

THE UNIVERSITY of EDINBURGH

Edinburgh Research Explorer

MSH2 is the very young onset ovarian cancer predisposition gene, not BRCA1

Citation for published version:

Flaum, N, Crosbie, EJ, Woodward, ER, Lalloo, F, Morgan, R, Ryan, N & Evans, DG 2023, 'MSH2 is the very young onset ovarian cancer predisposition gene, not BRCA1', Journal of Medical Genetics. https://doi.org/10.1136/jmg-2022-109055

Digital Object Identifier (DOI):

10.1136/jmg-2022-109055

Link:

Link to publication record in Edinburgh Research Explorer

Document Version: Publisher's PDF, also known as Version of record

Published In: Journal of Medical Genetics

General rights

Copyright for the publications made accessible via the Edinburgh Research Explorer is retained by the author(s) and / or other copyright owners and it is a condition of accessing these publications that users recognise and abide by the legal requirements associated with these rights.

Take down policy The University of Edinburgh has made every reasonable effort to ensure that Edinburgh Research Explorer content complies with UK legislation. If you believe that the public display of this file breaches copyright please contact openaccess@ed.ac.uk providing details, and we will remove access to the work immediately and investigate your claim.



MSH2 is the very young onset ovarian cancer predisposition gene, not *BRCA1*

Referenced paragraph

The genetic cause of very young onset ovarian cancer (VYOC), diagnosed under 30 years of age, is unclear.¹ The histology and underlying genetics in VYOC is significantly different from the overall epithelial ovarian cancer (EOC) population; we aimed to explore this in VYOC cases known to the North-West of England. We found mismatch repair genes to be the most commonly affected in VYOCs, especially *MSH2*. The cumulative likelihood of an EOC in *MSH2* heterozygotes is >2% by age 35, with this likelihood still below 0.5% for *BRCA1* and rare for *BRCA2*.²

Article

The inherited landscape of epithelial ovarian cancer (EOC) is well established with contributions from homologous recombination deficiency (HRD) genes, particularly BRCA1 and BRCA2, and mismatch repair deficiency (MMRD) genes MSH2, MLH1, MSH6 and PMS2.¹ Highgrade serous ovarian cancer (HGSOC) is associated with HRD, accounting for up to 23% of HGSOC.² Approximately 3% of EOC cases occur in <30 years of age, described as very young onset ovarian cancer (VYOC).¹ The pathology in VYOC differs from overall EOC; a study of 114 VYOC cases found only 28% were serous, while 59% had mucinous pathology. Among the 101 tested cases, no BRCA1 or BRCA2 pathogenic variant (PV) was identified, only 2 MLH1 PVs.3 VYOC seems associated with MMRD-related EOC as opposed to homologous recombination deficiency (HRD)-related EOC as is seen

Table 1 Breakdown of VYOC cases by histological subtype									
Histological subtype	n (%)	Carriers of PV detected (%)	PV details						
HGSOC	6 (7.8)	0	NA						
LGSOC	4 (5.2)	0	NA						
Poorly differentiated	1 (1.3)	0	NA						
Serous NOS	12 (15.6)	1 (8.3)	MSH2x1						
Endometrioid	7 (9.1)	2 (28.6)	MSH2x1, PMS2x1						
Clear cell	2 (2.6)	1 (50)	MSH2x1						
Mucinous	29 (37.7)	0	NA						
Borderline	9 (11.7)	0	NA						
Granulosa cell	1 (1.3)	0	NA						
Adenocarcinoma NOS	1 (1.3)	0	NA						
Unknown	5 (6.5)	1 (20)	MSH2x1						
Total	77	5 (6.5%)	MSH2x4, PMS2x1						

HGSOC, high-grade serous ovarian cancer; LGSOC, low-grade serous ovarian cancer; NA, Not Applicable; NOS, not otherwise specified; PV, pathogenic variant; VYOC, very young onset ovarian cancer.

in *BRCA1/2* carriers, in whom the risk of EOC increases from 35 years for *BRCA1* carriers and from 45 years in *BRCA2* carriers.²⁴

We retrospectively assessed the presence of PVs in VYOC cases aged <30 and 30-34 years before the main risk period is associated with PVs in BRCA1/2. These women had been referred to Manchester Centre for Genomic Medicine (MCGM) within the last two decades following diagnosis with VYOC in the North-West of England. The genetic testing described was performed as part of standard diagnostic testing within MCGM.⁵ A series of 77 women with ovarian cancer (9 borderline and 1 granulosa cell tumour in addition to EOCs) were screened for MMRD and HRD germline PVs by sequencing, multiple ligation-dependent probe amplification and a prescreen for MMR immunohistochemistry (IHC) as previously described.5 We also assessed the proportion of known carriers that developed EOC at age <30 and 30-34 years from our extensive dataset of >4000 female BRCA1/2 carriers and 910 MMRD heterozygotes.

Of the 77 ovarian cancer cases aged <30 years, no *BRCA1/2* PV was identified. However, five MMRD PVs (four *MSH2*, one *PMS2*) were detected, making

up 6.5% of all cases and 7.5% of epithelial cases, as shown in table 1. Of the 69 invasive tumours, 5 were of unspecified histological subtype (1xMSH2), 2 clear cell (1xMSH2), 7 endometrioid (1xMSH2, 1xPMS2), 29 mucinous and 12 serous tumours not otherwise specified (1xMSH2). Age range was 15.0–29.9; mean age=25.00, median age=25.77, IQR=23.23–28.00. The age of the four MSH2 heterozygotes was 23.6, 25.1, 26.2 and 27.5 years and that of PMS2 homozygote was 26.8 years.

When assessing the proportions of VYOC in PV carriers of HRD and MMRD genes, we included all tested and obligate carriers. There were 2005 female BRCA1, 1999 BRCA2 and 393 MSH2 PV heterozygotes (table 2). One BRCA1 PV carrier (obligate carrier) was identified with an EOC at age <30(0.05%) (26 years old at diagnosis) but pathology subtype was unavailable. In contrast to the low rate in BRCA1/2 carriers, 4 out of 393 VYOC cases (1%) were found to carry the same MSH2 PV. This proportion was significantly greater in MSH2 PV carriers than either BRCA1 (p=0.003) or BRCA2 (p=0.0007) by χ^2 testing.

Of the 2005 BRCA1 and 1999 BRCA2 carriers, six and three cases of ovarian cancer were diagnosed, respectively,

Table 2	Proportion of women with VYOC with heterozygous PVs in BRCA1/2 and MMRD genes							
Gene	Female carriers (n)	OC <30 (%)	OR	P value	OC 30–34 (%)	OR	P value	
BRCA1	2005	1 (0.05)	0.05	0.003	6 (0.30)	0.23	0.02	
BRCA2	1999	0 (0.00)	NA	0.007	3 (0.15)	0.12	0.004	
MSH2	393	4 (1.02)	Ref	NA	5 (1.29)	Ref	NA	
MLH1	278	0 (0.00)	NA	NA	0 (0.00)	NA	0.013	
MSH6	169	0 (0.00)	NA	NA	0 (0.00)	NA	0.06	
PMS2	70	1 (1.43)	NA	NA	0 (0.00)	NA	0.36	
MMADD minimately reprint definition on NA Net Applicables OC evening concern DV anther price visite to finite the VVOC version expect evening concern								

Cancer genetics

between 30 and 34 years. This was significantly less than the five cases found in 393 *MSH2* PV carriers (p=0.02; 0.004, respectively). The proportion of ovarian cancer in *MSH2* PV carriers <35 years was significantly higher than 0 out of 278 found for *MLH1* PV carriers. In addition to the previous four cases, there were two clear cell, one endometrioid, one serous and one yolk sac tumour (non-EOC, but with MSH2 loss detected by IHC in tumour) histological subtypes. Only one *MSH2* PV carrier died from EOC with 77% surviving 10 years and 66% surviving >15 years.

In contrast, 7 out of 10 *BRCA1/2* PV heterozygotes had died, with 4 died <5 years of diagnosis. Two long-term *BRCA2* survivors both diagnosed at aged 32 had mucinous tumours (the remaining histologies were high-grade serous: *BRCA1=2*, adenocarcinoma not otherwise specified *BRCA1=5*; endometrioid *BRCA2=1*), raising doubts as to whether these were HRD-driven tumours. Hypothesising that these mucinous cases were not HRD-driven, survival at 12 years was significantly better in *MSH2* (77%) than *BRCA1/2* heterozygotes (15%; p=0.01).

Very few studies have addressed the contribution of HRD and MMRD genes to VYOC. In addition to the study of 101 cases aged <30,³ we identified a study of 47 women diagnosed at age \leq 40 with EOC who underwent germline screening for 11 genes associated with ovarian cancer. This identified PVs in 13 (28%) of women (*BRCA1*: 10, *BRCA2*: 1, *MSH2*: 1, *RAD51D*: 1).⁶ This study included only two women diagnosed under 30 years of age, neither of whom had an identifiable PV.

Our study has shown that while the genetic predisposition for many early onset ovarian cancers is still unknown, MSH2 is the most important EOC predisposition gene at age <35 years. The cumulative likelihood of an EOC in MSH2 heterozygotes would appear to be >2% by 35, with this likelihood still below 0.5% for BRCA1 and rare for $BRCA2^2$; indeed, two-thirds of cases identified in BRCA2

carriers may not have been driven by HRD. This increased incidence despite the good long-term survival in *MSH2* should prompt awareness of the increased risk and consideration for early risk-reduction strategies.

Nicola Flaum ⁽²⁾, ^{1,2} Emma J Crosbie ⁽³⁾, ^{3,4} Emma Roisin Woodward ⁽³⁾, ⁵ Fiona Lalloo, ⁵ Robert Morgan, ⁶ Neil Ryan, ⁷ D Gareth Evans ⁽⁵⁾, ^{1,2,6,8,9}

 ¹Division of Evolution, Infection and Genomics, School of Biological Sciences, Faculty of Biology, Medicine and Health, University of Manchester, Manchester Academic Health Science Centre, Manchester, UK
²North West Genomics Laboratory Hub, Manchester Centre for Genomic Medicine, St Mary's Hospital, Manchester University NHS Foundation Trust, Manchester Academic Health Science Centre, Manchester, UK
³Division of Cancer Sciences, School of Medical Sciences, Faculty of Biology, Medicine and Health,

University of Manchester, Manchester, UK ⁴Division of Gynaecology, St Mary's Hospital, Manchester University NHS Foundation Trust, Manchester, UK

⁵Clinical Genetics Service, Manchester Centre for Genomic Medicine, St Mary's Hospital, Manchester University NHS Foundation Trust, Manchester, UK ⁶The Christie NHS Foundation Trust, Manchester, UK ⁷The Academic Women's Health Unit, Translational Health Sciences, Bristol Medical School, University of Bristol, Bristol, UK

⁸Prevention Breast Cancer Centre and Nightingale Breast Screening Centre, University Hospital of South Manchester, Manchester, UK

⁹Manchester Breast Centre, Manchester Cancer Research Centre, University of Manchester, Manchester, UK

Correspondence to Dr Nicola Flaum; nicola.flaum@manchester.ac.uk

Twitter Nicola Flaum @drnikiflaum, Emma J Crosbie @DrEmmaCrosbie and Emma Roisin Woodward @ ER_Woodward

Collaborators Not applicable.

Contributors The project was conceived by DGE and NF, and primary manuscript was written by NF. Statistics was performed by DGE and NF. All authors commented on and edited the manuscript.

Funding DGE and EJC are supported by the National Institute for Health Research (NIHR) Manchester Biomedical Research Centre (IS-BRC-1215-20007). EJC is an NIHR Advanced Fellow (NIHR300650). NF is supported by CRUK via the funding to Cancer Research UK Manchester Centre: (C147/A18083) and (C147/ A25254).

Competing interests None declared.

Patient consent for publication Not applicable.

Provenance and peer review Not commissioned; externally peer reviewed.



Open access This is an open access article distributed in accordance with the Creative Commons Attribution 4.0 Unported (CC BY 4.0) license, which permits others to copy, redistribute, remix, transform and build upon this work for any purpose, provided the original work is properly cited, a link to the licence is given, and indication of whether changes were made. See: https:// creativecommons.org/licenses/by/4.0/.

© Author(s) (or their employer(s)) 2023. Re-use permitted under CC BY. Published by BMJ.



To cite Flaum N, Crosbie EJ, Woodward ER, *et al. J Med Genet* Epub ahead of print: [please include Day Month Year]. doi:10.1136/jmg-2022-109055

Received 14 November 2022 Accepted 22 December 2022

J Med Genet 2023;**0**:1–2. doi:10.1136/jmg-2022-109055

ORCID iDs

Nicola Flaum http://orcid.org/0000-0001-8900-0645 Emma J Crosbie http://orcid.org/0000-0003-0284-8630 Emma Roisin Woodward http://orcid.org/0000-0002-6297-2855

D Gareth Evans http://orcid.org/0000-0002-8482-5784

REFERENCES

- Cancer Research UK. Ovarian cancer incidence by age. 2022. Available: https://www.cancerresearchuk.org/ health-professional/cancer-statistics/statistics-by-cancertype/ovarian-cancer/incidence#heading-One
- Kuchenbaecker KB, Hopper JL, Barnes DR, et al. Risks of breast, ovarian, and contralateral breast cancer for BRCA1 and BRCA2 mutation carriers. JAMA 2017;317:2402–16.
- 3 Stratton JF, Thompson D, Bobrow L, et al. The genetic epidemiology of early-onset epithelial ovarian cancer: a population-based study. Am J Hum Genet 1999;65:1725–32.
- 4 Ryan NAJ, Evans DG, Green K, et al. Pathological features and clinical behavior of lynch syndromeassociated ovarian cancer. *Gynecol Oncol* 2017;144:491–5.
- 5 Woodward ER, Green K, Burghel GJ, et al. 30 year experience of index case identification and outcomes of cascade testing in high-risk breast and colorectal cancer predisposition genes. Eur J Hum Genet 2022;30:413–9.
- 6 Bernards SS, Norquist BM, Harrell MI, et al. Genetic characterization of early onset ovarian carcinoma. *Gynecol Oncol* 2016;140:221–5.