

Hereditary Breast, Ovarian and Pancreatic VCEP

Start Date	Your Name	Do you work for a laboratory that offers fee-for-service testing related to the work of your Expert Panel?	Have you made substantial contributions to the literature implicating a gene:disease relationship that relates to the work of your Expert Panel?		Do you have any other existing or planned independent efforts that will potentially overlap with the scope of your ClinGen work? If so, please describe and also send an email describing the project(s) to the co-chairs and coordinator:	Please disclose any other relevant conflicts of interest (e.g. patents, intellectual property ownership, or paid consultancies related to any variants or genes associated with the work of your Expert Panel):
	Open-Ended Response	Response	Response	If yes, please list the genes:	Open-Ended Response	Open-Ended Response
2021-01-26 10:02:45	Alvaro Monteiro	No	Yes	BRCA1, BRCA2, CHEK2, PALB2	Yes. My group has ongoing efforts on the integration of functional data for the classification of variants in breast and ovarian predisposition genes. Much of this work is also performed in the context of the ENIGMA consortium.	None to disclose
2020-06-18 15:58:32	Melissa Cline	No	No		Yes. I am the PI on a grant to develop federated analysis and data integration methods to support variant interpretation (1U01CA242954-01). It is my hope and vision that these activities will support the work of this expert panel. At the same time, I recognize that where the aims of the expert panel differ from the aims of my U01, the aims of the expert panel will take precedence.	None
2020-03-12 14:12:50	Colin Young	Yes	No		I was postdoc for Sean Tavtigian and am working to finish up a publication describing strengths of evidence provided by computational tools using AGVGD on ATM and CHEK2 cases and controls.	None
2020-03-04 16:44:48	Shannon McNulty	No	No		I am currently working to optimize an existing functional assay (double-strand break repair traffic light reporter system) for high-throughput analysis of PALB2 variants. The goal is to provide functional information that may aid in the reclassification of PALB2 VUS.	N/A
2019-09-10 14:59:41	Terra Brannan	Yes	No		Yes, I perform variant classifications as part of my employment	
2019-03-04 08:25:18	Sarah Brnich	No	No		I am curating some PALB2 missense variants from ClinVar for use in my dissertation, but would prefer for those curations to occur in the context of this expert panel.	N/A
2018-10-24 14:43:20	Thomas Slavin, M.D.	No	Yes	ATM, PALB2, CHEK2, NBN, RAD51C, RAD51D, BARD1	No	None
2018-08-29 17:39:25	Sean Tavtigian	No	Yes	ATM, CHEK2	PI of R01CA121245, which includes functional assays for ATM and CHEK2. Results could roll into ClinGen	None
2018-06-26 10:17:23	Holly LaDuca	Yes	Yes	ATM, BARD1, BRIP1, CHEK2, PALB2, RAD51C, RAD51D	No	employed by Ambry Genetics, a commercial genetic testing laboratory
2018-06-25 12:58:06	Kelly McGoldrick	Yes	No		I am participating in the "Developing consensus guidance on how to annotate low penetrance sequence and copy number variants for Mendelian disorders" ClinGen working group.	none
2018-06-23 21:23:58	Fergus Couch	No	Yes	PALB2 BRIP1 BARD1 RAD51C RAD51D ATM CHEK2	Functional studies of PALB2, RAD51C, RAD51D Penetrance studies of deleterious mutations in ATM, BARD1, BRIP1, CHEK2, PALB2, RAD51C, and RAD51D. In silico prediction studies for missense variants in ATM, BARD1, BRIP1, CHEK2, PALB2, RAD51C, and RAD51D.	
2018-06-12 15:45:51	Michael J. Anderson	Yes	No		I work in a large commercial lab that provides testing for all of the genes covered by this Expert Group.	None, other than those stated in Question 4.
2018-06-12 02:34:03	Logan Walker	No	Yes	BRCA1, BRCA2 'Other genes' projects within ENIGMA SWG, including BARD1, PALB2, and BRIP1 (Data not yet published)	No	No conflicts to disclose
2018-06-11 03:48:02	Arjen Mensenkamp	Yes	No		Membership of ENIGMA expert panel for BRCA1/2. and ClinGen CDH1 expert panel membership	
2018-06-10 08:40:26	Melissa Southey	No	Yes	ATM, CHEK2, PALB2	No	None to disclose
2018-06-10 04:08:29	Amanda Spurdle	No	No		No - my other work relates to BRCA1/2, TP53, CDH1, STK11 and MMR genes	none
2018-06-08 21:21:22	Jean-Yves Masson	No	Yes	PALB2, RAD51 paralogs, BRCA2	No	None
2018-06-08 14:06:07	Miguel de la Hoya	No	No		No	Naturally occurring alternative splicing analysis for some of these genes
2018-06-08 10:30:49	Lauren Yackowski	Yes	No		As an employee of a commercial laboratory, I am involved in ongoing variant and gene curation efforts utilizing published and internal data which may result in independent manuscripts during the course of this working group; however, we aim to not compete with any working group efforts that are established.	none
2018-06-08 10:26:04	Susan Hiraki	Yes	No		n/a	n/a
2018-06-08 10:19:12	Clare Turnbull	No	Yes	RAD51C RAD51D	CVIG-UK (Cancer Variant Interpretation Group): UK curation/classification effort for cancer susceptibility genes	Nil
2018-06-08 10:06:11	Sarah Nielsen	No	No			
2018-06-08 07:30:21	Huma Rana	No	No		TP53 genotype-phenotype work pending	N/a
2018-06-08 07:00:17	Tina Pesaran	Yes	No		I curate variants as a member of variant assessment team at Ambry Genetics	None
2018-06-08 04:09:29	Marc Tischkowitz	No	Yes	PALB2	I Lead the PALB2 Interest Group	none
2018-06-07 21:45:57	Marcy E. Richardson	Yes	No		No	None