demand is supported by the evidence that gBRCA testing in breast cancer management is a cost-effective strategy. However, without a substantial increase in personnel, traditional, genetics-led models of counseling and consenting are unable to meet the growing demand. A case can be made to increase the number of genetically trained HCPs but, even if possible, there will be a certain time lag before they are available. Mainstreaming models and the use of digital tools have demonstrated potential in providing efficient, patient-centered care that can meet the increasing needs of patients. In the future, we may see a move toward more widespread and comprehensive testing for germline and tumor alterations, raising further challenges as to how this can be effectively incorporated into comprehensive cancer care.

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Additional information

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