

1 **British Gynaecological Cancer Society/British Association of Gynaecological**  
2 **Pathology consensus for germline and tumour testing for *BRCA1/2* variants in**  
3 **ovarian cancer in the United Kingdom**

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41

42 **Abstract**

43 The British Gynaecological Cancer Society and the British Association of Gynaecological  
44 Pathologists established a multidisciplinary consensus group comprising experts in surgical  
45 gynaecological oncology, medical oncology, genetics, laboratory science and clinical nurse  
46 specialists to identify the optimal pathways to *BRCA* germline and tumour testing in patients  
47 with ovarian cancer in routine clinical practice. In particular, the group explored models of  
48 consent, quality standards identified at pathology, laboratory and experience/data from  
49 pioneering cancer centres. The group liaised with representatives from ovarian cancer charities  
50 to also identify patient perspectives that would be important to implementation.  
51 Recommendations from this consensus group deliberations are presented in this manuscript.

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## 55 Introduction

56 Pathogenic germline *BRCA1/2* variants play a key role in the etiology of epithelial ovarian  
57 cancer. Recent studies showing the prevalence of pathogenic *BRCA* germline mutations in  
58 patients with high-grade serous ovarian cancer of 13-15% as well as the recognition of the  
59 clinically significant role of therapeutic poly-ADP ribose polymerase (PARP) inhibition in *BRCA*  
60 deficient tumours has led to an expansion in demand for germline *BRCA* testing.<sup>1-6</sup> The Cancer  
61 Genome Atlas (TCGA) identified somatic and germline *BRCA* pathogenic variants in ~22% of  
62 high-grade serous ovarian cancers.<sup>7</sup>

63 To manage this increased demand and ensure timely access to testing early on in the patient  
64 care pathway, models of delivery using surgeons, oncologists or clinical nurse specialists to  
65 “mainstream” germline testing have been developed in many centres. In these models, cancer  
66 clinicians counsel and offer germline *BRCA* testing to all ovarian cancer patients and only  
67 patients with pathogenic variants or variants of uncertain significant are referred to genetics  
68 services.

69 Different models have developed across the UK with variable testing criteria, availability and  
70 access.<sup>4, 8, 9</sup> Some models restrict testing to defined histological criteria (high-grade serous or  
71 endometrioid), others restrict testing to age groups (under 70 years). However, there is  
72 considerable variability in implementation of mainstream germline *BRCA* testing worldwide  
73 with some centres still relying on individual clinicians referring patients to regional genetics  
74 centres and approximately 30% of eligible patients not being offered testing.<sup>10</sup>

75

76 Until 2018, the evidence base for maintenance PARP inhibition strategies was restricted to  
77 women with relapsed ovarian cancer. However, following publication of the SOLO-1 trial, the  
78 evidence for benefit has been demonstrated in the first-line setting with women with *BRCA*-  
79 deficient advanced stage IIIC/IV ovarian cancer having significantly longer progression-free  
80 survival with maintenance olaparib compared to placebo.<sup>11</sup>

81

82 There are currently two methods by which *BRCA* testing may be undertaken, each of which  
83 detects slightly different pathogenic variants due to the pathogenesis of the mutations and the  
84 limitations of the analytical techniques. *Germline testing* is undertaken on blood samples and  
85 will detect inherited pathogenic variants, including the large duplications/deletions which are  
86 not reliably detectable on tumour testing. Thus, germline testing results carries implications  
87 for family members. *Tumour testing* involves extracting DNA from the ovarian tumour and  
88 subjected to test for pathogenic variants. Approximately two-thirds of the mutations detected  
89 in tumour will be of *germline* (inherited) origin, however nearly one-third will be found to be  
90 *somatic* (tumour only – not inherited) mutations. Therefore, tumour testing results may have  
91 implications for family members in some, but not all instances.

92 Crucially, PARP inhibition increases progression-free survival in patients with somatic *BRCA*  
93 mutation.<sup>11</sup> Therefore, patients and clinicians need as much information as possible to guide  
94 treatment choices in the first-line setting.

95 Thus, there is an urgent clinical need to clearly identify women whose tumours contain  
96 deleterious *BRCA* mutations early in their ovarian cancer treatment journey to maximize the  
97 population of women afforded the opportunity of PARP inhibitor treatment upon completion  
98 of first-line chemotherapy. Additionally, unselected germline testing identifies approximately

99 50% more women whose families may benefit from predictive testing and subsequent  
100 screening and prevention in unaffected individuals.<sup>12</sup>

101 Implementing these tests into routine practice at first-line treatment of ovarian cancer  
102 requires careful consideration of issues around scheduling of both tests, the timing of testing  
103 in relation to first-line therapy, counselling of patients, costs involved, sample management  
104 processes, quality controls and audit trails. This guidance document evaluates the underlying  
105 evidence and sets out recommendations for implementation into clinical practice in the United  
106 Kingdom.

#### 107 *Detection of different DNA variants in germline testing*

108 Next generation sequencing based technologies are used for detection of *BRCA* ‘point  
109 mutations’ (single nucleotide variants or small insertion/deletion variants typically <40 bp in  
110 size) in both blood (germline) and tumour samples. Although pathogenic large genomic  
111 rearrangements can be detected in germline samples using next generation sequencing, the  
112 algorithms show reduced sensitivity for smaller, single exon large genomic rearrangements.  
113 Consequently, pathogenic large genomic rearrangements in *BRCA* are typically detected in  
114 clinical laboratories using multiplex ligation dependent probe amplification in blood samples.  
115 However, multiplex ligation dependent probe amplification has a high analytical failure rate in  
116 formalin fixed paraffin embedded derived tumour DNA due to poor DNA quality and genomic  
117 instability present in many ovarian tumours and is consequently not routinely employed.

#### 118 *Scheduling of germline and tumour BRCA testing*

119 The consensus group carefully reviewed the emerging evidence summarised below to  
120 formulate its recommendation on scheduling of testing.

121 Evidence from the SIGNPOST study

122 A concomitant/parallel panel germline and tumour genetic testing pathway for all high-grade  
123 non-mucinous epithelial ovarian cancer was initially introduced at Barts Health (North East  
124 London Cancer Network) in 2016. This involved an initial period of training of clinical staff  
125 (surgeons, medical oncologists, clinical nurse specialists, design of patient information  
126 materials and was undertaken within the SIGNPOST (Systematic Genetic Testing for  
127 Personalised Ovarian Cancer Therapy) study (ISRCTN 16988857). Germline testing included  
128 testing for *BRCA1*, *BRCA2*, *RAD51C*, *RAD51D*, *BRIP1*. Tumour testing was undertaken for *BRCA1*  
129 and *BRCA2* genes. Both germline and tumour testing were done in parallel. This was offered  
130 both prospectively and retrospectively to those with a pre-existing diagnosis.

131 Pathogenic variant rates identified in the SIGNPOST study were consistent with what has been  
132 previously reported in the literature. Critically, this study shows that 10% of *BRCA* mutation  
133 carriers (those individuals with large genomic rearrangements) would not have been identified  
134 without concomitant parallel testing for both germline and somatic mutations (personal  
135 communication Prof Manchanda, unpublished data).

136 Evidence from Imperial College Healthcare NHS Trust and the Royal Marsden Hospital

137 At Imperial College Healthcare NHS Trust, parallel germline and tumour *BRCA* genetic testing  
138 is offered to all eligible ovarian cancer patients. The cancer team discuss the pathways and  
139 possibility of genetic testing and its implications with the patient at initial presentation. If  
140 consent is obtained, germline and tumour tests are requested from the gynaecological  
141 oncology clinic.

142 The Royal Marsden Hospital initiated mainstream germline *BRCA* testing in 2012 for all patients  
143 with non-mucinous ovarian cancer through the oncology teams as standard of care.  
144 Subsequently, reflex tumour testing was introduced for all patients with high-grade serous  
145 ovarian cancer. Currently, the data (unpublished) from The Royal Marsden Hospital has  
146 identified 9% of patients with pathogenic variants present only in the tumour; and 15% of  
147 patients with germline pathogenic variants that were not detected in the tumour testing. All  
148 of the latter represent large genomic rearrangements (duplications or deletions) that are not  
149 reliably detectable during tumour *BRCA* testing due to DNA fragmentation.

150 Evidence from Public Health England

151 Data from Public Health England shows that as of end of February 2020, from a total of 17,384  
152 pathogenic *BRCA* variants reported by all labs in England, 1,830 were large genomic  
153 rearrangements. (Personal communication from Fiona McDonald, Programme Manager,  
154 Molecular, Genomic and Research Data National Disease Registration, Public Health England).  
155 See Figure -1. However, it is widely accepted in England, that there are several 'hotspots' for  
156 large genomic rearrangements, which also coincide with less access to testing, thus, the true  
157 proportion of large genomic rearrangements in this population may be closer to 15-17% of  
158 pathogenic variants. This would be consistent with data from the Manchester and Royal  
159 Marsden labs (unpublished).

160

161 In England, given above results, a parallel testing would be the most effective strategy and  
162 would avoid missing a proportion of patients (roughly 10%), as tumour testing alone using 'next  
163 generation sequencing' technology is likely to miss the proportion of patients with germline

164 pathogenic large genomic rearrangements of *BRCA*. Conversely, germline testing alone will miss  
165 a proportion of patients with only somatic variants in *BRCA*. Ongoing studies in Scotland will  
166 provide information for local populations.

167 Each health system will need to establish baseline rates to determine whether sequential  
168 testing or parallel testing is optimal for their patient groups. In patients with limited ethnicity  
169 specific data such as those from South Asian populations  
170 (<https://academic.oup.com/pcm/article/1/2/75/5106037>), parallel testing will be particularly  
171 important.

#### 172 *Timing of BRCA testing in relation to first-line treatment*

173 The consensus group reflected on two issues in this section; the first to preserve patient choice  
174 and autonomy in making an informed decision, the second the crucial utility of knowledge of  
175 *BRCA* status in decisions for neoadjuvant/adjuvant/maintenance treatments at first-line  
176 settings. The consensus group also had discussions with ovarian cancer charities representing  
177 patient perspectives. The consensus group agreed that preserving patient choice in timing of  
178 testing was key. However, discussions around *BRCA* testing should start at the earliest available  
179 opportunity in a patient's cancer diagnosis journey.

180 In the ideal scenario, earliest testing at the time of diagnosis of ovarian cancer is vital so that  
181 *BRCA* status is available when it is clinically most relevant to the patient and should factor in  
182 the local turnaround time for testing and the potential need for genetic counselling. It is  
183 recognized that patients may feel ready to undergo testing at different points in their cancer  
184 journey. The counselling and consenting can be carried out by a trained gynaecological  
185 oncologist, the referring gynaecologist with expertise in gynaecological oncology (cancer unit

186 lead in the UK), oncologist or adequately trained clinicians (Clinical Nurse Specialist). Some  
187 patients may need to access the genetics service for pre-test counselling and this should be  
188 supported where possible.

189 Initial consultation

190 *BRCA* tumour testing can be discussed with patients who present with a high clinical suspicion  
191 of ovarian cancer (carcinomatosis on CT (computerized tomography) scan with CA125/CEA  
192 ratio >25) at initial presentation to a referring gynaecologist (cancer unit lead in the UK) or  
193 gynaecological oncologist, prior to confirmatory histological or cytological diagnosis.

194 Consultation before primary cytoreductive surgery

195 As part of the counselling and consenting for primary cytoreductive surgery, informed consent  
196 should be sought for tumour *BRCA* mutation testing; this can be in the form of a verbal  
197 discussion which is documented. Although undertaken by some centres (and considered good  
198 practice), currently tumour testing does not necessitate written consent in the UK. Information  
199 on whether the patient has provided or declined consent for tumour testing should be  
200 communicated with the pathology team receiving the surgical specimens after primary  
201 cytoreductive surgery, by being recorded in the pathology request form or communicated via  
202 other means. This will enable a streamlined process wherein the pathology team can identify  
203 the representative tumour block (or slides) and arrange transfer of the specimen to the  
204 genomic laboratory hub once a diagnosis of high-grade serous carcinoma or high-grade  
205 endometrioid cancer of tubo-ovarian or peritoneal origin is confirmed.

206 Consultation after primary cytoreductive surgery

207 If the pathology of the surgery reveals non-mucinous high-grade epithelial ovarian cancer, the  
208 patient should be counselled about germline *BRCA* mutation testing and written consent must  
209 be obtained. If consenting for tumour *BRCA* mutation testing was not obtained prior to surgery,  
210 this should be done and the nominated pathologist should be informed.

211 Consultation before biopsy in patients planned to receive neoadjuvant chemotherapy:

212 If the patient is not suitable for primary cytoreductive surgery (or in cases of diagnostic  
213 uncertainty) counselling about tumour *BRCA* testing should be performed before the imaging-  
214 guided biopsy or diagnostic laparoscopy. Informed consent should be obtained either in the  
215 form of a verbal discussion which is documented or through a formal consent form. The fact  
216 whether the patient has provided or declined consent for tumour testing should be recorded  
217 in the pathology request form after biopsy or conveyed to the pathologist by other means  
218 (electronic records, letter or email).

219 *Special Considerations:*

220 Imaging-guided biopsy

221 In order to obtain adequate amount of chemotherapy naïve tissue, extra cores of tumour tissue  
222 should be obtained for the purpose of successful tumour *BRCA* mutation testing. This must be  
223 recorded in the histopathology request form. Experience from the BRITROC study suggests that  
224 image guided biopsy using an 18-gauge needle and two passes are feasible and acceptable to  
225 patients and results in adequate tissue sampling.<sup>13</sup> If the pre-chemotherapy biopsy does not  
226 yield adequate tissue sample for *BRCA* testing, tumour testing should be reconsidered from  
227 the interval debulking surgery specimens in patients with negative germline testing. As the

228 success rate of tumour sequencing from post chemotherapy specimens is lower (impaired DNA  
229 yield) compared to chemotherapy naïve tissue, maximum attempt should be made to obtain  
230 adequate amount of tissue during pre-treatment biopsy. If debulking surgery is not performed  
231 after neoadjuvant chemotherapy, repeat imaging-guided biopsy for tumour testing should be  
232 considered.

233 Diagnostic laparoscopy

234 Adequate biopsy should be taken to provide the genetic laboratories with a sufficient amount  
235 of tissue for tumour testing.

236 Ascites cytology (in rare cases where tissue cannot be obtained)

237 Ascitic fluid should be sent to the pathology laboratory to obtain a tumour cell-rich block. A  
238 summary of indications, timing, sequence of testing and consent process is summarised in  
239 Table 1.

240 *Pathology - Tissue handling and pathways for tumour BRCA testing*

241 The mutation testing relies on detecting a mutant allele in a background of wild type alleles. It  
242 is important that adequate numbers of malignant cells are available to provide DNA for the  
243 test. Therefore, maximising the tissue available in a diagnostic biopsy is of paramount  
244 importance. Any biopsy done with suspicion of tubo-ovarian cancer must be sampled in at least  
245 two blocks. One block (with the lesser volume of tumour) should have an H&E (hematoxylin  
246 and eosin) stain with a confirmatory panel of PAX8, WT1, ER and p53. In context of  
247 morphology, PAX8 +ve, WT1 +ve, ER +ve and p53 mutation/aberrant staining  
248 (<https://www.thebagp.org/download/bagp-uknegas-project-p53-interpretation-guide-2016/>) is  
249 confirmatory for tubal/ovarian high-grade serous carcinoma. In case of diagnostic uncertainty,

250 in order to preserve tissue, the case should be sent to a cancer centre for review before further  
251 tissue is used for immunohistochemistry. The second block should have an H&E stain to  
252 confirm presence of malignancy. This is the tissue that needs to be sent to the nominated  
253 pathologist/s. In resection specimens, the reporting pathologist should send one block of  
254 primary or metastatic carcinoma containing maximum viable and well-fixed tumour with its  
255 H&E-stained slide to the a designated pathologist. Cellblock from cytology received with  
256 suspicion of ovarian cancer should be sent to pathologist if confirmatory of tubal/ovarian high-  
257 grade serous carcinoma.

258 Pathology teams and clinical teams should jointly establish pathways for communication of  
259 requests for tumour testing. This communication should clearly document patient consent for  
260 testing. The nominated pathologist should mark tumour areas on H&E slide and estimate  
261 tumour volume. The tissue (as required by the genomic laboratory hub), marked slide and  
262 completed form are sent to the genomic laboratory hub. This should be recorded securely and,  
263 where possible, this record should be accessible to the clinical team. When result received, the  
264 result should be added to the initial pathology report as a supplementary and/or upload report  
265 on electronic patient record.

#### 266 *Genomic Laboratory Hub considerations*

267 The NHS Genomic Laboratory Hub network has limited capacity to undertake assessment of  
268 pathology samples for adequacy for somatic *BRCA* analysis from ovarian cancer patients. Their  
269 specialist expertise is the analysis of nucleic acids. It is the primary responsibility of the  
270 pathology laboratory holding the tissue sample to undertake an assessment of the adequacy  
271 of tissue samples for tumour *BRCA* analysis. This should include an assessment of the  
272 neoplastic cell content of the sample. It is recommended that the neoplastic cell content of

273 samples should be at least twice the limit of detection of the assay used. For next generation  
274 sequencing based assays, the typical minimum neoplastic cell content for reliable detection of  
275 pathogenic variants is 20%. Formalin fixed paraffin embedded samples with less than 20%  
276 neoplastic cell content and regions of higher neoplastic cell content may be 'rescued' by  
277 macrodissection in the genomic laboratory. Macrodissection by the referring pathologist  
278 should, therefore, be considered for any samples where the neoplastic cell content is less than  
279 the minimum recommended by the genomics laboratory. A clearly marked H&E-stained guide  
280 slide with areas of neoplasia ringed using an indelible marker should be sent along with  
281 unstained slide mounted sections. The H&E guide slide should be derived from a serial section  
282 next to the sections sent for genomic analysis. Tissue morphology can change as successive  
283 sections are cut from the block and a neighbouring section mitigates against macrodissecting  
284 an inappropriate region of the tissue section.

285 Genomic target test turnaround times for genomic laboratory hubs in England are set by  
286 National Health Service England. The key turnaround times appropriate for ovarian cancer are  
287 21 calendar days for tumour *BRCA* analysis and 42 calendar days for germline *BRCA* analysis.  
288 Genomic laboratories are expected to meet these in at least 90% of the cases.

#### 289 *Consent issues*

290 With the roll-out of the NHS Genomic Medicine Service, patients across England gain equity of  
291 access to genomic testing for the first time, including whole genome sequencing for certain  
292 rare diseases and cancers. Healthcare professionals will need to be equipped to facilitate  
293 patient consent to these tests, and provide the information and support required.

294 To support this, the Genomics Education Programme has developed a competency framework  
295 that identifies eight areas of proficiency to facilitate and consent patients to genomic tests.  
296 (<https://www.genomicseducation.hee.nhs.uk/consent-a-competency-framework/>). It is  
297 intended as a cross-professional guide for best practice and has been designed around four  
298 categories of healthcare professionals based on their training and experience with genomics.  
299 The competency framework can be used by individual healthcare professionals as a guide to  
300 help them identify their learning needs. For educators, the framework provides a mechanism  
301 to recognise the training needs of health professional groups, and to structure training so that  
302 consent conversations about genomic testing can be delivered consistently across different  
303 specialties. In addition, the competencies can be used to evaluate how consent is being  
304 facilitated in different practice areas to enhance the delivery of genomic medicine.

305 Crucially, with the new framework, consent is rightly seen as a process whereby an 'offer' is  
306 made, adequate information provided and discussions to enable informed choice by patients  
307 are provided. Until the 'patient choice' forms are readily available in the UK (as detailed in the  
308 Genomics education programme), the current consent forms can be used and adapted to  
309 indicate if a patient has provided consent for somatic/germline/or combination (parallel)  
310 testing. It must be recorded in the patient notes that the discussion about opting to have a  
311 BRCA test has taken place over different points in the diagnostic/treatment work up. The  
312 consenting process should comply with General Medical Council standards. ([https://www.gmc-](https://www.gmc-uk.org/ethical-guidance/ethical-guidance-for-doctors/consent)  
313 [uk.org/ethical-guidance/ethical-guidance-for-doctors/consent](https://www.gmc-uk.org/ethical-guidance/ethical-guidance-for-doctors/consent))

314 In all cases, high quality, culturally appropriate information must be provided to patients so  
315 they can make an informed decision. Please see Appendix 2-4 for template letters.

316 *Recording of BRCA status and multidisciplinary team meeting outputs*

317 Consistency of terminology is important to avoid confusion. For instance, use of the term  
318 “BRCA positive” should be avoided as it can be interpreted to mean the diametric opposites of  
319 the positive presence of a mutation or the positive presence of protein. To avoid confusion the  
320 following terms should therefore be used: germline variant – a variant detected in the blood  
321 sample vs. tumour variant – a variant detected in the tumour. Importantly, without reference  
322 to the blood sample, a tumour variant could be either germline or somatic. Somatic variant –  
323 a pathogenic variant detected in the tumour sample which is not present in the blood sample.  
324 To define a somatic variant therefore requires that both a blood and a tumour sample have  
325 been analysed.

326 For ease of recording a common notation is to use a prefix to define the type of variant  
327 described and a suffix to describe the result. Using these notations, g, t, s are used to describe  
328 germline, tumour and somatic, respectively. Additionally, m, vus & wt are used to describe  
329 pathogenic or likely-pathogenic variant (mutation), variant of unknown significance and wild  
330 type respectively. For example, gBRCA1m would describe a germline variant (pathogenic or  
331 likely-pathogenic variant) of BRCA1, in contrast to sBRCA2wt which would describe a somatic  
332 wild type (no pathogenic variant) BRCA2. For more information on classes of variant. Table 2

333 *Patient perspectives*

334 Conversations with gynaecological cancer charities have highlighted issues of concern and  
335 importance for patients that need to be considered when implementing *BRCA* testing.  
336 Critically, patients should feel reassured that the timing of *BRCA* testing is their decision as

337 patients may feel ready to undergo testing at different points in their journey. High quality,  
338 culturally appropriate information is vital to this. Table 3

## 339 Conclusions

340 Germline testing has significant implications for patients, in terms of therapy choices, but also  
341 for their families in terms of risk management and the development of additional tumours.  
342 Tumour *BRCA* testing identifies an additional subgroup of women who have benefit from PARP  
343 inhibitors. Recommendations for testing are summarised in Table 4. It remains of critical  
344 importance to stratify patients and identify those who do not have a *BRCA* (germline/somatic)  
345 pathogenic variant as this group of women are least likely to benefit from PARP inhibitors and  
346 should therefore be considered for studies of novel therapies/combinations going forward.  
347 Additionally, family members who have a pathogenic/likely pathogenic variant can opt for a  
348 range of interventions such as reproductive choices, prenatal genetic diagnosis, planning a  
349 family, risk reduction surgery, screening or chemoprevention to minimize their ovarian cancer  
350 and breast cancer risk.

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356

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358 Sudha Sundar has received honoraria from Astra Zeneca outside the submitted work.  
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367 nothing to disclose.  
368  
369  
370

371 **List of Figures and Tables**

372 Figure 1: Proportion of germline pathogenic variants from hereditary Breast and Ovarian cancer  
373 patients that are large genomic re-arrangements.

374

375 Table – 1. Summary of testing of BRCA genes in Ovarian cancer in the UK

376 Table – 2. Classes of variants.

377 Table – 3. Patients perspectives on BRCA testing

378 Table -4. Recommendations for BRCA testing in ovarian cancer the UK

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