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UK consensus recommendations for clinical management of cancer risk for women with germline pathogenic variants in cancer predisposition genes: *RAD51C*, *RAD51D*, *BRIP1* and *PALB2*

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ABSTRACT

Germline pathogenic variants (GPVs) in the cancer predisposition genes *BRCA1*, *BRCA2*, *MLH1*, *MSH2*, *MSH6*, *BRIP1*, *PALB2*, *RAD51D* and *RAD51C* are identified in approximately 15% of patients with ovarian cancer (OC). While there are clear guidelines around clinical management of cancer risk in patients with GPV in *BRCA1*, *BRCA2*, *MLH1*, *MSH2* and *MSH6*, there are few guidelines on how to manage the more moderate OC risk in patients with GPV in *BRIP1*, *PALB2*, *RAD51D* and *RAD51C*, with clinical questions about appropriateness and timing of risk-reducing gynaecological surgery. Furthermore, while recognition of *RAD51C* and *RAD51D* as OC predisposition genes has been established for several years, an association with breast cancer (BC) has only more recently been described and clinical management of this risk has been unclear. With expansion of genetic testing of these genes to all patients with non-mucinous OC, new data on BC risk and improved estimates of OC risk, the UK Cancer Genetics Group and CanGene-CanVar project convened a 2-day meeting to reach a national consensus on clinical management of *BRIP1*, *PALB2*, *RAD51D* and *RAD51C* carriers in clinical practice. In this paper, we present a summary of the processes used to reach and agree on a consensus, as well as the key recommendations from the meeting.

BACKGROUND

Tube-ovarian cancer (OC) is the sixth most common cancer in women in the UK with over 7500 new diagnoses per year.¹ The risk of OC in first-degree relatives of patients with OC has been estimated to be threefold greater compared with the population risk. Most of this excess familial risk is due to germline pathogenic variants (GPVs) in the cancer predisposition genes *BRCA1* and *BRCA2*, which are associated with high lifetime risks of breast cancer (BC) and OC.² However, other genes which have lower lifetime risks of OC have been identified. These include *MLH1*, *MSH2*, *MSH6*, *BRIP1*,

PALB2, *RAD51D* and *RAD51C* as well as common, low-risk OC genetic susceptibility variants identified through genome-wide association studies.^{3,4}

While guidelines around clinical management of cancer risk in patients with GPV in *BRCA1* and *BRCA2*^{5,6} and the mismatch repair genes; *MLH1*, *MSH2*, *MSH6*⁷ have been published, there are few guidelines on how to manage patients with GPV in genes associated with more moderate risks of OC: *BRIP1*, *PALB2*, *RAD51D* and *RAD51C*.

Population-based studies of genetic testing in patients with BC or OC suggest that GPV in these genes are present in a smaller but clinically important number of patients compared with GPV in *BRCA1* and *BRCA2*. Detection rates in a large population-based study of patients with BC were 274/47 522 (0.58%) for *PALB2*, 54/47 522 (0.11%) for *RAD51C*, 51/47 522 (0.11%) for *RAD51D* and 86/47 522 (0.18%) for *BRIP1*, compared with 515/48 826 (1.05%) for *BRCA1* and 754/48 826 (1.54%) for *BRCA2*.⁸ For a population-based series of patients with OC undergoing genetic testing in two states in the USA, detection rates were 0.4% (95% CI 0.11% to 1.0%), 0.58% (95% CI 0.19% to 1.3%), 0.48% (95% CI 0.13% to 1.2%) and 0.92% (95% CI 0.4% to 1.8%) for *PALB2*, *RAD51C*, *RAD51D* and *BRIP1*, compared with 8.7% (95% CI 7.5% to 10.1%) and 5.8% (5% CI 4.7% to 6.9%) for *BRCA1* and *BRCA2*.⁹ In a smaller series of 303 patients with high-grade non-mucinous OC tested through the North East London Cancer Network, the prevalence of *RAD51C*, *RAD51D* and *BRIP1* carriers was slightly higher: 0.7%, 1.0% and 0.7% for *RAD51C*, *RAD51D* and *BRIP1*, respectively, but still lower than the prevalence of *BRCA1* and *BRCA2* carriers (11% and 4.68%), respectively.¹⁰

There are controversies about both the appropriateness and timing of risk-reducing gynaecological surgery and also BC risk associated with GPV in these genes. International guidelines for management of individuals with *PALB2* GPV have recently been published,¹¹ and *PALB2* carriers in the UK undergo breast screening via the National



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Table 1 Summary of key studies assessing association of BC risk in *RAD51C* and *RAD51D* GPV carriers (modified and updated from Yang *et al*¹⁶)

Study	Cases	Controls	RR (95% CI)	
			<i>RAD51C</i>	<i>RAD51D</i>
Dorling <i>et al</i> ⁶	48 826 population based	50 703	OR 1.93 (1.20 to 3.11)	1.8 (1.11 to 2.93)
Hu <i>et al</i> ¹⁸	32 247 population based	32 544	1.20 (0.75 to 1.93)*	1.72 (0.88 to 3.51)*
Yang <i>et al</i> ¹⁶	6178 families, 125 with <i>RAD51C</i> GPV, and 6690 families, 60 with <i>RAD51D</i> GPV	–	1.99 (1.39 to 2.85)	1.83 (1.24 to 2.72)
Li <i>et al</i> ¹⁶	3080 with BC/EOC(Epithelial ovarian cancer)	4840	8.7 (1.9 to 80.5)	Not applicable
Susynska <i>et al</i> ⁷⁷	Meta-analysis of published estimates	–	1.13 (0.88 to 1.44)	1.25 (0.9 to 1.75)
Castera <i>et al</i> ⁷⁸	5131 with family history (FH) of BC/EOC	571	1.92 (0.71 to 3.85)	2.42 (0.36 to 7.39)
Huake <i>et al</i> ⁷⁹	5589 eligible for mutation screening	ExAC (–27K)	OR: 1.29 to 5.91	3.04 (0.99 to 9.30)
Couch <i>et al</i> ⁸⁰	38 326 eligible for mutation screening	ExAC (–27K)	0.78 (0.47 to 1.37)	3.07 (1.21 to 7.88)
Slavin <i>et al</i> ⁸¹	2135 with BC/EOC FH	ExAC (–27K)	0.39 (0.02 to 2.41)	8.33 (2.2 to 30.5)
Loveday <i>et al</i> ⁸²	1132 families with BC/EOC FH	–	0.91 (0.45 to 1.86)	NA
Loveday <i>et al</i> ¹⁷	911 families with BC/EOC FH	–	NA	1.32 (0.58 to 2.96)

*When assessing ER-negative BC cases (n=3805), the risk association was stronger *RAD51C* 2.19 (0.97–4.49) and *RAD51D* 3.93 (1.40–10.29). BC, breast cancer; GPV, germline pathogenic variant; NA, not applicable.

Health Service (NHS) Very High Risk (VHR) Breast Screening programme.¹² While some guidelines have addressed management of OC in individuals with *RAD51C*, *RAD51D* or *BRIP1* GPV,¹³ there are no UK guidelines available for the management of all cancer risk in this patient group.

A previous UK Cancer Genetics Group (UKCGG) Consensus meeting held in 2018 agreed that *PALB2* should be included on a BC predisposition gene panel and *RAD51C*, *RAD51D* and *BRIP1* on an OC predisposition gene panel.¹⁴ The national genomic test directory (NHS England) was first published in August 2020 and included *PALB2* in the panel for inherited BC and OC (R208) and *RAD51C*, *RAD51D* and *BRIP1* in the inherited familial OC panel (R207). The test directory is updated each year with input from the clinical and scientific community incorporating changes to both eligibility criteria and genes on a panel (<https://www.england.nhs.uk/wp-content/uploads/2018/08/Rare-and-inherited-disease-eligibility-criteria-version-3.1-August-2022.pdf>). Similar panel testing is available in Wales, Scotland and Northern Ireland. Recently, evidence substantiating the role of *RAD51C* and *RAD51D* in BC predisposition was published,⁸ which suggests further amendments to the test directory may be required (see table 1 for a summary of studies assessing BC risk for *RAD51C* and *RAD51D*).

On the morning of Thursday, 30 September and the morning of Friday, 1 October 2021, a virtual consensus meeting was hosted as a collaboration between CanGene-CanVar (Cancer Research UK-funded catalyst project) and the UKCGG. The aim of the meeting was to develop guidelines around clinical management of patients with GPV in moderate-risk OC genes *PALB2*, *RAD51C*, *RAD51D* and *BRIP1* (henceforth referred to as carriers) to clarify surveillance and risk-reducing options.

CONSENSUS MEETING

The meeting was held virtually via Zoom and was moderated by HH, FL, AK and MT on behalf of CanGene-CanVar and UKCGG. There were 65 attendees: clinical geneticists (31), genetic counsellors (9), gynaecologists (10), menopause specialists (1), clinical nurse specialists (1), breast surgeons and oncologists (5), radiologists (2), clinical scientists (1), primary care (1) and patient representatives (4). There was representation from the four devolved nations of the UK. The meeting also included commissioners (individuals involved in funding decisions at both NHSE (National Health Service England) and the devolved nations), although they did not take part in clinical discussions.

Prior to the meeting, a review of relevant literature concerning GPV in *PALB2*, *RAD51C*, *RAD51D* and *BRIP1* was undertaken, and a background document circulated summarising published evidence for associated cancers, cancer risks, surveillance and risk-reducing surgery (background document available via <https://www.ukcgg.org/information-education/ukcgg-consensus-meetings/>). Participants were also asked to complete a survey (via SurveyMonkey) prior to the meeting, assessing opinions on genetic testing for GPV in moderate-risk OC genes, surveillance that should be offered to carriers and options around risk-reducing surgery.

The two sessions of the meeting were divided into consideration of BC risks and management on the first morning and OC risks and management on the second morning. At the start of each session, the results from the premeeting survey were presented followed by short lectures covering risk assessment, surveillance and risk-reducing surgery by expert speakers. Following these lectures, open discussions were held around specific statements (as discussed further) to inform the clinical recommendations. Voting on each statement was undertaken using Slido, which allows real-time voting online. Statements were displayed and five options for answers from ‘strongly agree’ to ‘strongly disagree’ were proposed. Within this process, it is important to set a consensus level at the beginning.¹⁵ A consensus was taken as 80% of respondents agreeing (voting agree and strongly agree). Alternative statements were debated until consensus was reached. The numbers of participants voting for each statement varied, depending on the expertise of the attendees. Notes were taken throughout the meeting and a draft document of the meeting outcomes was written and edited by FL and HH and then circulated via the core group for further input and agreement.

Question 1: should *RAD51C* and *RAD51D* be included on a BC predisposition panel?

GPV in both *RAD51C* and *RAD51D* has been identified in BC and OC families.^{16 17} For both genes, the association appears to be strongest with triple negative or ER (estrogen receptor) -negative BC.^{8 18} The largest study to date¹⁶ analysed data from 125 families with GPV in *RAD51C* and 60 families with *RAD51D*. This reported a relative risk (RR) of BC of 1.99 (95% CI 1.39 to 2.85) and 1.83 (95% CI 1.24 to 2.72), respectively. A large case-control analysis of 113 000 women via the Breast Cancer Association Consortium (BCAC)⁸ demonstrated similar levels of BC

risk with an OR of 1.93 (95% CI 1.2 to 3.11) for *RAD51C* and 1.8 (95% CI 1.11 to 2.93) for *RAD51D*. These risks are associated with truncating GPV in these genes. An association was not demonstrated for missense variants in either *RAD51C* (OR=0.9, 95% CI 0.76 to 1.14; $p=0.49$) or *RAD51D* (OR=1.05, 95% CI 0.86 to 1.27; $p=0.64$).⁸

The cumulative lifetime risk of BC associated with truncating GPV in these genes is approximately 20% for both genes.^{8 16} However, lifetime BC risk can be significantly modified by family history with a risk as high as 44%–46% for carriers, with two first-degree relatives diagnosed with BC.¹⁶

Poll statement: *RAD51C* and *RAD51D* should be included on a BC predisposition panel.

Poll results: 33% strongly agree; 67% agree (100% consensus, $n=48$).

Recommendation: consensus reached to include *RAD51C* and *RAD51D* on a BC predisposition panel.

Question 2: should *BRIP1* be included on a BC predisposition panel?

In 2006 *BRIP1* was reported as a low-penetrance BC gene.¹⁹ However, subsequent larger and more comprehensive studies have suggested that there is not a significant association with BC predisposition.^{18 20 21} The recent BCAC study did not identify an association between truncating variants in *BRIP1* and BC risk (OR=1.11, 95% CI 0.80 to 1.53; $p=0.54$),⁸ and it is now widely considered that *BRIP1* is not a BC predisposition gene.

Poll statement: *BRIP1* should not be included on a BC predisposition panel.

Poll results: 36% strongly agree; 58% agree (94% consensus, $n=50$).

Recommendation: consensus reached that *BRIP1* should not be included on a BC predisposition panel.

Question 3: should the genes on a germline BC predisposition panel be the same whether the test is requested from mainstream specialty or clinical genetics?

Traditionally, diagnostic genetic testing has taken place within regional genetic services with the likelihood of identification of a GPV in a cancer predisposition gene calculated based on the family history of cancer. The main driver of testing was to facilitate predictive testing and subsequent surveillance and risk reduction strategies for unaffected family members. However, it is known that a high proportion *BRCA1* and *BRCA2* carriers do not have a significant family history of BC or OC.²² More recent developments in personalised cancer management for individuals with GPV have resulted in lowered thresholds for germline testing with increasing numbers of individuals now eligible. As a result, mainstreaming testing (genetic testing through a non-genetics specialty at the time of new cancer diagnosis) has been widely adopted. Predictive genetic testing in an unaffected family member for a known familial GPV in a cancer predisposition gene is undertaken through Clinical Genetics, which can provide pretest counselling and assessment of residual risk based on family history and other risk factors if an individual has not inherited a familial variant.

There have been a number of studies assessing mainstreaming pathways both from the UK^{23 24} and internationally.^{25 26} While there have been some initial concerns from non-genetic specialists around offering genomic testing,²⁷ many studies have found that the mainstreaming of testing is acceptable to both patients and healthcare professionals.^{10 28–31} The group considered whether the addition of lower-risk genes to a BC predisposition

panel would be of concern to mainstreaming clinicians due to unfamiliarity or uncertainty about associated cancer risks. This was weighed against the practicality of having multiple different panels, along with the consideration that all patients identified with a GPV would be referred into a genetics service for detailed discussions regarding cancer risk and surveillance/risk-reducing strategies alongside discussion of cascade testing for other at-risk family members.

Poll statement: Genes on a BC predisposition panel should be the same whether requested from mainstream specialty or clinical genetics.

Poll results: 24% strongly agree; 67% agree (91% consensus, $n=54$).

Recommendation: consensus reached that genes on a BC predisposition panel should be the same, whether requested from mainstream specialty or clinical genetics.

Question 4: what BC surveillance should be offered to patients with a GPV in *RAD51C* or *RAD51D*?

While Yang *et al*¹⁶ estimated lifetime risks of BC of 21% and 20% for *RAD51C* and *RAD51D* carriers, respectively, these risks apply for female carriers without a significant family history. Their study also demonstrated that family history may modify this risk. With two first-degree relatives affected with BC, these lifetime risks increase to 44% and 40%, respectively.¹⁷ Similarly, the BCAC study suggested higher OR estimates when the comparison was made between cases with BC family history and controls. One of the more commonly used risk algorithms, CanRisk (www.canrisk.org), incorporates both family history and carrier status into individual risk assessments, and this or similar models should be used to provide an individualised risk assessment. While clinical judgement can also be used to assess the extent of family history, CanRisk provides a more detailed risk assessment and estimated 5-year, 10-year and lifetime risks, which are helpful in counselling patients and in shared decision making. The CanRisk tool can also include questionnaire-based factors (eg, hormonal and lifestyle factors), polygenic risk scores and mammographic density, although the latter two are not currently routinely assessed in standard clinical practice.

Poll statement: Breast surveillance for *RAD51C* and *RAD51D* carriers should be based on an individual risk assessment.

Poll results: 40% strongly agree; 60% agree (100% consensus, $n=53$).

Recommendation: consensus reached that recommendations for breast surveillance in carriers of GPVs in *RAD51C* and *RAD51D* should be based on an individual risk assessment.

Question 5: what breast surveillance should be offered to *RAD51C* and *RAD51D* carriers with a lifetime BC risk of 17%–30% (National Institute for Health and Care Excellence (NICE) moderate-risk category)? and question 6: what breast surveillance should be offered to *RAD51C* and *RAD51D* carriers with a lifetime BC risk of >30% but <40% (NICE high-risk category)

NICE guidelines on familial BC (CG164)⁶ stratify the level of lifetime BC risk at which mammography should be offered outside of the NHS Population Breast Screening Programme (NHSBSP). Individuals are classified as having a moderate lifetime risk of BC (as opposed to an average risk) when the risk is 17%–30% or the 10-year risk is 3%–8% aged 40–50 years. The NICE guidelines for patients at moderate risk suggest annual mammography between the ages of 40 years and 49 years followed by entry into the NHS Breast Screening Programme (mammography every 3

years from age 50). These guidelines also suggest that patients with a lifetime risk of BC between 30% and 40% (10-year risk of 8%–12% aged 40–50 years) (high risk) should undergo annual mammography until the age of 59 years and then revert to population screening.

As previously mentioned, BC risk of *RAD51C* and *RAD51D* carriers can be modified by family history and other BC risk factors. Based on lifetime risk calculations undertaken in the CanRisk web tool which incorporates BOADICEA V.6 (www.canrisk.org) and considering the multifactorial model (including lifestyle/hormonal risk factors, mammographic density and polygenic risk scores), carriers can be classified into different risk categories. For an 'average' 20-year-old woman with a *RAD51C* GPV (without considering cancer family history), based on the multifactorial model, approximately 38% of carriers would fall into a population risk category; 43% of carriers would fall into a moderate-risk category; and 19% would fall into a high-risk category. However, for a 20-year-old *RAD51C* carrier and a mother affected with BC at age 50, based on the multifactorial model, 15% of carriers would fall into a population risk category; 42% of carriers would fall into a moderate-risk category; and 43% would fall into a high-risk category.³²

It was noted that, in some areas of the UK, despite NICE guidelines, there remains patchy provision of moderate-risk breast screening³³ and that access to this needs to be improved, as well as more standardised quality of reporting, as this screening currently lies outside the NHSBSP.

It was recognised that, at present, risk assessment is based predominantly on family history, but that other factors such as mammographic density and polygenic risk score could also modify risk and consequently recommendations for surveillance. The group commented that future work should focus on developing and implementing new clinical pathways that incorporate these additional risk factors.

Poll statement: *RAD51C* and *RAD51D* carriers with a lifetime BC risk of 17%–30% should be offered moderate-risk surveillance according to NICE guidelines.

Poll results: 27% strongly agree; 71% agree (98% consensus, n=52).

Recommendation: consensus reached that *RAD51C* and *RAD51D* carriers with a lifetime BC risk of 17%–30% should be offered moderate-risk surveillance: annual mammograms 40–49 years then NHSBSP.

Poll statement: *RAD51C* and *RAD51D* carriers with a lifetime BC risk of >30% but <40% should be offered high-risk surveillance according to NICE guidelines.

Poll results: 20% strongly agree; 65% agree (85% consensus, n=49).

Recommendations: consensus reached that *RAD51C* and *RAD51D* carriers with a lifetime BC risk of >30% but <40% should be offered high-risk surveillance annual mammograms 40–59 years then NHSBSP.

Question 7: what breast surveillance should be offered to *RAD51C* and *RAD51D* carriers with a lifetime BC risk of 40% or greater?

The NHS VHR screening programme offers a combination of annual mammography and MRI screening to patients at VHR of BC between the ages of 25–30 years and 70 years.¹² This is defined as 'women with a lifetime risk of 40% or greater due to a specific genetic abnormality in the woman or her family'. To access the VHR screening programme, an individualised risk assessment using an NHS endorsed computer risk modelling

software program such as CanRisk needs to be undertaken to demonstrate that 10-year BC risks are greater than 8% between the ages of 25 years and 29 years, 8% between the ages of 30 years and 39 years, or 12% between the ages of 40 years and 49 years.

While most *RAD51C* and *RAD51D* carriers are unlikely to reach this level of BC risk, a small number of patients may reach this level of risk based on the strength of their family history and/or other modifying factors.¹⁶

Poll statement: *RAD51C* and *RAD51D* carriers with a lifetime BC risk of ≥40% should be referred to the VHR Breast Screening programme.

Poll results: 23% strongly agree; 75% agree (98% consensus, n=48).

Recommendations: consensus reached that *RAD51C* and *RAD51D* carriers with a lifetime BC risk of ≥40% should be referred to the NHS VHR Breast Screening programme at the appropriate age following an individualised risk assessment.

Question 8: when should risk-reducing mastectomy (RRM) be discussed with *RAD51C* and *RAD51D* carriers?

NICE guidelines include RRM as part of the pathway for managing patients with a VHR of BC. There is a strong body of evidence demonstrating a greater than 90% risk reduction associated with RRM in patients with GPV in *BRCA1* and *BRCA2*.³⁴ There is also emerging evidence that surgery will increase survival.^{35–37} However, no formal study of *RAD51D* or *RAD51C* carriers has been undertaken. The group discussed that as for patients with a strong family history of BC who meet a 30% or greater lifetime risk of BC, discussion of the option of RRM with a patient is appropriate. However, discussion with the patient should be based on individual circumstance and shared decision making. Given the lack of studies for *RAD51C* and *RAD51D* carriers, detailed counselling with patients should include, but not be restricted to, individualised cancer risk assessment, personal circumstance and preferences of the counsellee. Non-genetic risk factors such as dense breast tissue, hormonal/lifestyle modifiers and other pre-existing medical conditions should also be considered. Importantly, age-specific risks and 5 and 10 year risks, should be communicated to the patient to help in their decision making.

It was noted that some patients may fall into a level of risk where RRM, but not surveillance within the VHR surveillance programme, is offered (lifetime risk 30%–39% inclusive). In this situation, particular consideration should be paid to detailed discussion of age-specific risks.

Poll statement: RRM should be discussed with *RAD51C* and *RAD51D* carriers with a lifetime BC risk of ≥30%.

Poll results: 16% strongly agree, 78% agree (94% consensus, n=50).

Recommendation: consensus reached that RRM should be discussed with *RAD51C* and *RAD51D* carriers with a lifetime BC risk of ≥30%, in conjunction with an individualised risk assessment, appropriate counselling and shared decision making.

Question 9: should OC surveillance be offered to *RAD51C*, *RAD51D*, *BRIP1* and *PALB2* carriers?

Currently, population screening for OC is not recommended in the UK due to lack of evidence of a mortality benefit.³⁸ While there have been several studies assessing screening in those at increased risk of OC (carriers of GPV in *BRCA1* and *BRCA2*), these have also not demonstrated a clear utility. The UK FOCSS

(UK Familial Ovarian Cancer Screening Study) reported that risk of ovarian cancer algorithm (ROCA)-based screening (screening with CA-125, interpreted using the ROCA and transvaginal sonography (TVS)) demonstrated a stage shift for patients at high risk of OC. However, it remains unknown whether this surveillance would improve survival in screened high-risk patients.³⁹ The recently published Avoiding Late Diagnosis of Ovarian Cancer (ALDO) project, evaluated the utility of ROCA in *BRCA1* and *BRCA2* carriers (not available at the time of the consensus meeting), confirmed previous research that although this approach cannot prevent ovarian cancer, in women who defer RRSO, it may facilitate detection of ovarian cancer at earlier stages, resulting in less complex surgery and reduced need for pre-op chemotherapy.⁴⁰ Recent guidelines on the management of individuals with *PALB2* GPV did not recommend OC surveillance.¹¹ Studies specifically assessing surveillance in individuals with GPV in *RAD51C*, *RAD51D* or *BRIP1* have not yet been undertaken. Therefore, currently, there are no national recommendations for OC surveillance for patients at increased risk based on family history and/or genetic status, and currently, no test has been shown to detect the majority of high-grade serous OC prior to metastatic disease, either in a population or high-risk setting. General population randomised controlled trials have not shown a mortality benefit,³⁸ and there is currently no national OC screening programme.

Poll statement: OC surveillance should not routinely be offered to *RAD51C*, *RAD51D*, *BRIP1* and *PALB2* carriers.

Poll results: 13% strongly agree; 70% agree (83% consensus, n=47).

Recommendation: consensus reached that OC surveillance should not routinely be offered to *RAD51C*, *RAD51D*, *BRIP1* and *PALB2* carriers.

Question 10: should OC surveillance be offered to *RAD51C*, *RAD51D*, *BRIP1* and *PALB2* carriers within a research study?

Given the lack of specific studies of the utility of OC surveillance in *RAD51C*, *RAD51D*, *BRIP1* and *PALB2* carriers, the group discussed whether surveillance could or should be offered within the context of a research study. While the majority of the group agreed with this approach, there was also concern that if OC surveillance in *BRCA1* and *BRCA2* carriers was supported in the future, then it would be difficult not to extend this to moderate-risk gene carriers, as OC in *RAD51C*, *RAD51D*, *BRIP1* and *PALB2* carriers is similarly, most likely to be high-grade serous OC, compared with other OC pathologies.⁴¹ While there is no formal analysis, it is likely that the cost per OC case detected would be higher for the moderate-risk genes compared with *BRCA* carriers, given the overall incidence of OC is lower. However, the absolute cost of adding *RAD51C*, *RAD51D*, *BRIP1* and *PALB2* carriers to any pre-existing high-risk programme would be low due to relatively small numbers.⁴¹ Overall, there was consensus that surveillance should only be offered in the context of a research study at present, but that this could be reviewed if national recommendations in the future support surveillance in a high-risk population.

Poll statement: OC surveillance should only be offered to *RAD51C*, *RAD51D*, *BRIP1* and *PALB2* carriers within an ethically approved research study.

Poll results: 20% strongly agree; 76% agree (96% consensus, n=47).

Recommendation: consensus reached that OC surveillance should only be offered to *RAD51C*, *RAD51D*, *BRIP1* and *PALB2* carriers within the context of a research study.

Question 11: what should the lifetime risk of OC for a *RAD51C*, *RAD51D*, *BRIP1* and *PALB2* carrier be based on for clinical discussions?

Yang *et al*¹⁶ reported a RR of OC of 7.55 (95% CI 5.6 to 10.19) for *RAD51C* GPV and 7.6 (95% CI 5.61 to 10.3) for *RAD51D* GPV. The cumulative lifetime risks of OC (to age 80 years) are estimated to be 11% (95% CI 6 to 21%) and 13% (95% CI 7 to 23%), respectively. This risk has been shown to be modified by family history of OC, with a risk exceeding 30% for carriers with two first-degree relatives with OC. The risks for *RAD51C* and *RAD51D* carriers are largely conferred after the age of 50 years.

Data from a meta-analysis of carriers of GPV in *BRIP1*⁴² calculated an OR of 4.94 (95% CI 4.07 to 6.00) for OC. A further study⁴³ calculated a cumulative risk of OC to the age of 80 years of 5.8% (95% CI 3.6 to 9.1%). This study gave a larger range of OR for risks which may be indicative of the influence of family history of OC on risk.

Presence of a GPV in *PALB2* was demonstrated in a study of 852 carriers⁴⁴ to give an RR of OC of 2.91 (95% CI 1.4 to 6.04). Age-specific risks were then calculated with the lifetime risk to 80 years being quoted as 5% (95% CI 2% to 10%). A further study of risks estimated the cumulative lifetime risk to age 80 years to be 3.2% (95% CI 1.8% to 5.7%).⁴⁵ As with *RAD51C* and *RAD51D*, the risk of OC associated with GPV in *PALB2* appears to be modified by family history of OC. For example, female carriers with a mother and sister with OC diagnosed at 50 years have a lifetime risk of 16% (95% CI 8% to 28%).⁴⁴

In addition to family history, it is recognised that other hormonal and lifestyle factors can modify OC risk, including use of oral contraception, hormonal replacement therapy (HRT), parity, body mass index, tubal ligation and endometriosis. All these factors can be incorporated into the CanRisk model alongside family history to provide an individualised risk assessment. In addition, polygenic risk scores may also modify risk in either direction but at present are not available in routine clinical practice.

Poll statement: Lifetime risk of OC for a *RAD51C*, *RAD51D*, *BRIP1* and *PALB2* carrier should be based on an individualised risk assessment taking family history into consideration.

Poll results: 35% strongly agree; 63% agree (98% consensus, n=51).

Recommendation: consensus reached that lifetime risk of OC of a *RAD51C*, *RAD51D*, *BRIP1* and *PALB2* carrier should be based on an individualised risk assessment taking family history into consideration.

Question 12: when discussing risk-reducing bilateral salpingo-oophorectomy (RRSO) with a patient, what risks should be considered?

The risks of OC for *RAD51D* and *RAD51D* carriers are largely conferred after the age of 50 years. The highest 10-year risk with GPV in *RAD51C* is between the ages of 50 and 60 with the highest risks between 50 and 70 years in *RAD51D* carriers (Yang, personal communication, 2021). These risks as previously described vary with family history.¹⁶ To discuss RRSO, both lifetime and 5-10 year OC risks should be considered, balancing risks versus benefits of surgery, with particular consideration to the average age of menopause in the population and individual menopause status of the patient. Other issues such as fertility and the impact of premature menopause may also affect timing of surgery and patient decision making.

For *BRIP1* carriers, the majority of OC risk also occurs after the age of 50 years. The average age of diagnosis for a *BRIP1* carrier was at age 63.8 years compared with 58 years in non-carriers.⁴³ This

Position statement

was replicated in a study assessing 222 patients with OC and *BRIP1* GPV.⁴⁶ In this study, 90% of cases occurred after the age of 50 years with a median age of 65 years. A recent study suggested that the *BRIP1* variant c.1045G>C is a higher-risk allele,⁴⁷ although this still demonstrated a mean age of diagnosis of 62.5 years.

Yang *et al*,⁴⁴ in a study of 852 female *PALB2* carriers, demonstrated that the majority of OC risk is over the age of 50 years, with an estimated cumulative risk below this age of less than 1%. This was supported by a further study⁴⁵ with a cumulative risk of OC of less than 1% under the age of 50.

Poll statement: Discussion of RRSO should consider both lifetime risk and risks of 5–10 years.

Poll results: 23% strongly agree; 68% agree (91% consensus, n=53).

Recommendation: consensus reached that discussion of RRSO should consider both lifetime risk and 5-10 year risks

Question 13: what discussions should take place when considering premenopausal RRSO?

It is recognised that in young women undergoing surgical oophorectomy, there is an impact on both morbidity and mortality.⁴⁸ The sequelae can include vasomotor symptoms, decrease in sexual function, osteoporosis, increased risk of cardiovascular disease, depression, anxiety, dementia and cognitive decline, and multimorbidity.^{49–53} While HRT is routinely recommended and a number of these outcomes are attenuated by the use of HRT, not all of the sequelae are fully mitigated by this.^{51 53–62} Other issues that need to be considered include fertility. RRSO is only recommended once childbearing is complete.

Studies have demonstrated need for information about the postsurgical effects of RRSO.^{63 64} The group considered the need for detailed discussions with patients to ensure they are aware of the potential side effects so that these can be balanced with individualised discussion of risk.

Poll statement: Discussion of premenopausal RRSO should include a full detailed discussion of OC risk versus potential sequelae of early menopause.

Poll results: 72% strongly agree, 28% agree (100 consensus, n=53).

Recommendation: consensus reached that discussion of premenopausal RRSO should include a full detailed discussion of OC risk versus potential sequelae of early menopause.

Question 14: at what level of risk should RRSO be offered?

Historically, in UK practice, a lifetime OC risk of 10% has been used as the threshold of risk for discussion of RRSO. However, prior to comprehensive risk assessment models for OC, both with and without a recognised causative GPV in a family, calculation of individualised risk has been complex. As a result, most typically, discussion of RRSO has been based on either a genetic diagnosis in a family or clinically based criteria. For example, two or more cases of OC in a family. Recent studies in the UK^{65 66} have demonstrated the cost-effectiveness of RRSO above a threshold of 4%–5% lifetime risk. These studies suggest that patients with a lifetime risk above this threshold should be offered the opportunity to discuss RRSO. However, while surgery will decrease OC risk, there are potentially long-term sequelae associated with surgical oophorectomy, as described previously.

Considering the aforementioned, the risk threshold for offering RRSO was discussed. The group considered that counselling of patients should include an individualised risk assessment with discussion of both lifetime risk and 5-10 year risks (see question 12) that takes genetic test results and family history as a minimum into the risk assessment. Discussion should also include counselling on the possible sequelae of an early menopause (see question 13).

Poll question: At what threshold of risk should RRSO be offered? (options: <5%, 5%–10% and >10%)

Poll results: 4% <5% lifetime risk, 79% 5%–10% lifetime risk, 13% over 10% lifetime risk, 4% uncertain (n=53).

Recommendation: consensus reached that RRSO should be discussed at a lifetime risk of OC of $\geq 5\%$.

Question 15: at what age should RRSO be considered for *RAD51C* carriers?

Yang *et al* demonstrated¹⁶ that the cumulative risk of OC up until 50 years was 1% (95% CI 0.6% to 2.0%) with a risk to 80 years of 11% (95% CI 6% to 21%). CanRisk data presented in the meeting by AA demonstrated that the average OC risk for a *RAD51C* carrier is ~11% to age 80 years, with only ~5% carriers falling below a lifetime risk of 5% based on the multifactorial OC risk model. When considering risk to age 50 years, the average risk is 1.1% and based on the multifactorial model, 99.6% of carriers fall below a 3% risk before age 50 years^{32 67} (table 2A). The risk-to-benefit ratio for surgery therefore changes over the age of 50 years. However, risk can also be modified by family history, and the risk classifications

Table 2 OC risk categorisation to ages 50 and 80 years for *RAD51C*, *RAD51D*, *BRIP1* and *PALB2* GPV carriers based on the multifactorial OC model⁶⁷ implemented in CanRisk (A Antoniou, personal communication 2022, and adapted from data in Lee *et al*³²)

Gene	Considering OC risk to age 80 years			Considering OC risk to age 50 years				
	Average risk to age 80 (%)*	% carriers in risk category†			Average risk to age 50 (%)*	% carriers in risk category†		
		<5% risk	5%–10% risk	>10% risk		<3% risk	3–5% risk	>5% risk
(A) For an unaffected GPV carrier, unselected due to family history of OC								
<i>RAD51D</i>	13	2	33	65	0.9	99.9	0.1	0
<i>RAD51C</i>	11	5	44	51	1.1	99.6	0.4	0
<i>BRIP1</i>	6	47	47	6	1	99.7	0.2	0
<i>PALB2</i>	5	62	35	3	0.8	100	0	0
(B) For a GPV carrier with a mother affected with OC, age 50								
<i>RAD51D</i>	23	0	98	1.9	95.4	4.4		0.2
<i>RAD51C</i>	20	0.1	94.6	2.3	89.2	10.1		0.8
<i>BRIP1</i>	11	5	49	2	91.6	7.9		0.5
<i>PALB2</i>	10	11	33	1.5	97.5	2.4		0.1

*Average risk for a GPV carrier, based only on the GPV.

†Based on the multifactorial risk model, including questionnaire-based/clinical risk factors and polygenic risk score. GPV, germline pathogenic variant; OC, ovarian cancer.

presented earlier, based on the multifactorial model, will also be family history specific (table 2B). Therefore, the group considered that an individualised risk assessment, as well as assessment of menopausal status, should be undertaken for all *RAD51C* carriers. The initial poll statement 'For *RAD51C* carriers RRSO should only rarely be considered <50 years and should include individualised risk assessment and shared decision making' was reworded following detailed group discussion to reflect the importance of these considerations to 'For *RAD51C* carriers with a 5% or greater lifetime risk, RRSO should be considered at 50. It can be considered in patients younger than 50 following individualised risk assessment including assessment of menopausal symptoms and shared decision making'.

Poll question: For *RAD51C* carriers with a 5% or greater lifetime risk, RRSO should be considered at 50. It can be considered in carriers younger than 50 following individualised risk assessment, including assessment of menopausal symptoms and shared decision making.

Poll results: 14% strongly agree; 86% agree (100% consensus, n=43).

Recommendation: consensus reached that for *RAD51C* carriers with a 5% or greater lifetime risk, RRSO should be considered at 50 years. It can be considered in carriers younger than 50 years following individualised risk assessment including assessment of menopausal symptoms and shared decision making.

Question 16: at what age should RRSO be considered for *RAD51D* carriers?

Yang *et al* estimated that the cumulative risk of OC for *RAD51D* carriers to 50 years is 0.8% (95% CI 0.5% to 2.0%) and 13% (95% CI 7% to 23%) to age 80 years. CanRisk data presented in the meeting by AA consider that the average OC risk for a *RAD51D* carrier is 13% to age 80 years. Based on the multifactorial model, 2% of *RAD51D* carriers fall below a lifetime risk of 5%. However, when considering risk to age 50 years, the average risk is 0.9%, and based on the multifactorial model, 99.9% of carriers fall below a 3% risk before age 50 years (table 2A). This risk can be influenced by family history, with risk rising with increasing number of first-degree relatives affected with OC and risk classification also dependent on cancer family history (table 2B). Like the discussions for *RAD51C* carriers, the group felt that the age at which to consider RRSO should reflect both an individualised risk assessment and also menopausal status, and rewording of the poll question for voting was undertaken in real time during the meeting to reflect this.

Poll question: For *RAD51D* carriers with a 5% or greater lifetime risk, RRSO should be considered at 50. It can be considered in carriers younger than 50 following individualised risk assessment including assessment of menopausal symptoms and shared decision making.

Poll results: 14% strongly agree; 86% agree (100% consensus, n=44).

Recommendations: consensus reached that for *RAD51D* carriers with a 5% or greater lifetime risk, RRSO should be considered at 50 years. It can be considered in carriers younger than 50 years following individualised risk assessment, including assessment of menopausal symptoms and shared decision making.

Question 17: at what age should RRSO be considered for *BRIP1* carriers?

The cumulative OC risk associated with pathogenic variants in *BRIP1* is 5.8% (95% CI 3.6% to 9.1%). It would appear that the risk is

highest after the age of 50 years, with the average diagnosis being at 63 years.⁴³ A further study of *BRIP1* carriers demonstrated that 90% developed OC after the age of 50, with a median age of diagnosis of 65 years.⁴⁶ CanRisk data presented in the meeting model the average OC risk for a *BRIP1* carrier to be ~6% to age 80 years, and based on the multifactorial model, 47% carriers fall below a lifetime risk of 5%. When considering risk to age 50 years, the average risk is 1.0%, with 99.7% of carriers falling below a 3% risk before age 50 years when using the multifactorial model (table 2A). Again, reflecting the previous discussions for *RAD51C* and *RAD51D* carriers, the group felt that the age at which to consider RRSO for *BRIP1* carriers should include both an individualised risk assessment and also menopausal status, and rewording of the poll question for voting was undertaken in real time during the meeting to reflect this. It was also noted that unlike *RAD51C* and *RAD51D*, many *BRIP1* carriers, in the absence of OC family history, may not reach a lifetime OC risk of 5%, and calculation of an individualised risk assessment was therefore fundamental to patient discussions (table 2A,B).

Poll question: For *BRIP1* carriers with a 5% or greater lifetime risk, RRSO should be considered at 50. It can be considered in carriers younger than 50 following individualised risk assessment, including assessment of menopausal symptoms and shared decision making.

Poll results: 9% strongly agree; 91% agree (100% consensus, n=43).

Recommendation: consensus reached that for *BRIP1* carriers with a 5% or greater lifetime risk, RRSO should be considered at 50 years. It can be considered in carriers younger than 50 years following individualised risk assessment, including assessment of menopausal symptoms and shared decision making.

Question 18: at what age should RRSO be considered for *PALB2* carriers?

A study of 852 *PALB2* carriers calculated age-specific OC risks. It was estimated that there is 0.6% risk (95% CI 0.3% to 1.0%) to 50 years and 5% (95% CI 2% to 10%) to age 80 years.⁴⁴ A further study suggested the risk is 3.2% until 80 years.⁴⁵ Family history can modify this risk. Given low cumulative risk under the age of 50 years, recent guidelines from the American College of Medical Genetics and Genomics suggested that RRSO be discussed only from 50 years onwards.¹¹ CanRisk data presented in the meeting model the average OC risk for a *PALB2* carrier to be ~5% to age 80 years, with many *PALB2* (62%) carriers falling below a lifetime risk of 5%, based on the multifactorial model and absence of OC family history (table 2A). When considering risk to age 50 years, the average risk is 0.8%, with all carriers (without family history of OC) falling below a 3% risk before age 50 years (table 2A). The group felt, in general, more cautious about recommendations for RRSO for *PALB2* carriers, compared with *RAD51C* and *RAD51D*. The importance of individualised risk assessment was emphasised, given that only a small number of *PALB2* carriers, without OC family history are likely to reach a lifetime OC risk of 5%, but if there is a family history of OC, a larger proportion of carriers will reach a lifetime risk of 5% (table 2A,B).

Poll question: For *PALB2* carriers with a 5% or greater lifetime risk, RRSO should be considered at 50. It can be considered in carriers younger than 50 following individualised risk assessment, including assessment of menopausal symptoms and shared decision making.

Poll results: 12% strongly agree; 86% agree (98% consensus, n=42).

Box 1 Summary of clinical recommendations for genes discussed in consensus meeting

Recommendations for RAD51C and RAD51D carriers

- ⇒ *RAD51C* and *RAD51D* should be included on a BC predisposition gene panel.
- ⇒ Breast surveillance for *RAD51C* and *RAD51D* carriers should be based on an individualised risk assessment.
 - ⇒ Carriers with a lifetime BC risk of 17%–30% should be offered moderate-risk breast surveillance: annual mammograms 40–49 then NHSBSP.
 - ⇒ Carriers with a lifetime BC risk of >30% but <40% should be offered high-risk breast surveillance annual mammograms 40–59 years then NHSBSP.
 - ⇒ Carriers with a lifetime BC risk of >40% should be referred to the VHR Breast Screening programme.
- ⇒ Risk reducing mastectomy can be discussed with *RAD51C* and *RAD51D* carriers with a lifetime BC risk of ≥30%, in conjunction with an individualised risk assessment, appropriate counselling and shared decision making.
- ⇒ OC surveillance should not routinely be offered to *RAD51C* or *RAD51D* carriers outside a research study
- ⇒ Discussion of lifetime risk of OC for *RAD51C* and *RAD51D* carriers should be based on an individualised risk assessment, considering both lifetime risk and risks of 5–10 years and taking family history into consideration.
- ⇒ For *RAD51C* and *RAD51D* carriers, with a 5% or greater lifetime risk of OC, RRSO should be considered at 50 years. It can be considered in carriers younger than 50 years following individualised risk assessment including assessment of menopausal symptoms and shared decision making.
- ⇒ Discussion of premenopausal RRSO should include a full detailed discussion of risk versus side effects due to an early menopause.
- ⇒ Risk reducing early salpingectomy with delayed oophorectomy should only be offered to *RAD51C* and *RAD51D* carriers within a research study.

Recommendations for BRIP1 carriers

- ⇒ *BRIP1* should *not* be included on a BC gene predisposition panel.
- ⇒ Breast surveillance for *BRIP1* carriers should be based on family history and not *BRIP1* carrier status.
- ⇒ OC surveillance should not routinely be offered to *BRIP1* carriers outside a research study.
- ⇒ Discussion of lifetime risk of OC for *BRIP1* carriers should be based on an individualised risk assessment, considering both lifetime risk and risks of 5–10 years and taking family history into consideration.
- ⇒ For *BRIP1* carriers with a 5% or greater lifetime risk, RRSO should be considered at 50 years. It can be considered in carriers younger than 50 years following individualised risk assessment, including assessment of menopausal symptoms and shared decision making.
- ⇒ Discussion of premenopausal RRSO should include a full detailed discussion of risk versus side effects due to an early menopause.
- ⇒ Risk reducing early salpingectomy with delayed oophorectomy should only be offered to *BRIP1* carriers within a research study.

Recommendations for PALB2 carriers

- ⇒ *PALB2* should be included on a breast cancer predisposition gene panel.

Box 1 Continued

- ⇒ Breast surveillance for *PALB2* carriers should be based on an individualised risk assessment with carriers referred to the NHSBSP VHR screening programme at age 25–30, depending on risk.
- ⇒ RRM can be discussed with *PALB2* carriers with a lifetime BC risk ≥30%, in conjunction with an individualised risk assessment, appropriate counselling and shared decision making
- ⇒ OC surveillance should not routinely be offered to *PALB2* carriers outside a research study.
- ⇒ Discussion of lifetime risk of OC for *PALB2* carriers should be based on an individualised risk assessment, considering both lifetime risk and risks of 5–10 years and taking family history into consideration.
- ⇒ For *PALB2* carriers, with a 5% or greater lifetime risk, RRSO should be considered at 50 years. It can be considered in carriers younger than 50 years following individualised risk assessment, including assessment of menopausal symptoms and shared decision making.
- ⇒ Discussion of premenopausal RRSO should include a full detailed discussion of risk versus side effects due to an early menopause.
- ⇒ Risk-reducing early salpingectomy with delayed oophorectomy should only be offered to *PALB2* carriers within a research study.

Recommendation: consensus reached that for *PALB2* carriers with a 5% or greater lifetime risk, RRSO should be considered at 50 years. It can be considered in carriers younger than 50 years following individualised risk assessment, including assessment of menopausal symptoms and shared decision making.

Question 19: should the option of risk-reducing early salpingectomy with delayed oophorectomy be considered for *RAD51C*, *RAD51D*, *BRIP1* and *PALB2* carriers?

There is now evidence that OC can arise from the fallopian tubes and that removal of the fallopian tubes may therefore decrease the risk of OC.^{68–70} This raises the possibility of using salpingectomy with delayed oophorectomy to decrease OC risk while delaying surgical menopause in carriers at increased risk of OC.

A recent study from the Netherlands assessed quality of life after risk-reducing salpingectomy versus RRSO in 577 *BRCA1* and *BRCA2* carriers. This demonstrated a better quality of life after risk-reducing salpingectomy irrespective of HRT.⁷¹ Within the UK, the PROTECTOR study is evaluating the option of risk-reducing early salpingectomy with delayed oophorectomy in patients at high risk of OC. *BRCA1*, *BRCA2*, *PALB2*, *BRIP1*, *RAD51C* and *RAD51D* carriers are eligible for this study.⁷²

While there are several ongoing studies about the acceptability of this approach, there are still no long-term data on outcomes, in particular around OC diagnoses in these cohorts. Recent reviews by Boerner *et al* and Gaba *et al* suggested that this surgery should be offered only in the context of a clinical trial.^{72 73}

Poll question: Risk-reducing early salpingectomy with delayed oophorectomy should be offered only to *RAD51C*, *RAD51D*, *BRIP1* and *PALB2* carriers within the context of a research study.

Poll results: 30% strongly agree; 63% agree (93% consensus, n=40).

Continued

Recommendation: consensus reached that risk-reducing early salpingectomy with delayed oophorectomy should currently be offered only to *RAD51C*, *RAD51D*, *BRIP1* and *PALB2* carriers within the context of a research study until further data are available.

Question 20: is there a role for OC surveillance for *RAD51C*, *RAD51D*, *BRIP1* and *PALB2* carriers who have opted not to pursue risk-reducing surgery?

The current evidence for surveillance for OC suggests that it is ineffective in the general population. The long-term follow-up results for the UKTOCS (UK Collaborative Trial of Ovarian Cancer Screening) have recently been published³⁸ and demonstrate that surveillance does not decrease deaths from ovarian or tubal cancers. The UKFOCSS study reported that ROCA-based screening (screening with CA-125, interpreted using the ROCA, and TVS) for patients at high risk of OC demonstrated a stage shift. However, it remains unknown whether this screening would improve survival in screened high-risk patients.³⁹ Previous discussion in the meeting had reached consensus that OC surveillance should only be offered to *RAD51C*, *RAD51D*, *BRIP1* and *PALB2* carriers within a research study (see question 10). The group also considered whether there was any role for surveillance for carriers who had opted not to pursue risk-reducing surgery.

Poll question: OC surveillance can be considered for *RAD51C*, *RAD51D*, *BRIP1* and *PALB2* carriers who have opted not to pursue risk-reducing surgery.

Poll results: 84% disagree; 2% strongly disagree (86% consensus, n=49).

Recommendation: Carriers of GPVs in *RAD51C*, *RAD51D*, *BRIP1* and *PALB2* should not be offered surveillance (outside the setting of a research study) even if they have opted not to pursue risk-reducing gynaecological surgery.

DISCUSSION

Clear management guidelines exist for *BRCA1* and *BRCA2* carriers with guidance around surveillance and risk-reducing surgery for BC and OC. While the contribution of GPV in *PALB2*, *BRIP1*, *RAD51C* and *RAD51D* is smaller than for *BRCA1* and *BRCA2*, the inclusion of these genes on breast and OC gene panels has resulted in the need for similar guidance around management of carriers for GPV in these genes. However, there are no guidelines setting out surveillance or encompassing all cancer risk management available in the UK. As with much of clinical genetics, clear evidence of the optimal management of affected individuals is scarce due to the rarity of the disease and requirement for very long-term follow-up studies to generate data on which to base guidelines.

A multidisciplinary workshop was therefore convened to draw on expert clinical experience. By the end of the two-session workshop, a consensus (over 80% agreement) had been obtained for a majority of recommendations for best clinical practice for carriers of GPV in *RAD51C*, *RAD51D*, *PALB2* and *BRIP1*. A summary is presented in [Box 1](#).

In summary, carriers of GPV in these genes should have a detailed discussion about their family history, individualised risk assessment and offered RRSO at the appropriate level of risk and age. *RAD51C*, *RAD51D* and *PALB2* carriers should also have an individualised risk assessment for BC and be entered into the appropriate breast screening programme.

While these guidelines suggest 'best practice' management, there are a number of issues which impact on the implementation of the guidelines, including issues around

resources and geographical differences in the delivery of care. It is suggested that *RAD51C* or *RAD51D* carriers should have enhanced BC risk screening as defined in the NICE CG164 familial BC guidelines.⁶ However, from discussions in the workshop, it is apparent that, in line with a previous publication,³³ provision of services for moderate-risk and high-risk breast surveillance varies around the UK. This needs to be addressed, not least in response to the NHS Long Term Plan (2019), which aims to increase the number of early cancer diagnosis by screening. Populations at increased risk of malignancy, such carriers of GPV in these moderate-risk genes, are populations for targeted screening, fulfilling criteria set by the National Screening Committee (<https://www.gov.uk/government/publications/evidence-review-criteria-national-screening-programmes/criteria-for-a-targeted-screening-programme>).

VHR breast screening is included within the NHSBSP for England, and there is now agreement to offer colonoscopy surveillance for individuals with Lynch syndrome within the NHS Bowel Screening programme with roll-out anticipated in 2023. One possibility is that moderate-risk and high-risk screening could be incorporated into the NHSBSP (as for VHR) so current inequities for access and reporting can be improved and standardised. However, there are issues with the funding and services offered to the devolved nations resulting in disparate provision across the UK.

The workshop has produced clear guidance around recommendations of risk levels at which to consider RRSO. However, it should be remembered that the risk estimates based on multifactorial risk models may be associated with wide CIs based on the uncertainty related to input parameters. Risk estimates are also dependent on the accuracy, quality, validity and extent of the information input into the tool. External, independent validation studies provide useful guidance on these aspects.⁷⁴ As such, discussions with patients will require explanation of the variation in risk estimations along with detailed discussions about timing of surgery and potential sequelae. This will then facilitate shared decision making with each individual patient to optimise care as per NICE guidance 197.⁷⁵ These sequelae may include menopausal symptoms, depending on the time of the surgery. Women should have access to discussions around HRT or alternative therapies when appropriate. However, it was highlighted in the workshop that while some centres of excellence exist, access to specialised menopausal care is variable around the UK.

Overall, with increased numbers of *RAD51C*, *RAD51D*, *BRIP1* and *PALB2* carriers likely to be identified imminently with updates to the national test directory within all four UK nations, we believe that these guidelines represent a framework for consistent and best practice based on the current evidence. It is likely that new relevant information will be published in the next 5 years, both from larger studies of carriers and results from studies addressing ovarian surveillance and risk-reducing surgery; therefore, regular review and updates will be required and discussions with patients should also include the potential for clinical recommendations to change over time, as and when new evidence becomes available.

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REFERENCES

- 1 Cancer Research UK. Ovarian cancer incidence statistics, 2022. Available: <https://www.cancerresearchuk.org/health-professional/cancer-statistics/statistics-by-cancer-type/ovarian-cancer/incidence> [Accessed Sep 2022].
- 2 Jervis S, Song H, Lee A, Dicks E, Tyrer J, Harrington P, Easton DF, Jacobs IJ, Pharoah PPD, Antoniou AC. Ovarian cancer familial relative risks by tumour subtypes and by known ovarian cancer genetic susceptibility variants. *J Med Genet* 2014;51:108–13.
- 3 Pietragalla A, Arcieri M, Marchetti C, Scambia G, Fagotti A. Ovarian cancer predisposition beyond BRCA1 and BRCA2 genes. *Int J Gynecol Cancer* 2020;30:1803–10.
- 4 Jones MR, Kamara D, Karlan BY, Pharoah PDP, Gayther SA. Genetic epidemiology of ovarian cancer and prospects for polygenic risk prediction. *Gynecol Oncol* 2017;147:705–13.
- 5 Ferreira MA, Gamazon ER, Al-Ejeh F, Aittomäki K, Andrusis IL, Anton-Culver H, Arason A, Arndt V, Aronson KJ, Arun BK, Asseryanis E, Azzolini J, Balmaña J, Barnes DR, Barrowdale D, Beckmann MW, Behrens S, Benitez J, Bermisheva M, Bialkowska K, Blomqvist C, Bogdanova NV, Bojesen SE, Bolla MK, Borg A, Brauch H, Brenner H, Broeks A, Burwinkel B, Caldés T, Caligo MA, Campa D, Campbell I, Canzian F, Carter J, Carter BD, Castelao JE, Chang-Claude J, Chanock SJ, Christiansen H, Chung WK, Claes KBM, Clarke CL, Couch FJ, Cox A, Cross SS, Czene K, Daly MB, de la Hoya M, Dennis J, Devilee P, Diez O, Dörk T, Dunning AM, Dwek M, Eccles DM, Ejlertsen B, Ellberg C, Engel C, Eriksson M, Fasching PA, Fletcher O, Flyger H, Friedman E, Frost D, Gabrielson M, Gago-Dominguez M, Ganz PA, Gapstur SM, Garber J, Garcia-Closas M, García-Sánchez JA, Gaudet MM, Giles GG, Glendon G, Godwin AK, Goldberg MS, Goldgar DE, González-Neira A, Greene MH, Gronwald J, Guénel P, Haiman CA, Hall P, Hamann U, He W, Heyworth J, Hogervorst FBL, Hollestelle A, Hoover RN, Hopper JL, Hulick PJ, Humphreys K, Ilyanovitch EN, Isaacs C, Jakimovska M, Jakubowska A, James PA, Janavicius R, Jankowitz RC, John EM, Johnson N, Joseph V, Karlan BY, Khushnudinova E, Kiiski JJ, Ko Y-D, Jones ME, Konstantopoulou I, Kristensen VN, Laitman Y, Lambrechts D, Lazaro C, Leslie G, Lester J, Lesueur F, Lindström S, Long J, Loud JT, Lubiński J, Makalic E, Mannermaa A, Manoochehri M, Margolin S, Maurer T, Mavroudis D, McGuffog L, Meindl A, Menon U, Michailidou K, Miller A, Montagna M, Moreno F, Moserle L, Mulligan AM, Nathanson KL, Neuhausen SL, Nevanlinna H, Nevelsteen I, Nielsen FC, Nikitina-Zake L, Nussbaum RL, Offit K, Olah E, Olopade OI, Olsson H, Osorio A, Papp J, Park-Simon T-W, Parsons MT, Pedersen IS, Peixoto A, Peterlongo P, Pharoah PDP, Plaseska-Karanfilska D, Poppe B, Presneau N, Radice P, Rantala J, Rennett G, Risch HA, Saloustros E, Sanden K, Sawyer EJ, Schmidt MK, Schmutzler RK, Sharma P, Shu X-O, Simard J, Singer CF, Soucy P, Southey MC, Spinelli JJ, Spurdle AB, Stone J, Swerdlow AJ, Tapper WJ, Taylor JA, Teixeira MR, Terry MB, Teulé A, Thomassen M, Thöne K, Thull DL, Tischkowitz M, Toland AE, Torres D, Truong T, Tung N, Vachon CM, van Asperen CJ, van den Ouweland AMW, van Rensburg EJ, Vega A, Viel A, Wang Q, Wappenschmidt B, Weitzel JN, Wendt C, Winquist R, Yang XR, Yannoukakis D, Zogas A, Kraft P, Antoniou AC, Zheng W, Easton DF, Milne RL, Beesley J, Chenevix-Trench G, EMBRACE Collaborators, GC-HBOC Study Collaborators, GEMO Study Collaborators, ABCTB Investigators, HEBON Investigators, BCFR Investigators. Genome-wide association and transcriptome studies identify target genes and risk loci for breast cancer. *Nat Commun* 2019;10:1741.
- 6 National Institute of Clinical Excellence. Familial breast cancer: classification, care and managing breast cancer and related risks in people with a family history of breast cancer (CG164)
- 7 Vasen HFA, Blanco I, Aktan-Collan K, Gopie JP, Alonso A, Aretz S, Bernstein I, Bertario L, Burn J, Capella G, Colas C, Engel C, Fraying IM, Genuardi M, Heinemann K, Hes FJ, Hodgson SV, Karagiannis JA, Lalloo F, Lindblom A, Mecklin J-P, Möller P, Myrholm T, Nagengast FM, Parc Y, Ponz de Leon M, Renkonen-Sinisalo L, Sampson JR, Stormorken A, Sijmons RH, Tejpar S, Thomas HJW, Rahner N, Wijnen JT, Järvinen HJ, Möslin G, Mallorca group. Revised guidelines for the clinical management of Lynch syndrome (HNPCC): recommendations by a group of European experts. *Gut* 2013;62:812–23.
- 8 Dorling L, Carvalho S, Allen J, González-Neira A, Luccarini C, Wahlström C, Pooley KA, Parsons MT, Fortunato C, Wang Q, Bolla MK, Dennis J, Keeman R, Alonso MR, Álvarez N, Herraez B, Fernandez V, Núñez-Torres R, Osorio A, Valcich J, Li M, Törngren T, Harrington PA, Baynes C, Conroy DM, Decker B, Fachal L, Mavaddat N, Ahearn T, Aittomäki K, Antonenkova NN, Arnold N, Arveux P, Ausems MGEM, Auvinen P, Becher H, Beckmann MW, Behrens S, Bermisheva M, Bialkowska K, Blomqvist C, Bogdanova NV, Bogdanova-Markov N, Bojesen SE, Bonanni B, Borresen-Dale A-L,

- Brauch H, Bremer M, Briceno I, Brüning T, Burwinkel B, Cameron DA, Camp NJ, Campbell A, Carracedo A, Castella JE, Cessna MH, Chanock SJ, Christiansen H, Collé JM, Cordina-Duverger E, Cornelissen S, Czene K, Dörk T, Ekici AB, Engel C, Eriksson M, Fasching PA, Figueroa J, Flyger H, Försti A, Gabrielson M, Gago-Dominguez M, Georgoulas V, Gil F, Giles GG, Glendon G, Garcia EBG, Alnæs GIG, Guénel P, Hadjisavvas A, Haerle L, Hahnen E, Hall P, Hamann U, Harkness EF, Hartikainen JM, Hartman M, He W, Heemskerk-Gerritsen BAM, Hillemanns P, Hogervorst FBL, Hollestelle A, Ho WK, Hoening MJ, Howell A, Humphreys K, Idris F, Jakubowska A, Jung A, Kapoor PM, Kerin MJ, Khushnutdinova E, Kim S-W, Ko Y-D, Kosma V-M, Kristensen VN, Kyriacou K, Lakeman IMM, Lee JW, Lee MH, Li J, Lindblom A, Lo W-Y, Loizidou MA, Lophatananon A, Lubiński J, MacLennan RJ, Madsen MJ, Mannerman A, Manoochehri M, Manoukian S, Margolin S, Martinez ME, Maurer T, Mavroudis D, McLean C, Meindl A, Mensenkamp AR, Michailidou K, Miller N, Mohd Taib NA, Muir K, Mulligan AM, Nevanlinna H, Newman WG, Nordestgaard BG, Ng P-S, Oosterwijk JC, Park SK, Park-Simon T-W, Perez JIA, Peterlongo P, Porteous DJ, Prasad C, Prokofyeva D, Radice P, Rashid MU, Rhenius V, Rookus MA, Rüdiger T, Saloustros E, Sawyer EJ, Schmutzler RK, Schneeweiss A, Schürmann P, Shah M, Sohn C, Southey MC, Surowy H, Suvanto M, Thanassitichai S, Tomlinson I, Torres D, Truong T, Tzardi M, Valova Y, van Asperen CJ, Van Dam RM, van den Ouweland AMW, van der Kolk LE, van Veen EM, Wendt C, Williams JA, Yang XR, Yoon S-Y, Zamora MP, Evans DG, de la Hoya M, Simard J, Antoniou AC, Borg Åke, Andrulis IL, Chang-Claude J, Garcia-Closas M, Chenevix-Trench G, Milne RL, Pharoah PDP, Schmidt MK, Spurdle AB, Vreeswijk MPG, Benitez J, Dunning AM, Kvist A, Teo SH, Devilee P, Easton DF, Breast Cancer Association Consortium. Breast cancer risk genes - Association analysis in more than 113,000 women. *N Engl J Med* 2021;384:428–39.
- 9 Kurian AW, Ward KC, Howlander N, Deapen D, Hamilton AS, Mariotto A, Miller D, Penberthy LS, Katz SJ. Genetic testing and results in a population-based cohort of breast cancer patients and ovarian cancer patients. *J Clin Oncol* 2019;37:1305–15.
- 10 Chandrasekaran D, Sobocan M, Blyuss O, Miller RE, Evans O, Crusz SM, Mills-Baldock T, Sun L, Hammond RFL, Gaba F, Jenkins LA, Ahmed M, Kumar A, Jeyarajah A, Lawrence AC, Brockbank E, Phadnis S, Quigley M, El Khouly F, Wuntakal R, Faruqi A, Trevisan G, Casey L, Burghel GJ, Schlecht H, Bulman M, Smith P, Bowers NL, Legood R, Lockley M, Wallace A, Singh N, Evans DG, Manchanda R. Implementation of multigene germline and parallel somatic genetic testing in epithelial ovarian cancer: SIGNPOST study. *Cancers* 2021;13. doi:10.3390/cancers13174344. [Epub ahead of print: 27 08 2021].
- 11 Tischkowitz M, Balmaña J, Foulkes WD, James P, Ngeow J, Schmutzler R, Voian N, Wick MJ, Stewart DR, Pal T. Management of individuals with germline variants in PALB2: a clinical practice resource of the American College of medical genetics and genomics (ACMG). *Genet Med* 2021;23:1416–23.
- 12 Public Health England. Tests and frequency of testing of women at very high risk - Breast Screening: very high risk women surveillance protocols.
- 13 Manchanda R, Gaba F, Talaulikar V, Pundir J, Gessler S, Davies M, Menon U, Royal College of Obstetricians and Gynaecologists. Risk-reducing salpingo-oophorectomy and the use of hormone replacement therapy below the age of natural menopause: scientific impact paper No. 66 October 2021: scientific impact paper No. 66. *BJOG* 2022;129:e16–34.
- 14 Taylor A, Brady AF, Frayling IM, Hanson H, Tischkowitz M, Turnbull C, Side L, UK Cancer Genetics Group (UK-CGG). Consensus for genes to be included on cancer panel tests offered by UK genetics services: guidelines of the UK cancer genetics group. *J Med Genet* 2018;55:372–7.
- 15 Diamond IR, Grant RC, Feldman BM, Pencharz PB, Ling SC, Moore AM, Wales PW. Defining consensus: a systematic review recommends methodologic criteria for reporting of Delphi studies. *J Clin Epidemiol* 2014;67:401–9.
- 16 Yang X, Song H, Leslie G, Engel C, Hahnen E, Auber B, Horváth J, Kast K, Niederacher D, Turnbull C, Houlston R, Hanson H, Loveday C, Dolinsky JS, LaDuca H, Ramus SJ, Menon U, Rosenthal AN, Jacobs J, Gayther SA, Dicks E, Nevanlinna H, Aittomäki K, Pelttari LM, Ehrencrona H, Borg Åke, Kvist A, Rivera B, Hansen TVO, Djursby M, Lee A, Dennis J, Bowtell DD, Traficante N, Diez O, Balmaña J, Gruber SB, Chenevix-Trench G, Investigators kConFab, Jensen A, Kjær SK, Høgdall E, Castéra L, Garber J, Janavicius R, Osorio A, Golmard L, Vega A, Couch FJ, Robson M, Gronwald J, Domchek SM, Culver JO, de la Hoya M, Easton DF, Foulkes WD, Tischkowitz M, Meindl A, Schmutzler RK, Pharoah PDP, Antoniou AC. Ovarian and breast cancer risks associated with pathogenic variants in RAD51C and RAD51D. *J Natl Cancer Inst* 2020;112:1242–50.
- 17 Loveday C, Turnbull C, Ramsay E, Hughes D, Ruark E, Frankum JR, Bowden G, Kalmyrzaev B, Warren-Perry M, Snape K, Adlard JW, Barwell J, Berg J, Brady AF, Brewer C, Brice G, Chapman C, Cook J, Davidson R, Donaldson A, Douglas F, Greenhalgh L, Henderson A, Izatt L, Kumar A, Lalloo F, Miedzybrodzka Z, Morrison PJ, Paterson J, Porteous M, Rogers MT, Shanley S, Walker L, Eccles D, Evans DG, Renwick A, Seal S, Lord CJ, Ashworth A, Reis-Filho JS, Antoniou AC, Rahman N, Breast Cancer Susceptibility Collaboration (UK). Germline mutations in RAD51D confer susceptibility to ovarian cancer. *Nat Genet* 2011;43:879–82.
- 18 Hu C, Hart SN, Gnanaolivu R, Huang H, Lee KY, Na J, Gao C, Lilyquist J, Yadav S, Boddicker NJ, Samara R, Klebba J, Ambrosone CB, Anton-Culver H, Auer P, Bandera EV, Bernstein L, Bertrand KA, Burnside ES, Carter BD, Eliassen H, Gapstur SM, Gaudet M, Haiman C, Hodge JM, Hunter DJ, Jacobs EJ, John EM, Kooperberg C, Kurian AW, Le Marchand L, Lindstrom S, Lindstrom T, Ma H, Neuhausen S, Newcomb PA, O'Brien KM, Olson JE, Ong IM, Pal T, Palmer JR, Patel AV, Reid S, Rosenberg L, Sandler DP, Scott C, Tamimi R, Taylor JA, Trentham-Dietz A, Vachon CM, Weinberg C, Yao S, Ziogas A, Weitzel JN, Goldgar DE, Domchek SM, Nathanson KL, Kraft P, Polley EC, Couch FJ. A population-based study of genes previously implicated in breast cancer. *N Engl J Med* 2021;384:440–51.
- 19 Seal S, Thompson D, Renwick A, Elliott A, Kelly P, Barfoot R, Chagtai T, Jayatilake H, Ahmed M, Spanova K, North B, McGuffog L, Evans DG, Eccles D, Easton DF, Stratton MR, Rahman N, Breast Cancer Susceptibility Collaboration (UK). Truncating mutations in the Fanconi anemia J gene BRIP1 are low-penetrance breast cancer susceptibility alleles. *Nat Genet* 2006;38:1239–41.
- 20 Couch FJ, Shimelis H, Hu C, Hart SN, Polley EC, Na J, Hallberg E, Moore R, Thomas A, Lilyquist J, Feng B, McFarland R, Pesaran T, Huether R, LaDuca H, Chao EC, Goldgar DE, Dolinsky JS. Associations between cancer predisposition testing panel genes and breast cancer. *JAMA Oncol* 2017;3:1190–90.
- 21 Weber-Lassalle N, Hauke J, Ramser J, Richters L, Grob E, Blümcke B, Gehrig A, Kahler T-A, Müller CR, Hackmann K, Honisch E, Weber-Lassalle K, Niederacher D, Borde J, Thiele H, Ernst C, Altmüller J, Neidhardt G, Nürnberg P, Klaschik K, Schroeder C, Platzer K, Volk AE, Wang-Gohrke S, Just W, Auber B, Kubisch C, Schmidt G, Horvath J, Wappenschmidt B, Engel C, Arnold N, Dworniczak B, Rhiem K, Meindl A, Schmutzler RK, Hahnen E. BRIP1 loss-of-function mutations confer high risk for familial ovarian cancer, but not familial breast cancer. *Breast Cancer Res* 2018;20:7.
- 22 Alsop K, Fereday S, Meldrum C, deFazio A, Emmanuel C, George J, Dobrovic A, Birrer MJ, Webb PM, Stewart C, Friedlander M, Fox S, Bowtell D, Mitchell G. BRCA mutation frequency and patterns of treatment response in BRCA mutation-positive women with ovarian cancer: a report from the Australian ovarian cancer study group. *J Clin Oncol* 2012;30:2654–63.
- 23 Flaum N, Morgan RD, Burghel GJ, Bulman M, Clamp AR, Hasan J, Mitchell CL, Badea D, Moon S, Hogg M, Hadjiyiannakis D, Clancy T, Schlecht H, Woodward ER, Crosbie EJ, Edmondson RJ, Wallace AJ, Jayson GC, Lalloo FI, Harkness EF, Evans DGR. Mainstreaming germline BRCA1/2 testing in non-mucinous epithelial ovarian cancer in the North West of England. *Eur J Hum Genet* 2020;28:1541–7.
- 24 Rahman B, Lanceley A, Kristeleit RS, Ledermann JA, Lockley M, McCormack M, Mould T, Side L. Mainstreamed genetic testing for women with ovarian cancer: first-year experience. *J Med Genet* 2019;56:195–8.
- 25 Buchanan AH, Rahm AK, Williams JL. Alternate service delivery models in cancer genetic counseling: a mini-review. *Front Oncol* 2016;6:120–20.
- 26 Kentwell M, Dow E, Antill Y, Wrede CD, McNally O, Higgs E, Hamilton A, Ananda S, Lindeman GJ, Scott CL. Mainstreaming cancer genetics: a model integrating germline BRCA testing into routine ovarian cancer clinics. *Gynecol Oncol* 2017;145:130–6.
- 27 Hallowell N, Wright S, Stirling D, Gourley C, Young O, Porteous M. Moving into the mainstream: healthcare professionals' views of implementing treatment focussed genetic testing in breast cancer care. *Fam Cancer* 2019;18:293–301.
- 28 Bokkers K, Zweemer RP, Koudijs MJ, Stehouwer S, Velthuisen ME, Bleiker EMA, Ausems MGE. Positive experiences of healthcare professionals with a mainstreaming approach of germline genetic testing for women with ovarian cancer. *Fam Cancer* 2022;21:295–304.
- 29 Kemp Z, Turnbull A, Yost S, Seal S, Mahamdallie S, Poyastro-Pearson E, Warren-Perry M, Eccleston A, Tan M-M, Teo SH, Turner N, Strydom A, George A, Rahman N. Evaluation of cancer-based criteria for use in mainstream BRCA1 and BRCA2 genetic testing in patients with breast cancer. *JAMA Netw Open* 2019;2:e194428–e28.
- 30 Plaskocinska I, Shipman H, Drummond J, Thompson E, Buchanan V, Newcombe B, Hodgkin C, Barter E, Ridley P, Ng R, Miller S, Dann A, Licence V, Webb H, Tan LT, Daly M, Ayers S, Rufford B, Earl H, Parkinson C, Duncan T, Jimenez-Linan M, Sagoo GS, Abbs S, Hulbert-Williams N, Pharoah P, Crawford R, Brenton JD, Tischkowitz M. New paradigms for BRCA1/BRCA2 testing in women with ovarian cancer: results of the genetic testing in epithelial ovarian cancer (GTEOC) study. *J Med Genet* 2016;53:655–61.
- 31 Meiser B, Woodward P, Gleeson M, Kentwell M, Fan HM, Antill Y, Butow PN, Boyle F, Best M, Taylor N, Bell K, Tucker K. Pilot study of an online training program to increase genetic literacy and communication skills in oncology healthcare professionals discussing BRCA1/2 genetic testing with breast and ovarian cancer patients. *Fam Cancer* 2022;21:157–66.
- 32 Lee A, Mavaddat N, Cunningham A, Carver T, Ficorella L, Archer S, Walter FM, Tischkowitz M, Roberts J, Usher-Smith J, Simard J, Schmidt MK, Devilee P, Zadnik V, Jürgens H, Mouret-Fourme E, De Pauw A, Rookus M, Mooij TM, Pharoah PP, Easton DF, Antoniou AC. Enhancing the BOADICEA cancer risk prediction model to incorporate new data on RAD51C, RAD51D, BARD1 updates to tumour pathology and cancer incidence. *J Med Genet* 2022;jmedgenet-2022-108471.
- 33 Evans DG, Edwards M, Duffy SW, Tischkowitz M, Cancer Genetics Group clinical leads. Sporadic implementation of UK familial mammographic surveillance guidelines 15 years after original publication. *Br J Cancer* 2020;122:329–32.
- 34 De Felice F, Marchetti C, Musella A, Palaia I, Perniola G, Musio D, Muzii L, Tombolini V, Benedetti Panici P. Bilateral risk-reduction mastectomy in BRCA1 and BRCA2 mutation carriers: a meta-analysis. *Ann Surg Oncol* 2015;22:2876–80.
- 35 Ludwig KK, Neuner J, Butler A, Geurts JL, Kong AL. Risk reduction and survival benefit of prophylactic surgery in BRCA mutation carriers, a systematic review. *Am J Surg* 2016;212:660–9.

- 36 Ingham SL, Sperrin M, Baildam A, Ross GL, Clayton R, Lalloo F, Buchan I, Howell A, Evans DGR. Risk-reducing surgery increases survival in BRCA1/2 mutation carriers unaffected at time of family referral. *Breast Cancer Res Treat* 2013;142:611–8.
- 37 Carbine NE, Lostumbo L, Wallace J, Ko H. Risk-reducing mastectomy for the prevention of primary breast cancer. *Cochrane Database Syst Rev* 2018;4:CD002748–CD48.
- 38 Menon U, Gentry-Maharaj A, Burnell M, Singh N, Ryan A, Karpinskyj C, Carlino G, Taylor J, Massingham SK, Raikou M, Kalsi JK, Woolas R, Manchanda R, Arora R, Casey L, Dawnay A, Dobbs S, Leeson S, Mould T, Seif MW, Sharma A, Williamson K, Liu Y, Fallowfield L, McGuire AJ, Campbell S, Skates SJ, Jacobs IJ, Parmar M. Ovarian cancer population screening and mortality after long-term follow-up in the UK Collaborative trial of ovarian cancer screening (UKCTOCS): a randomised controlled trial. *Lancet* 2021;397:2182–93.
- 39 Rosenthal AN, Fraser LSM, Philpott S, Manchanda R, Burnell M, Badman P, Hadwin R, Rizzuto I, Benjamin E, Singh N, Evans DG, Eccles DM, Ryan A, Liston R, Dawnay A, Ford J, Gunu R, Mackay J, Skates SJ, Menon U, Jacobs IJ, United Kingdom Familial Ovarian Cancer Screening Study collaborators. Evidence of stage shift in women diagnosed with ovarian cancer during phase II of the United Kingdom familial ovarian cancer screening study. *J Clin Oncol* 2017;35:1411–20.
- 40 Philpott S, Raikou M, Manchanda R, Lockley M, Singh N, Scott M, Evans DG, Adlard J, Ahmed M, Edmondson R, Woodward ER, Lamnisos A, Balega J, Brady AF, Sharma A, Izzat L, Kulkarni A, Tripathi V, Solomons JS, Hayes K, Hanson H, Snape K, Side L, Skates S, McGuire A, Rosenthal AN. The avoiding late diagnosis of ovarian cancer (ALDO) project; a pilot national surveillance programme for women with pathogenic germline variants in *BRCA1* and *BRCA2*. *J Med Genet* 2022. doi:10.1136/jmg-2022-108741. [Epub ahead of print: 01 Nov 2022].
- 41 Pavanello M, Chan IH, Ariff A, Pharoah PD, Gayther SA, Ramus SJ. Rare germline genetic variants and the risks of epithelial ovarian cancer. *Cancers* 2020;12. doi:10.3390/cancers12103046. [Epub ahead of print: 19 10 2020].
- 42 Suszynska M, Ratajska M, Kozlowski P. BRIP1, RAD51C, and RAD51D mutations are associated with high susceptibility to ovarian cancer: mutation prevalence and precise risk estimates based on a pooled analysis of ~30,000 cases. *J Ovarian Res* 2020;13:50.
- 43 Ramus SJ, Song H, Dicks E, Tyrer JP, Rosenthal AN, Intermaggio MP, Fraser L, Gentry-Maharaj A, Hayward J, Philpott S, Anderson C, Edlund CK, Conti D, Harrington P, Barrowdale D, Bowtell DD, Alsop K, Mitchell G, Cicek MS, Cunningham JM, Fridley BL, Alsop J, Jimenez-Linan M, Poblete S, Lele S, Sucheston-Campbell L, Moysich KB, Sieh W, McGuire V, Lester J, Bogdanova N, Dürst M, Hillemanns P, Odunsi K, Whittemore AS, Karlan BY, Dörk T, Goode EL, Menon U, Jacobs IJ, Antoniou AC, Pharoah PDP, Gayther SA, AOC Study Group, Ovarian Cancer Association Consortium. Germline mutations in the BRIP1, BARD1, PALB2, and NBN genes in women with ovarian cancer. *J Natl Cancer Inst* 2015;107. doi:10.1093/jnci/djv214. [Epub ahead of print: 27 08 2015].
- 44 Yang X, Leslie G, Doroszuk A, Schneider S, Allen J, Decker B, Dunning AM, Redman J, Scarth J, Plaskocinica I, Luccarini C, Shah M, Pooley K, Dorling L, Lee A, Adank MA, Adlard J, Aittomäki K, Andrulis IL, Ang P, Barwell J, Bernstein JL, Bobolis K, Borg Åke, Blomqvist C, Claes KBM, Concannon P, Cuggia A, Culver JO, Damiola F, de Pauw A, Diez O, Dolinsky JS, Domchek SM, Engel C, Evans DG, Fostira F, Garber J, Golmard L, Goode EL, Gruber SB, Hahnen E, Hake C, Heikkinen T, Hurley JE, Janavicius R, Kleibl Z, Kleiblova P, Konstantopoulou I, Kvist A, Laduca H, Lee ASG, Lesueur F, Maher ER, Mannema A, Manoukian S, McFarland R, McKinnon W, Meindl A, Metcalfe K, Mohd Taib NA, Moilanen J, Nathanson KL, Neuhausen S, Ng PS, Nguyen-Dumont T, Nielsen SM, Obermair F, Offit K, Olopade OI, Ottini L, Penkert J, Pylkäs K, Radice P, Ramus SJ, Rudaitis V, Side L, Silva-Smith R, Silvestri V, Skytte A-B, Slavin T, Soukupova J, Tondini C, Trainer AH, Unzeitig G, Usha L, van Overeem Hansen T, Whitworth J, Wood M, Yip CH, Yoon S-Y, Yussuf A, Zogopoulos G, Goldgar D, Hopper JL, Chenevix-Trench G, Pharoah P, George SHL, Balmaña J, Houdayer C, James P, El-Haffaf F, Ehrencrona H, Janatova M, Peterlongo P, Nevanlinna H, Schmutzler R, Teo S-H, Robson M, Pal T, Couch F, Weitzel JN, Elliott A, Southey M, Winquist R, Easton DF, Foulkes WD, Antoniou AC, Tischkowitz M. Cancer risks associated with Germline *PALB2* pathogenic variants: an International study of 524 families. *J Clin Oncol* 2020;38:674–85.
- 45 Song H, Dicks EM, Tyrer J, Intermaggio M, Chenevix-Trench G, Bowtell DD, Traficante N, Group A, Brenton J, Goranova T, Hosking K, Piskorz A, van Oudenhove E, Doherty J, Harris HR, Rossing MA, Duerst M, Dork T, Bogdanova NV, Modugno F, Moysich K, Odunsi K, Ness R, Karlan BY, Lester J, Jensen A, Krüger Kjaer S, Høgdall E, Campbell IG, Lázaro C, Pujara MA, Cunningham J, Vierkant R, Winham SJ, Hildebrandt M, Huff C, Li D, Wu X, Yu Y, Permut JB, Levine DA, Schildkraut JM, Riggan MJ, Berchuck A, Webb PM, Group OS, Cybulski C, Gronwald J, Jakubowska A, Lubinski J, Alsop J, Harrington P, Chan I, Menon U, Pearce CL, Wu AH, de Fazio A, Kennedy CJ, Goode E, Ramus S, Gayther S, Pharoah P. Population-based targeted sequencing of 54 candidate genes identifies *PALB2* as a susceptibility gene for high-grade serous ovarian cancer. *J Med Genet* 2021;58:305–13.
- 46 Cummings S, Roman SS, Saam J, Bernhisel R, Brown K, Lancaster JM, Usha L. Age of ovarian cancer diagnosis among BRIP1, RAD51C, and RAD51D mutation carriers identified through multi-gene panel testing. *J Ovarian Res* 2021;14:61.
- 47 Flaum N, van Veen EM, Smith O, Amico S, Newman WG, Crosbie EJ, Edmondson R, Smith MJ, Evans DG. Dominant-negative pathogenic variant BRIP1 c.1045G>C is a high-risk allele for non-mucinous epithelial ovarian cancer: a case-control study. *Clin Genet* 2022;101:48–54.
- 48 Parker WH, Feskanich D, Broder MS, Chang E, Shoupe D, Farquhar CM, Berek JS, Manson JE. Long-term mortality associated with oophorectomy compared with ovarian conservation in the nurses' health study. *Obstet Gynecol* 2013;121:709–16.
- 49 Kershaw V, Hickey I, Wylid L, Jha S. The impact of risk reducing bilateral salpingo-oophorectomy on sexual function in BRCA1/2 mutation carriers and women with Lynch syndrome: a systematic review and meta-analysis. *Eur J Obstet Gynecol Reprod Biol* 2021;265:7–17.
- 50 Shuster LT, Gostout BS, Grossardt BR, Rocca WA. Prophylactic oophorectomy in premenopausal women and long-term health. *Menopause Int* 2008;14:111–6.
- 51 Levine ME, Lu AT, Chen BH, Hernandez DG, Singleton AB, Ferrucci L, Bandinelli S, Salfati E, Manson JE, Quach A, Kusters CDJ, Kuh D, Wong A, Teschendorff AE, Widschwendter M, Ritz BR, Absher D, Assimes TL, Horvath S. Menopause accelerates biological aging. *Proc Natl Acad Sci U S A* 2016;113:9327–32.
- 52 Rocca WA, Gazzuola Rocca L, Smith CY, Grossardt BR, Faubion SS, Shuster LT, Kirkland JL, Stewart EA, Miller VM. Bilateral oophorectomy and accelerated aging: cause or effect? *J Gerontol A Biol Sci Med Sci* 2017;72:1213–7.
- 53 Rocca WA, Grossardt BR, Geda YE, Gostout BS, Bower JH, Maraganore DM, de Andrade M, Melton LJ. Long-term risk of depressive and anxiety symptoms after early bilateral oophorectomy. *Menopause* 2008;15:(Electronic):1050–9.
- 54 Rocca WA, Grossardt BR, Miller VM, Shuster LT, Brown RD. Premature menopause or early menopause and risk of ischemic stroke. *Menopause* 2012;19:272–7.
- 55 Kattah AG, Smith CY, Gazzuola Rocca L, Grossardt BR, Garovic VD, Rocca WA. CKD in patients with bilateral oophorectomy. *Clin J Am Soc Nephrol* 2018;13:1649–58.
- 56 Georgakis MK, Beskou-Kontou T, Theodoridis I, Skalkidou A, Petridou ET. Surgical menopause in association with cognitive function and risk of dementia: a systematic review and meta-analysis. *Psychoneuroendocrinology* 2019;106:9–19.
- 57 Hall E, Finch A, Jacobson M, Rosen B, Metcalfe K, Sun P, Narod SA, Kotsopoulos J. Effects of bilateral salpingo-oophorectomy on menopausal symptoms and sexual functioning among women with a BRCA1 or BRCA2 mutation. *Gynecol Oncol* 2019;152:145–50.
- 58 Philp L, Alimena S, Ferris W, Saini A, Bregar AJ, Del Carmen MG, Eisenhauer EL, Growdon WB, Goodman A, Dorney K, Mazina V, Sisodia RC. Patient reported outcomes after risk-reducing surgery in patients at increased risk of ovarian cancer. *Gynecol Oncol* 2022;164:421–7.
- 59 Vermeulen RFM, Beurden Mvan, Kieffer JM, Bleiker EMA, Valdimarsdottir HB, Massuger LFAG, Mourits MJE, Gaarenstroom KN, van Dorst EBL, van der Putten HWHM, Aaronson NK. Hormone replacement therapy after risk-reducing salpingo-oophorectomy minimises endocrine and sexual problems: a prospective study. *Eur J Cancer* 2017;84:159–67.
- 60 Huber D, Seitz S, Kast K, Emons G, Ortmann O. Hormone replacement therapy in BRCA mutation carriers and risk of ovarian, endometrial, and breast cancer: a systematic review. *J Cancer Res Clin Oncol* 2021;147:2035–45.
- 61 Vermeulen RFM, Korse CM, Kenter GG, Brood-van Zanten MMA, Beurden Mvan. Safety of hormone replacement therapy following risk-reducing salpingo-oophorectomy: systematic review of literature and guidelines. *Climacteric* 2019;22:352–60.
- 62 Johansen N, Liavaag AH, Mørkrid L, Michelsen TM. Hormone levels and sexual functioning after risk-reducing salpingo-oophorectomy. *Sex Med* 2018;6:143–53.
- 63 Hollowell N, Baylock B, Heinger L, Butow PN, Patel D, Meiser B, Saunders C, Price MA, KConFab Psychosocial Group on behalf of the KConFab Investigators. Looking different, feeling different: women's reactions to risk-reducing breast and ovarian surgery. *Fam Cancer* 2012;11:215–24.
- 64 Hollowell N. A qualitative study of the information needs of high-risk women undergoing prophylactic oophorectomy. *Psychooncology* 2000;9:486–95.
- 65 Manchanda R, Legood R, Antoniou AC, Gordeev VS, Menon U. Specifying the ovarian cancer risk threshold of 'premenopausal risk-reducing salpingo-oophorectomy' for ovarian cancer prevention: a cost-effectiveness analysis. *J Med Genet* 2016;53:591–9.
- 66 Manchanda R, Menon U. Setting the threshold for surgical prevention in women at increased risk of ovarian cancer. *Int J Gynecol Cancer* 2018;28:34–42.
- 67 Lee A, Yang X, Tyrer J, Gentry-Maharaj A, Ryan A, Mavaddat N, Cunningham AP, Carver T, Archer S, Leslie G, Kalsi J, Gaba F, Manchanda R, Gayther S, Ramus SJ, Walter FM, Tischkowitz M, Jacobs I, Menon U, Easton DF, Pharoah P, Antoniou AC. Comprehensive epithelial tubo-ovarian cancer risk prediction model incorporating genetic and epidemiological risk factors. *J Med Genet* 2022;59:632–43.
- 68 Samuel D, Diaz-Barbe A, Pinto A, Schlumberg M, George S. Hereditary ovarian carcinoma: cancer pathogenesis looking beyond *BRCA1* and *BRCA2*. *Cells* 2022;11. doi:10.3390/cells11030539. [Epub ahead of print: 04 02 2022].
- 69 Falconer H, Yin L, Grönberg H, Altman D. Ovarian cancer risk after salpingectomy: a nationwide population-based study. *J Natl Cancer Inst* 2015;107. doi:10.1093/jnci/dju410. [Epub ahead of print: 27 01 2015].
- 70 Kindelberger DW, Lee Y, Miron A, Hirsch MS, Feltmate C, Medeiros F, Callahan MJ, Garner EO, Gordon RW, Birch C, Berkowitz RS, Muto MG, Crum CP. Intraepithelial carcinoma of the fimbria and pelvic serous carcinoma: evidence for a causal relationship. *Am J Surg Pathol* 2007;31:161–9.

- 71 Steenbeek MP, Harmsen MG, Hoogerbrugge N, de Jong MA, Maas AHEM, Prins JB, Bulten J, Teerenstra S, van Bommel MHD, van Doorn HC, Mourits MJE, van Beurden M, Zweemer RP, Gaarenstroom KN, Slangen BFM, Brood-van Zanten MMA, Vos MC, Piek JMJ, van Lonkhuijzen LRCW, Apperloo MJA, Coppus SFPJ, Massuger L FAG, Int'Hout J, Hermens R PMG, de Hullu JA. Association of Salpingectomy with delayed oophorectomy versus salpingo-oophorectomy with quality of life in BRCA1/2 pathogenic variant carriers: a nonrandomized controlled trial. *JAMA Oncol* 2021;7:1203–12.
- 72 Gaba F, Robbani S, Singh N, McCluggage WG, Wilkinson N, Ganesan R, Bryson G, Rowlands G, Tyson C, Arora R, Saridogan E, Hanson H, Burnell M, Legood R, Evans DG, Menon U, Manchanda R, PROTECTOR team. Preventing ovarian cancer through early excision of tubes and late ovarian removal (protector): protocol for a prospective non-randomised multi-center trial. *Int J Gynecol Cancer* 2021;31:286–91.
- 73 Boerner T, Long Roche K. Salpingectomy for the risk reduction of ovarian cancer: is it time for a Salpingectomy-alone approach? *J Minim Invasive Gynecol* 2021;28:403–8.
- 74 Yang X, Eriksson M, Czene K, Lee A, Leslie G, Lush M, Wang J, Dennis J, Dorling L, Carvalho S, Mavaddat N, Simard J, Schmidt MK, Easton DF, Hall P, Antoniou AC. Prospective validation of the BOADICEA multifactorial breast cancer risk prediction model in a large prospective cohort study. *J Med Genet* 2022;jmg-2022-108806.
- 75 NICE. Shared decision making NICE guideline; 2021.
- 76 Li N, McInerney S, Zethoven M, Cheasley D, Lim BWX, Rowley SM, Devereux L, Grewal N, Ahmadloo S, Byrne D, Lee JEA, Li J, Fox SB, John T, Antill Y, Goringe KL, James PA, Campbell IG. Combined tumor sequencing and case-control analyses of RAD51C in breast cancer. *J Natl Cancer Inst* 2019;111:1332–8.
- 77 Suszynska M, Klonowska K, Jasinska AJ, Kozłowski P. Large-scale meta-analysis of mutations identified in panels of breast/ovarian cancer-related genes - Providing evidence of cancer predisposition genes. *Gynecol Oncol* 2019;153:452–62.
- 78 Castéra L, Harter V, Muller E, Krieger S, Goardon N, Ricou A, Rousselin A, Paimparay G, Legros A, Bruet O, Quesnelle C, Domin F, San C, Brault B, Fouillet R, Abadie C, Béra O, Berthet P, Frébourg T, Vaur D, French Exome Project Consortium. Landscape of pathogenic variations in a panel of 34 genes and cancer risk estimation from 5131 HBOC families. *Genet Med* 2018;20:1677–86.
- 79 Hauke J, Horvath J, Groß E, Gehrig A, Honisch E, Hackmann K, Schmidt G, Arnold N, Faust U, Sutter C, Hentschel J, Wang-Gohrke S, Smogavec M, Weber BHF, Weber-Lassalle N, Weber-Lassalle K, Borde J, Ernst C, Altmüller J, Volk AE, Thiele H, Hübel V, Nürnberg P, Keupp K, Versmold B, Pohl E, Kubisch C, Grill S, Paul V, Herold N, Lichey N, Rhiem K, Ditsch N, Ruckert C, Wappenschmidt B, Auber B, Rump A, Niederacher D, Haaf T, Ramser J, Dworniczak B, Engel C, Meindl A, Schmutzler RK, Hahnen E. Gene panel testing of 5589 BRCA1/2-negative index patients with breast cancer in a routine diagnostic setting: results of the German Consortium for hereditary breast and ovarian cancer. *Cancer Med* 2018;7:1349–58.
- 80 Couch FJ, Shimelis H, Hu C, Hart SN, Polley EC, Na J, Hallberg E, Moore R, Thomas A, Lilyquist J, Feng B, McFarland R, Pesaran T, Huether R, LaDuca H, Chao EC, Goldgar DE, Dolinsky JS. Associations between cancer predisposition testing panel genes and breast cancer. *JAMA Oncol* 2017;3:1190–6.
- 81 Slavin TP, Maxwell KN, Lilyquist J, Vijai J, Neuhausen SL, Hart SN, Ravichandran V, Thomas T, Maria A, Villano D, Schrader KA, Moore R, Hu C, Wubbenhorst B, Wenz BM, D'Andrea K, Robson ME, Peterlongo P, Bonanni B, Ford JM, Garber JE, Domchek SM, Szabo C, Offit K, Nathanson KL, Weitzel JN, Couch FJ. The contribution of pathogenic variants in breast cancer susceptibility genes to familial breast cancer risk. *NPJ Breast Cancer* 2017;3:22.
- 82 Loveday C, Turnbull C, Ruark E, Xicola RMM, Ramsay E, Hughes D, Warren-Perry M, Snape K, Eccles D, Evans DG, Gore M, Renwick A, Seal S, Antoniou AC, Rahman N, Breast Cancer Susceptibility Collaboration (UK). Germline RAD51C mutations confer susceptibility to ovarian cancer. *Nat Genet* 2012;44:475–6.