

Breast cancer polygenic risk score and contralateral breast cancer risk

Iris Kramer¹, Maartje J. Hoening², Nasim Mavaddat³, Michael Hauptmann^{4, 5}, Renske Keeman¹, Ewout W Steyerberg^{6, 7}, Daniele Giardiello^{1, 6}, Antonis C. Antoniou³, Paul D.P. Pharoah^{3, 8}, Sander Canisius^{1, 9}, Zumuruda Abu-Ful¹⁰, Irene L. Andrulis^{11, 12}, Hoda Anton-Culver¹³, Kristan J. Aronson¹⁴, Annelie Augustinsson¹⁵, Heiko Becher^{16, 17}, Matthias W. Beckmann¹⁸, Sabine Behrens¹⁹, Javier Benitez^{20, 21}, Marina Bermisheva²², Natalia V. Bogdanova²³⁻²⁵, Stig E. Bojesen²⁶⁻²⁸, Manjeet K. Bolla³, Bernardo Bonanni²⁹, Hiltrud Brauch³⁰⁻³², Michael Bremer²³, Sara Y. Brucker³³, Barbara Burwinkel^{34, 35}, Jose E. Castelao³⁶, Tsun L. Chan^{37, 38}, Jenny Chang-Claude^{19, 39}, Stephen J. Chanock⁴⁰, Georgia Chenevix-Trench⁴¹, Ji-Yeob Choi^{42, 43}, Christine L. Clarke⁴⁴, NBCS Collaborators⁴⁵⁻⁵⁶, J. Margriet Collée⁵⁷, Fergus J. Couch⁵⁸, Angela Cox⁵⁹, Simon S. Cross⁶⁰, Kamila Czene⁶¹, Mary B. Daly⁶², Peter Devilee^{63, 64}, Thilo Dörk²⁴, Isabel dos-Santos-Silva⁶⁵, Alison M. Dunning⁸, Miriam Dwek⁶⁶, Diana M. Eccles⁶⁷, D. Gareth Evans^{68, 69}, Peter A. Fasching^{18, 70}, Henrik Flyger⁷¹, Manuela Gago-Dominguez^{72, 73}, Montserrat García-Closas⁴⁰, José A. García-Sáenz⁷⁴, Graham G. Giles⁷⁵⁻⁷⁷, David E. Goldgar⁷⁸, Anna González-Neira²¹, Christopher A. Haiman⁷⁹, Niclas Håkansson⁸⁰, Ute Hamann⁸¹, Mikael Hartman^{82, 83}, Bernadette A.M. Heemskerk-Gerritsen², Antoinette Hollestelle², John L. Hopper⁷⁶, Ming-Feng Hou⁸⁴, Anthony Howell⁸⁵, ABCTB Investigators⁸⁶, kConFab Investigators^{87, 88}, Hidemi Ito^{89, 90}, Milena Jakimovska⁹¹, Anna Jakubowska^{92, 93}, Wolfgang Janni⁹⁴, Esther M. John⁹⁵, Audrey Jung¹⁹, Daehee Kang^{42, 43, 96}, C. Marleen Kets⁹⁷, Elza Khusnutdinova^{22, 98}, Yon-Dschun Ko⁹⁹, Vessela N. Kristensen^{45, 56}, Allison W. Kurian^{95, 100}, Ava Kwong^{37, 101, 102}, Diether Lambrechts^{103, 104}, Loic Le Marchand¹⁰⁵, Jingmei Li¹⁰⁶, Annika Lindblom^{107, 108}, Jan Lubiński⁹², Arto Mannermaa¹⁰⁹⁻¹¹¹, Mehdi Manoochehri⁸¹, Sara Margolin^{112, 113}, Keitaro Matsuo^{89, 90}, Dimitrios Mavroudis¹¹⁴, Alfons Meindl¹¹⁵, Roger L. Milne⁷⁵⁻⁷⁷, Anna Marie Mulligan^{116, 117}, Taru A. Muranen¹¹⁸, Susan L. Neuhausen¹¹⁹, Heli Nevanlinna¹¹⁸, William G. Newman^{68, 69}, Andrew F. Olshan¹²⁰, Janet E.

Olson¹²¹, Håkan Olsson¹⁵, Tjong-Won Park-Simon²⁴, Julian Peto⁶⁵, Christos Petridis¹²², Dijana Plaseska-Karanfilska⁹¹, Nadege Presneau⁶⁶, Katri Pylkäs^{123, 124}, Paolo Radice¹²⁵, Gad Rennert¹⁰, Atocha Romero¹²⁶, Rebecca Roylance¹²⁷, Emmanouil Saloustros¹²⁸, Elinor J. Sawyer¹²⁹, Rita K. Schmutzler¹³⁰⁻¹³², Lukas Schwentner⁹⁴, Christopher Scott¹²¹, Mee-Hoong See¹³³, Mitul Shah⁸, Chen-Yang Shen^{134, 135}, Xiao-Ou Shu¹³⁶, Sabine Siesling^{137, 138}, Susan Slager¹²¹, Christof Sohn¹³⁹, Melissa C. Southey^{75, 77, 140}, John J. Spinelli^{141, 142}, Jennifer Stone^{76, 143}, William J. Tapper⁶⁷, Maria Tengström^{109, 144, 145}, Soo Hwang Teo^{146, 147}, Mary Beth Terry¹⁴⁸, Rob A.E.M. Tollenaar¹⁴⁹, Ian Tomlinson^{150, 151}, Melissa A. Troester¹²⁰, Celine M. Vachon¹⁵², Chantal van Ongeval¹⁵³, Elke M. van Veen^{68, 69}, Robert Winqvist^{123, 124}, Alicja Wolk^{80, 154}, Wei Zheng¹³⁶, Argyrios Ziogas¹³, Douglas F. Easton^{3, 8}, Per Hall^{61, 112}, Marjanka K. Schmidt^{1, 155}

¹ The Netherlands Cancer Institute - Antoni van Leeuwenhoek Hospital, Division of Molecular Pathology, Amsterdam, 1066 CX, The Netherlands.

² Erasmus MC Cancer Institute, Department of Medical Oncology, Rotterdam, 3015 CN, The Netherlands.

³ University of Cambridge, Centre for Cancer Genetic Epidemiology, Department of Public Health and Primary Care, Cambridge, CB1 8RN, UK.

⁴ The Netherlands Cancer Institute - Antoni van Leeuwenhoek hospital, Department of Epidemiology and Biostatistics, Amsterdam, 1066 CX, The Netherlands.

⁵ Brandenburg Medical School Theodor Fontane, Institute of Biostatistics and Registry Research, Neuruppin, 16816, Germany.

⁶ Leiden University Medical Center, Department of Biomedical Data Sciences, Leiden, 2333 ZA, The Netherlands.

⁷ Erasmus MC Cancer Institute, Department of Public Health, Rotterdam, 3015 GD, The Netherlands.

⁸ University of Cambridge, Centre for Cancer Genetic Epidemiology, Department of Oncology, Cambridge, CB1 8RN, UK.

⁹ The Netherlands Cancer Institute - Antoni van Leeuwenhoek hospital, Division of Molecular Carcinogenesis, Amsterdam, 1066 CX, The Netherlands.

¹⁰ Carmel Medical Center and Technion Faculty of Medicine, Clalit National Cancer Control Center, Haifa, 35254, Israel.

¹¹ Lunenfeld-Tanenbaum Research Institute of Mount Sinai Hospital, Fred A. Litwin Center for Cancer Genetics, Toronto, ON, M5G 1X5, Canada.

¹² University of Toronto, Department of Molecular Genetics, Toronto, ON, M5S 1A8, Canada.

¹³ University of California Irvine, Department of Epidemiology, Genetic Epidemiology Research Institute, Irvine, CA, 92617, USA.

¹⁴ Queen's University, Department of Public Health Sciences, and Cancer Research Institute, Kingston, ON, K7L 3N6, Canada.

¹⁵ Lund University, Department of Cancer Epidemiology, Clinical Sciences, Lund, 222 42, Sweden.

¹⁶ University Medical Center Hamburg-Eppendorf, Institute of Medical Biometry and Epidemiology, Hamburg, 20246, Germany.

¹⁷ Charité –Universitätsmedizin Berlin, Institute of Biometry and Clinical Epidemiology, Berlin, 10117, Germany.

¹⁸ University Hospital Erlangen, Friedrich-Alexander-University Erlangen-Nuremberg, Department of Gynecology and Obstetrics, Comprehensive Cancer Center ER-EMN, Erlangen, 91054, Germany.

¹⁹ German Cancer Research Center (DKFZ), Division of Cancer Epidemiology, Heidelberg, 69120, Germany.

²⁰ Centro de Investigación en Red de Enfermedades Raras (CIBERER), Madrid, 28029, Spain.

- ²¹ Spanish National Cancer Research Centre (CNIO), Human Cancer Genetics Programme, Madrid, 28029, Spain.
- ²² Ufa Federal Research Centre of the Russian Academy of Sciences, Institute of Biochemistry and Genetics, Ufa, 450054, Russia.
- ²³ Hannover Medical School, Department of Radiation Oncology, Hannover, 30625, Germany.
- ²⁴ Hannover Medical School, Gynaecology Research Unit, Hannover, 30625, Germany.
- ²⁵ N.N. Alexandrov Research Institute of Oncology and Medical Radiology, Minsk, 223040, Belarus.
- ²⁶ Copenhagen University Hospital, Copenhagen General Population Study, Herlev and Gentofte Hospital, Herlev, 2730, Denmark.
- ²⁷ Copenhagen University Hospital, Department of Clinical Biochemistry, Herlev and Gentofte Hospital, Herlev, 2730, Denmark.
- ²⁸ University of Copenhagen, Faculty of Health and Medical Sciences, Copenhagen, 2200, Denmark.
- ²⁹ IEO, European Institute of Oncology IRCCS, Division of Cancer Prevention and Genetics, Milan, 20141, Italy.
- ³⁰ Dr. Margarete Fischer-Bosch-Institute of Clinical Pharmacology, Stuttgart, 70376, Germany.
- ³¹ University of Tübingen, iFIT-Cluster of Excellence, Tübingen, 72074, Germany.
- ³² German Cancer Research Center (DKFZ), German Cancer Consortium (DKTK), Partner Site Tübingen, Tübingen, 72074, Germany.
- ³³ University of Tübingen, Department of Gynecology and Obstetrics, Tübingen, 72076, Germany.
- ³⁴ German Cancer Research Center (DKFZ), Molecular Epidemiology Group, C080, Heidelberg, 69120, Germany.
- ³⁵ University of Heidelberg, Molecular Biology of Breast Cancer, University Womens Clinic Heidelberg, Heidelberg, 69120, Germany.

- ³⁶ Instituto de Investigacion Sanitaria Galicia Sur (IISGS), Xerencia de Xestion Integrada de Vigo-SERGAS, Oncology and Genetics Unit, Vigo, 36312, Spain.
- ³⁷ Cancer Genetics Centre, Hong Kong Hereditary Breast Cancer Family Registry, Happy Valley, Hong Kong.
- ³⁸ Hong Kong Sanatorium and Hospital, Department of Pathology, Happy Valley, Hong Kong.
- ³⁹ University Medical Center Hamburg-Eppendorf, Cancer Epidemiology Group, University Cancer Center Hamburg (UCCH), Hamburg, 20246, Germany.
- ⁴⁰ National Cancer Institute, National Institutes of Health, Department of Health and Human Services, Division of Cancer Epidemiology and Genetics, Bethesda, MD, 20850, USA.
- ⁴¹ QIMR Berghofer Medical Research Institute, Department of Genetics and Computational Biology, Brisbane, Queensland, 4006, Australia.
- ⁴² Seoul National University Graduate School, Department of Biomedical Sciences, Seoul, 03080, Korea.
- ⁴³ Seoul National University, Cancer Research Institute, Seoul, 03080, Korea.
- ⁴⁴ University of Sydney, Westmead Institute for Medical Research, Sydney, New South Wales, 2145, Australia.
- ⁴⁵ Oslo University Hospital-Radiumhospitalet, Department of Cancer Genetics, Institute for Cancer Research, Oslo, 0379, Norway.
- ⁴⁶ University of Oslo, Institute of Clinical Medicine, Faculty of Medicine, Oslo, 0450, Norway.
- ⁴⁷ Vestre Viken Hospital, Department of Research, Drammen, 3019, Norway.
- ⁴⁸ Oslo University Hospital-Ullevål, Section for Breast- and Endocrine Surgery, Department of Cancer, Division of Surgery, Cancer and Transplantation Medicine, Oslo, 0450, Norway.
- ⁴⁹ Oslo University Hospital, Department of Radiology and Nuclear Medicine, Oslo, 0379, Norway.
- ⁵⁰ Akershus University Hospital, Department of Pathology, Lørenskog, 1478, Norway.

⁵¹ Oslo University Hospital, Department of Tumor Biology, Institute for Cancer Research, Oslo, 0379, Norway.

⁵² Oslo University Hospital-Radiumhospitalet, Department of Oncology, Division of Surgery, Cancer and Transplantation Medicine, Oslo, 0379, Norway.

⁵³ Oslo University Hospital-Radiumhospitalet, National Advisory Unit on Late Effects after Cancer Treatment, Department of Oncology, Oslo, 0379, Norway.

⁵⁴ Akershus University Hospital, Department of Oncology, Lørenskog, 1478, Norway.

⁵⁵ Oslo University Hospital, Oslo Breast Cancer Research Consortium, Oslo, 0379, Norway.

⁵⁶ Oslo University Hospital and University of Oslo, Department of Medical Genetics, Oslo, 0379, Norway.

⁵⁷ Erasmus University Medical Center, Department of Clinical Genetics, Rotterdam, 3015 CN, The Netherlands.

⁵⁸ Mayo Clinic, Department of Laboratory Medicine and Pathology, Rochester, MN, 55905, USA.

⁵⁹ University of Sheffield, Sheffield Institute for Nucleic Acids (SInFoNiA), Department of Oncology and Metabolism, Sheffield, S10 2TN, UK.

⁶⁰ University of Sheffield, Academic Unit of Pathology, Department of Neuroscience, Sheffield, S10 2TN, UK.

⁶¹ Karolinska Institutet, Department of Medical Epidemiology and Biostatistics, Stockholm, 171 65, Sweden.

⁶² Fox Chase Cancer Center, Department of Clinical Genetics, Philadelphia, PA, 19111, USA.

⁶³ Leiden University Medical Center, Department of Pathology, Leiden, 2333 ZA, The Netherlands.

⁶⁴ Leiden University Medical Center, Department of Human Genetics, Leiden, 2333 ZA, The Netherlands.

⁶⁵ London School of Hygiene and Tropical Medicine, Department of Non-Communicable Disease Epidemiology, London, WC1E 7HT, UK.

- ⁶⁶ University of Westminster, School of Life Sciences, London, W1B 2HW, UK.
- ⁶⁷ University of Southampton, Faculty of Medicine, Southampton, SO17 1BJ, UK.
- ⁶⁸ University of Manchester, Manchester Academic Health Science Centre, Division of Evolution and Genomic Sciences, School of Biological Sciences, Faculty of Biology, Medicine and Health, Manchester, M13 9WL, UK.
- ⁶⁹ St Mary's Hospital, Manchester University NHS Foundation Trust, Manchester Academic Health Science Centre, North West Genomics Laboratory Hub, Manchester Centre for Genomic Medicine, Manchester, M13 9WL, UK.
- ⁷⁰ University of California at Los Angeles, David Geffen School of Medicine, Department of Medicine Division of Hematology and Oncology, Los Angeles, CA, 90095, USA.
- ⁷¹ Copenhagen University Hospital, Department of Breast Surgery, Herlev and Gentofte Hospital, Herlev, 2730, Denmark.
- ⁷² Grupo de Medicina Xenómica, Instituto de Investigación Sanitaria de Santiago de Compostela (IDIS), Fundación Pública Galega de Medicina Xenómica, Santiago de Compostela, 15706, Spain.
- ⁷³ University of California San Diego, Moores Cancer Center, La Jolla, CA, 92037, USA.
- ⁷⁴ Instituto de Investigación Sanitaria San Carlos (IdISSC), Centro Investigación Biomédica en Red de Cáncer (CIBERONC), Medical Oncology Department, Hospital Clínico San Carlos, Madrid, 28040, Spain.
- ⁷⁵ Cancer Council Victoria, Cancer Epidemiology Division, Melbourne, Victoria, 3004, Australia.
- ⁷⁶ The University of Melbourne, Centre for Epidemiology and Biostatistics, Melbourne School of Population and Global Health, Melbourne, Victoria, 3010, Australia.
- ⁷⁷ Monash University, Precision Medicine, School of Clinical Sciences at Monash Health, Clayton, Victoria, 3168, Australia.
- ⁷⁸ Huntsman Cancer Institute, University of Utah School of Medicine, Department of Dermatology, Salt Lake City, UT, 84112, USA.

- ⁷⁹ University of Southern California, Department of Preventive Medicine, Keck School of Medicine, Los Angeles, CA, 90033, USA.
- ⁸⁰ Karolinska Institutet, Institute of Environmental Medicine, Stockholm, 171 77, Sweden.
- ⁸¹ German Cancer Research Center (DKFZ), Molecular Genetics of Breast Cancer, Heidelberg, 69120, Germany.
- ⁸² National University of Singapore and National University Health System, Saw Swee Hock School of Public Health, Singapore, 119077, Singapore.
- ⁸³ National University Health System, Department of Surgery, Singapore, 119228, Singapore.
- ⁸⁴ Kaohsiung Medical University, Chung-Ho Memorial Hospital, Kaohsiung, 807, Taiwan.
- ⁸⁵ University of Manchester, Division of Cancer Sciences, Manchester, M13 9PL, UK.
- ⁸⁶ University of Sydney, Australian Breast Cancer Tissue Bank, Westmead Institute for Medical Research, Sydney, New South Wales, 2145, Australia.
- ⁸⁷ Peter MacCallum Cancer Center, Melbourne, Victoria, 3000, Australia.
- ⁸⁸ The University of Melbourne, Sir Peter MacCallum Department of Oncology, Melbourne, Victoria, 3000, Australia.
- ⁸⁹ Aichi Cancer Center Research Institute, Division of Cancer Epidemiology and Prevention, Nagoya, 464-8681, Japan.
- ⁹⁰ Nagoya University Graduate School of Medicine, Division of Cancer Epidemiology, Nagoya, 466-8550, Japan.
- ⁹¹ MASA, Research Centre for Genetic Engineering and Biotechnology 'Georgi D. Efremov', Skopje, 1000, Republic of North Macedonia.
- ⁹² Pomeranian Medical University, Department of Genetics and Pathology, Szczecin, 71-252, Poland.
- ⁹³ Pomeranian Medical University, Independent Laboratory of Molecular Biology and Genetic Diagnostics, Szczecin, 71-252, Poland.
- ⁹⁴ University Hospital Ulm, Department of Gynaecology and Obstetrics, Ulm, 89075, Germany.

- ⁹⁵ Stanford Cancer Institute, Stanford University School of Medicine, Department of Epidemiology & Population Health, Stanford, CA, 94304, USA.
- ⁹⁶ Seoul National University College of Medicine, Department of Preventive Medicine, Seoul, 03080, Korea.
- ⁹⁷ The Netherlands Cancer Institute - Antoni van Leeuwenhoek Hospital, Department of Clinical Genetics, Amsterdam, 1066 CX, The Netherlands.
- ⁹⁸ Bashkir State University, Department of Genetics and Fundamental Medicine, Ufa, 450000, Russia.
- ⁹⁹ Johanniter Krankenhaus, Department of Internal Medicine, Evangelische Kliniken Bonn gGmbH, Bonn, 53177, Germany.
- ¹⁰⁰ Stanford University School of Medicine, Department of Health Research and Policy, Stanford, CA, 94305, USA.
- ¹⁰¹ The University of Hong Kong, Department of Surgery, Pok Fu Lam, Hong Kong.
- ¹⁰² Hong Kong Sanatorium and Hospital, Cancer Genetics Center and Department of Surgery, Happy Valley, Hong Kong.
- ¹⁰³ VIB Center for Cancer Biology, Leuven, 3001, Belgium.
- ¹⁰⁴ University of Leuven, Laboratory for Translational Genetics, Department of Human Genetics, Leuven, 3000, Belgium.
- ¹⁰⁵ University of Hawaii Cancer Center, Epidemiology Program, Honolulu, HI, 96813, USA.
- ¹⁰⁶ Genome Institute of Singapore, Human Genetics Division, Singapore, 138672, Singapore.
- ¹⁰⁷ Karolinska Institutet, Department of Molecular Medicine and Surgery, Stockholm, 171 76, Sweden.
- ¹⁰⁸ Karolinska University Hospital, Department of Clinical Genetics, Stockholm, 171 76, Sweden.
- ¹⁰⁹ University of Eastern Finland, Translational Cancer Research Area, Kuopio, 70210, Finland.
- ¹¹⁰ University of Eastern Finland, Institute of Clinical Medicine, Pathology and Forensic Medicine, Kuopio, 70210, Finland.

- ¹¹¹ Kuopio University Hospital, Biobank of Eastern Finland, Kuopio, 70210, Finland.
- ¹¹² Södersjukhuset, Department of Oncology, Stockholm, 118 83, Sweden.
- ¹¹³ Karolinska Institutet, Department of Clinical Science and Education, Södersjukhuset, Stockholm, 118 83, Sweden.
- ¹¹⁴ University Hospital of Heraklion, Department of Medical Oncology, Heraklion, 711 10, Greece.
- ¹¹⁵ University of Munich, Campus Großhadern, Department of Gynecology and Obstetrics, Munich, 81377, Germany.
- ¹¹⁶ University of Toronto, Department of Laboratory Medicine and Pathobiology, Toronto, ON, M5S 1A8, Canada.
- ¹¹⁷ University Health Network, Laboratory Medicine Program, Toronto, ON, M5G 2C4, Canada.
- ¹¹⁸ Helsinki University Hospital, Department of Obstetrics and Gynecology, University of Helsinki, Helsinki, 00290, Finland.
- ¹¹⁹ Beckman Research Institute of City of Hope, Department of Population Sciences, Duarte, CA, 91010, USA.
- ¹²⁰ University of North Carolina at Chapel Hill, Department of Epidemiology, Gillings School of Global Public Health and UNC Lineberger Comprehensive Cancer Center, Chapel Hill, NC, 27599, USA.
- ¹²¹ Mayo Clinic, Department of Health Sciences Research, Rochester, MN, 55905, USA.
- ¹²² King's College London, Research Oncology, Guy's Hospital, London, SE1 9RT, UK.
- ¹²³ University of Oulu, Laboratory of Cancer Genetics and Tumor Biology, Cancer and Translational Medicine Research Unit, Biocenter Oulu, Oulu, 90220, Finland.
- ¹²⁴ Northern Finland Laboratory Centre Oulu, Laboratory of Cancer Genetics and Tumor Biology, Oulu, 90220, Finland.
- ¹²⁵ Fondazione IRCCS Istituto Nazionale dei Tumori (INT), Unit of Molecular Bases of Genetic Risk and Genetic Testing, Department of Research, Milan, 20133, Italy.

- ¹²⁶ Hospital Universitario Puerta de Hierro, Medical Oncology Department, Madrid, 28222, Spain.
- ¹²⁷ UCLH Foundation Trust, Department of Oncology, London, NW1 2PG, UK.
- ¹²⁸ University Hospital of Larissa, Department of Oncology, Larissa, 411 10, Greece.
- ¹²⁹ King's College London, School of Cancer & Pharmaceutical Sciences, Comprehensive Cancer Centre, Guy's Campus, London, SE1 1UL, UK.
- ¹³⁰ Faculty of Medicine and University Hospital Cologne, University of Cologne, Center for Familial Breast and Ovarian Cancer, Cologne, 50937, Germany.
- ¹³¹ Faculty of Medicine and University Hospital Cologne, University of Cologne, Center for Integrated Oncology (CIO), Cologne, 50937, Germany.
- ¹³² Faculty of Medicine and University Hospital Cologne, University of Cologne, Center for Molecular Medicine Cologne (CMMC), Cologne, 50931, Germany.
- ¹³³ University of Malaya, Breast Cancer Research Unit, University Malaya Cancer Research Institute, Faculty of Medicine, Kuala Lumpur, 50603, Malaysia.
- ¹³⁴ Academia Sinica, Institute of Biomedical Sciences, Taipei, 115, Taiwan.
- ¹³⁵ China Medical University, School of Public Health, Taichung, 40402, Taiwan.
- ¹³⁶ Vanderbilt University School of Medicine, Division of Epidemiology, Department of Medicine, Vanderbilt Epidemiology Center, Vanderbilt-Ingram Cancer Center, Nashville, TN, 37232, USA.
- ¹³⁷ Netherlands Comprehensive Cancer Organisation (IKNL), Department of Research, Utrecht, 3511 DT, The Netherlands.
- ¹³⁸ University of Twente, Department of Health Technology and Service Research, Technical Medical Center, Enschede, 7522 NB, The Netherlands.
- ¹³⁹ University Hospital and German Cancer Research Center, National Center for Tumor Diseases, Heidelberg, 69120, Germany.
- ¹⁴⁰ The University of Melbourne, Department of Clinical Pathology, Melbourne, Victoria, 3010, Australia.

- ¹⁴¹ BC Cancer, Population Oncology, Vancouver, BC, V5Z 1G1, Canada.
- ¹⁴² University of British Columbia, School of Population and Public Health, Vancouver, BC, V6T 1Z4, Canada.
- ¹⁴³ Curtin University and University of Western Australia, The Curtin UWA Centre for Genetic Origins of Health and Disease, Perth, Western Australia, 6000, Australia.
- ¹⁴⁴ Kuopio University Hospital, Department of Oncology, Cancer Center, Kuopio, 70210, Finland.
- ¹⁴⁵ University of Eastern Finland, Institute of Clinical Medicine, Oncology, Kuopio, 70210, Finland.
- ¹⁴⁶ Cancer Research Malaysia, Breast Cancer Research Programme, Subang Jaya, Selangor, 47500, Malaysia.
- ¹⁴⁷ University of Malaya, Department of Surgery, Faculty of Medicine, Kuala Lumpur, 50603, Malaysia.
- ¹⁴⁸ Columbia University, Department of Epidemiology, Mailman School of Public Health, New York, NY, 10032, USA.
- ¹⁴⁹ Leiden University Medical Center, Department of Surgery, Leiden, 2333 ZA, The Netherlands.
- ¹⁵⁰ University of Birmingham, Institute of Cancer and Genomic Sciences, Birmingham, B15 2TT, UK.
- ¹⁵¹ University of Oxford, Wellcome Trust Centre for Human Genetics and Oxford NIHR Biomedical Research Centre, Oxford, OX3 7BN, UK.
- ¹⁵² Mayo Clinic, Department of Health Science Research, Division of Epidemiology, Rochester, MN, 55905, USA.
- ¹⁵³ Leuven Cancer Institute, University Hospitals Leuven, Leuven Multidisciplinary Breast Center, Department of Radiology, Leuven, 3000, Belgium.
- ¹⁵⁴ Uppsala University, Department of Surgical Sciences, Uppsala, 751 05, Sweden.

¹⁵⁵ The Netherlands Cancer Institute - Antoni van Leeuwenhoek hospital, Division of Psychosocial Research and Epidemiology, Amsterdam, 1066 CX, The Netherlands.

***Correspondence:** Marjanka K Schmidt, Netherlands Cancer Institute, Plesmanlaan 121, 1066 CX Amsterdam, The Netherlands; +31205122767; mk.schmidt@nki.nl

Abstract

Previous research has shown that polygenic risk scores (PRS) can be used to stratify women according to their risk of developing primary invasive breast cancer. This study aimed to evaluate the association between a recently validated PRS of 313 germline variants (PRS_{313}) and contralateral breast cancer (CBC) risk. We included 56,068 women of European ancestry diagnosed with first invasive breast cancer from 1990 onwards with follow-up from the Breast Cancer Association Consortium. Metachronous CBC risk ($N=1,027$) according to the distribution of the PRS_{313} was quantified using Cox regression analyses. We assessed PRS_{313} interaction with age at first diagnosis, family history, morphology, ER-, PR-, and HER2-status, and (neo)adjuvant therapy. In Asian studies, with limited follow-up, CBC risk associated with PRS_{313} was assessed using logistic regression for 340 women with CBC compared with 12,133 women with unilateral breast cancer. Higher PRS_{313} was associated with increased CBC risk: hazard ratio per standard deviation (SD)=1.25 (95%CI=1.18-1.33) for Europeans, and an OR per SD=1.15 (95%CI=1.02-1.29) for Asians. The absolute lifetime risks of CBC, accounting for death as competing risk, were 12.4% for European women at the 10th percentile and 20.5% at the 90th percentile of the PRS_{313} . We found no evidence of confounding by, or interaction with patient characteristics, characteristics of the primary tumor, or treatment. The C-index for the PRS_{313} alone was 0.563 (95%CI=0.547-0.586). In conclusion, the PRS_{313} is an independent factor associated with CBC risk, and may be incorporated in CBC risk prediction models to help improve stratification of patients and optimize surveillance and treatment strategies.

Introduction

Due to the high incidence of breast cancer and improving survival, an increasing number of breast cancer survivors are at risk of developing contralateral breast cancer (CBC). The 10-year cumulative incidence of CBC is ~4%^{1; 2}, however estimates vary widely depending on factors such as germline genetics, family history, and (neo)adjuvant systemic therapy for the first breast cancer³. The risk of developing CBC is particularly high in women with rare mutations in certain genes including *BRCA1*, *BRCA2*, and *CHEK2*, with approximately two- to fourfold higher risks reported compared with women without these mutations³.

Recently, genome-wide association studies (GWAS) have identified multiple common germline variants that are associated with first primary breast cancer risk^{4; 5}. These are associated with small differences in risk individually, but their combined effects can be summarized in a polygenic risk score (PRS), which has been shown to stratify women according to their risk of developing breast cancer⁶⁻⁹. Using a large GWAS dataset from the Breast Cancer Association Consortium (BCAC), we previously developed and validated a 313-variant PRS (PRS₃₁₃) among women of European descent. In independent prospective studies, this PRS₃₁₃ predicted the risk of primary invasive breast cancer with an odds ratio (OR) per standard deviation (SD) of 1.61 (95% confidence interval (95%CI)=1.57-1.65)⁷. The PRS₃₁₃ has also been externally validated using the UK Biobank cohort.

The aim of the current study was to evaluate the association between PRS₃₁₃ and CBC risk, using data from BCAC. Other studies have shown associations between risk of CBC and both a 67-variant PRS¹⁰ and individual variants¹¹, but not yet with PRS₃₁₃, the most extensively validated PRS. Further, the dataset currently evaluated is larger than those previously tested. We carried out two types of analyses. We conducted a cohort study among studies of European ancestry women with follow-up data available, and performed Cox regression analyses to

estimate hazard ratios (HRs) for CBC. Potential confounding and interaction with patient characteristics, characteristics of the primary tumor, or treatment were tested. In addition, to directly compare with the OR reported for PRS₃₁₃ and first breast cancer, we selected case-case series and performed logistic regression analyses comparing the PRS₃₁₃ distribution in women with CBC versus those with unilateral breast cancer. These analyses were conducted separately in European and Asian women (follow-up was too limited to perform a cohort study for the Asian population). Use of PRS₃₁₃ may lead to more accurate CBC risk prediction to support decision making for women who may or may not benefit from additional surveillance and risk-reducing treatment strategies.

Material and Methods

Study subjects

Case-case series

We selected women who were diagnosed with breast cancer and women without any diagnosis of breast cancer from the BCAC including all women of European ancestry, based on genotyping data, selecting only those studies which reported on CBC (62 studies) (Figure S1A, Table S1-S2). BCAC database version freeze 12 was used. All women diagnosed with invasive breast cancer as a first cancer were included in the analysis; the small number of tumors with unknown invasiveness were considered invasive (Table S2). In the case-case series, a CBC was defined as a breast cancer (in situ or invasive) in the contralateral breast irrespective of the time since the first breast cancer. The case-case series comprised 81,000 women with unilateral breast cancer, 3,607 women with CBC, and 62,830 women without any diagnosis of breast cancer (Figure S1A). We also compared women with unilateral breast cancer to women without any diagnosis of breast cancer to reproduce the estimate that was previously reported for first breast cancer risk⁷ in our study selection.

We selected for a separate analysis women of Asian ancestry of the BCAC data comprising 12,133 women with unilateral breast cancer, 340 women with CBC, and 13,398 women without any diagnosis of breast cancer from eight studies (Figure S1B, Table S2).

European cohort

In the European cohort we used metachronous CBC as the outcome, defined as a breast cancer in the contralateral breast (in situ or invasive) diagnosed at least three months after the first breast cancer. We used a cut-off of three months to reduce the likelihood that these CBCs represent metastases rather than true second primary tumors. We selected all women diagnosed with breast cancer from the European case-case series and excluded four studies

that did not provide follow-up information on vital status (Figure S1A). We did not include Asian women since follow-up was too limited in these studies. We additionally excluded 6,207 women with no follow-up and 2,208 women who developed synchronous CBC, distant metastasis, or who died or last known to be alive within three months after the first breast cancer diagnosis. Since BCAC also included prevalent cases, we excluded 3,796 women who developed CBC or were censored before study entry. The case-case series included women diagnosed between 1947 and 2018. In the European cohort, we excluded 2,235 women who were diagnosed with their first breast cancer before 1990 or who had missing year of first diagnosis. We restricted to women diagnosed from 1990 onwards so that diagnostic procedures and treatment would be more representative of current practice. Moreover, clinico-pathological, treatment and follow-up data were more complete after 1990. In addition, we excluded 16 studies (9,783 women) without information about metachronous CBC events (Figure S1A). After these exclusions, the cohort for this analysis comprised data from 42 studies, including 56,068 women with invasive breast cancer among whom 1,027 metachronous CBC occurred (Table S2).

All individuals provided written informed consent, and all studies were approved by the relevant institutional review boards. BCAC data were centrally harmonized and cleaned in communication with the study data managers and principal investigators. Data collection for individual studies is described in Table S1.

Genotyping and PRS

DNA samples from participants were genotyped using the iCOGS array^{12; 13} or the OncoArray^{4; 14}, with genotypes for variants not on the arrays estimated by imputation^{4; 13}. The PRS₃₁₃ was calculated as a weighted sum of the minor allele dosages; the variant selection and weights are as given by Mavaddat et al.⁷. We also calculated estimates for a previously published PRS₇₇⁶, and estrogen receptor (ER)-specific PRSs (ER-positive PRS₃₁₃ and ER-negative PRS₃₁₃)⁷. The

ER-specific PRSs were constructed by defining subtype-specific weights for the 313 variants using a hybrid approach⁷. Variants and corresponding coefficients used to construct the PRS are shown in Table S3. We standardized the PRS in our analyses by dividing it by the SD of the PRS of the controls (PRS₇₇ SD=0.45; PRS₃₁₃ SD=0.61; ER-positive PRS₃₁₃ SD=0.65; ER-negative PRS₃₁₃ SD=0.59) exactly as was done in the analyses of the PRS and first breast cancer risk^{6: 7}. This allows a direct comparison of the magnitude of the CBC relative risk estimation to that of the first breast cancer.

For samples genotyped with both OncoArray and iCOGS array (9,071 samples), OncoArray data were used in preference as the imputation quality was generally higher. The intraclass correlation coefficient (ICC) between the PRS derived from the two platforms was 0.99 (95%CI=0.99-0.99) for the PRS₇₇, and 0.96 (95%CI=0.95-0.96) for PRS₃₁₃ (Figure S2). Given the high correlation between the two platforms, PRS measures from both platforms were used in the analyses without adjustment.

Statistical analysis

European cohort

The primary outcome in the European cohort was the development of metachronous CBC. Cox proportional hazards models were used to estimate HRs for metachronous CBC risk by PRS, stratified by country. Since previous studies have shown that age at first breast cancer diagnosis is an important predictor of CBC³, the analyses were performed with attained age as the time scale. Time at risk started three months after the first breast cancer diagnosis and ended at the age of CBC diagnosis, distant metastasis (where available), death, or end of follow-up, whichever came first. For patients that had a study entry more than three months after first breast cancer diagnosis, follow-up started at the age of study entry. We also performed a fixed-effect meta-analysis of country-specific effects using the STATA command

metan. We performed a fixed-effect meta-analysis over a random-effect meta-analysis since there was no evidence for heterogeneity in effect sizes between countries (I-squared=0%, Figure S3). For some analyses, only invasive CBC was used as the outcome; in these analyses we censored on in situ CBC. Separate analyses were conducted for ER-positive CBC (censored on ER-negative- and ER-unknown CBC) and ER-negative CBC (censored on ER-positive- and ER-unknown CBC).

We evaluated the linearity of the association between PRS₃₁₃ per unit SD and CBC risk using restricted cubic splines with three knots. There was no evidence for violation of the linearity assumption. Therefore, in the main analysis, the PRS₃₁₃ was treated as a continuous covariate, and estimated the HR per unit SD of the PRS₃₁₃. Violation of the proportional hazard assumption was assessed by inspection of the Schoenfeld residuals¹⁵. As a second analysis, we used the per SD log HR of the PRS₃₁₃ to calculate the predicted HR at different percentiles of the PRS₃₁₃, compared to the 50th percentile. Third, the PRS₃₁₃ was categorized into percentile groups (0th to 10th, 10th to 20th, 20th to 40th, 40th to 60th, 60th to 80th, 80th to 90th, 90th to 100th) to illustrate the differences between PRS₃₁₃ subgroups, with the middle quintile (40th to 60th) as the reference.

We also performed multivariable Cox regression analyses to determine whether the log HR of CBC risk by PRS changed when adjusting for year of first breast cancer diagnosis, family history of breast cancer in a first degree relative, and several clinical characteristics of the first breast cancer such as nodal status, tumor size, morphology, ER-, progesterone receptor (PR)- and human epidermal growth factor receptor 2 (HER2)-status, (neo)adjuvant chemotherapy, adjuvant endocrine therapy, and radiotherapy. These analyses were performed in all patients, a complete case set (excluding patients with unknown values for the covariates), and in a set excluding studies oversampling cases with family history. Potential effect modification of the

PRS₃₁₃ effect by the same variables was evaluated by fitting interaction terms in different models using complete case sets, including the standardized PRS₃₁₃, modifier, and interaction.

The discriminative ability of different models; ([model 1] PRS₃₁₃ alone, [model 2] other risk factors (the adjustment variables from the multivariable Cox regression analyses), [model 3] PRS₃₁₃ + other risk factors) was calculated using Harrell's C-index¹⁶. Since no standard performance measures are currently available to account for left-truncated follow-up time (*i.e.*, to start analyses at age at study entry), we used time since first breast cancer as the time scale to calculate the C-index.

Absolute risks

Absolute risks of developing CBC at PRS₃₁₃ percentiles were calculated using the estimated log HRs per SD from the breast cancer cohort (BCAC) under the log-linear model, assuming the PRS is normally distributed. The PRS₃₁₃- and age-specific incidences were constrained to the age-specific CBC incidences from women diagnosed with a first invasive breast cancer in the period 2003-2010 from the Netherlands Cancer Registry (NCR)¹. The procedure for constraining the incidences has been previously described¹⁷. The age-specific CBC incidences were calculated overall and for age-specific groups, censoring on death and distant metastasis. We used data from the NCR since this registry has complete coverage of all newly diagnosed cancers in the Netherlands. The NCR cohort included all females aged ≥ 18 years and follow-up for second cancers was complete until February 1, 2016¹. We then applied the competing risk of dying on the absolute CBC risks. The absolute CBC risk (AR_g) by age t in PRS₃₁₃ category g , taking into account the competing risk of dying was calculated by:

$$AR_g(t) = \sum_{u=0}^{t-1} \mu_g(u)S_g(u)S_m(u)$$

Where $\mu_g(t)$ is the CBC incidence associated with PRS₃₁₃ category g , $S_g(t)$ the probability of being free of CBC to age t , and $S_m(t)$ the probability of surviving to age t .

Case-case series

For the case-case series (European and Asian), logistic regression models were used to estimate the ORs for CBC risk (comparing with unilateral breast cancer) and for unilateral breast cancer risk (comparing with women without any diagnosis of breast cancer) associated with PRS₃₁₃. All analyses were adjusted for age and country (Table S1). For all unilateral- and contralateral breast cancer patients we used age at first breast cancer diagnosis, and for women without any diagnosis of breast cancer we used age at baseline questionnaire.

For direct comparison with the estimate reported for PRS₃₁₃ and first breast cancer, we also performed logistic regression analyses in the same BCAC study participants included in the validation of the association between PRS₃₁₃ and first breast cancer risk⁷. This validation set comprised a subsample from 24 studies and included 3,781 women with unilateral breast cancer, 94 women with CBC, and 3,753 women without any diagnosis of breast cancer (Table S2). For this analysis, we adjusted for 10 principal components, in line with Mavaddat et al.⁷.

For European women who had follow-up time available more than three months after the first breast cancer diagnosis, a sensitivity analysis was performed for metachronous CBC (1,702 CBCs). We also did a separate analysis for invasive CBC (N=3,246), by excluding CBC in situ.

All P-values are two sided; tests with $P < .05$ are referred to as statistically significant. Analyses were performed using STATA, version 13.1 (StataCorp) and R version 3.3.2.

Results

European (cohort) Cox regression analyses

The European cohort included 56,068 women diagnosed with first invasive breast cancer with 1,027 metachronous CBC events. Median follow-up was 8.4 years. Patient, tumor, and treatment characteristics are summarized in Table S4.

The associations between the different PRSs and CBC risk are shown in Table 1. The HR for CBC per SD of PRS₃₁₃ was 1.25 (95%CI=1.18-1.33). For comparison, the HR per SD for PRS₇₇ was 1.21 (95%CI=1.14-1.29). Women within the 0th to 10th and the 90th to 100th percentile of the PRS₃₁₃ had 0.59-fold (95%CI=0.45-0.78) and 1.38-fold (95%CI=1.13-1.69) risks of CBC, respectively, compared with women within the 40th to 60th percentile (Figure 1, Table S5). The predicted HRs of CBC for women at the 10th and 90th percentile of the PRS₃₁₃ were 0.75 and 1.33, respectively, compared to the 50th percentile (Figure 1). Since we observed evidence of departure from the proportional hazards assumption ($P=0.02$)¹⁵, we also calculated HRs stratified for follow-up duration (<five and ≥five years). The HR by SD of the PRS₃₁₃ was 1.21 (95%CI=1.10-1.32) for CBC diagnosed ≤five years after first breast cancer diagnosis (CBC N=428), and 1.28 (95%CI=1.18-1.38) for CBC diagnosed >five years after first diagnosis (CBC N=599).

The HR per SD of PRS₃₁₃ for ER-positive invasive CBC was 1.38 (95%CI=1.23-1.55), compared to a HR per SD of the ER-positive PRS₃₁₃ of 1.37 (95%CI=1.22-1.54) (Table 1). For ER-negative invasive CBC, the HR per SD was 0.92 (95%CI=0.75-1.12) for PRS₃₁₃ and 1.06 (95%CI=0.86-1.30) for the ER-negative PRS₃₁₃.

Sensitivity analysis using the overall PRS₃₁₃ showed a HR per SD of 1.24 (95%CI=1.16-1.32) for invasive CBC risk. When we used time since first breast cancer as the time scale, we found

similar results (HR per SD=1.25, 95%CI=1.18-1.33). Meta-analysis of country-specific effects showed a HR per SD of 1.25 (95%CI=1.18-1.33) for CBC risk by PRS₃₁₃ (Figure S3).

The association between the PRS₃₁₃ and CBC risk did not change when adjusting for patient, tumor, and treatment characteristics, nor when excluding studies oversampling cases with a family history (Table S6). When considering potential modifiers of the effect of the PRS₃₁₃ on CBC risk (Table 2), we found that the HR was the lowest in women aged <40 years at first breast cancer diagnosis (HR per SD=1.13; 95%CI=0.98-1.31), and tended to increase with age, although these effects were not statistically significant ($P_{\text{heterogeneity}}=.26$; $P_{\text{trend}}=.05$). We found no indication for effect modification by family history ($P_{\text{heterogeneity}}=.63$), morphology ($P_{\text{heterogeneity}}=.14$), ER-status ($P_{\text{heterogeneity}}=.13$), PR-status ($P=.26$), HER2-status ($P_{\text{heterogeneity}}=.42$), chemotherapy ($P_{\text{heterogeneity}}=.60$), endocrine therapy ($P_{\text{heterogeneity}}=.79$), or radiotherapy ($P_{\text{heterogeneity}}=.40$) (Table 2).

The C-index was 0.563 (95%CI=0.547-0.586) for the model only including PRS₃₁₃, 0.605 (95%CI=0.591-0.629) for the model only including other risk factors, and 0.623 (95%CI=0.608-0.645) for the complete model (Table 3).

Absolute risks

Based on the HR estimates for PRS₃₁₃, the predicted CBC risk by age 80 years was 12.4% at the 10th percentile of the PRS₃₁₃, compared with 20.5% at the 90th percentile of the PRS₃₁₃ (Figure 2), accounting for death as competing risk. When death was not taken into account as competing risk, the corresponding predicted risks by age 80 were 17.0% at the 10th percentile and 27.9% at the 90th percentile of the PRS₃₁₃ (Figure S4). Table 4 shows the five- and 10-year cumulative CBC risks by PRS₃₁₃ for different age groups, accounting for death as competing risk (Table S7 shows results without competing risks).

European and Asian (case-case series) logistic regression analyses

Figure 3 shows the distribution of the PRS₃₁₃ per SD in the European case-case series. Median PRS₃₁₃ was -0.4 (interquartile range [IQR]=1.35) for control women without any diagnosis of breast cancer (N=81,000), 0.2 (IQR=1.36) for women with unilateral breast cancer (N=62,830), and 0.5 (IQR=1.40) for women with CBC (N=3,607). The OR for unilateral breast cancer per SD of the PRS₃₁₃, compared to control women, was 1.82 (95%CI=1.80-1.84) (Table S8). The OR for CBC per SD of PRS₃₁₃, compared to unilateral breast cancer, was 1.30 (95%CI=1.26-1.35) .

In sensitivity analyses, the OR per SD of PRS₃₁₃ was 1.27 (95%CI=1.21-1.33) for metachronous CBC and the OR per SD was 1.29 (95%CI=1.24-1.33) for invasive CBC, compared to unilateral breast cancer. When analyses were restricted to the validation set of Mavaddat et al⁷, the OR for unilateral breast cancer per SD of the PRS₃₁₃ was 1.67 (95%CI=1.59-1.76) compared to control women, and the OR for CBC per SD of PRS₃₁₃ was 1.39 (95%CI=1.13-1.70) compared to unilateral breast cancer (Table S8).

For women of Asian descent, the OR for unilateral breast cancer per SD of the PRS₃₁₃ was 1.56 (95%CI=1.52-1.60) compared to control women, and the OR for CBC per SD of PRS₃₁₃ was 1.15 (95%CI=1.02-1.29) compared to women with unilateral breast cancer (Table S8).

Discussion

Previous studies have shown that a PRS, summarizing the effects of common germline variants, can be used to stratify women with respect to their risk to develop a primary breast cancer⁶⁻⁹. In this study, we observed a clear association between the PRS₃₁₃ and CBC risk in women of both European and Asian ancestry. The association was observed in both the case-case series and the European cohort. The HRs per SD of CBC for women at the 10th and 90th percentile of the continuous predicted PRS₃₁₃ were 0.75 and 1.33, respectively, compared to the 50th percentile. This translates to absolute risks at the 10th and the 90th percentile of the PRS₃₁₃ of 12.4% and 20.5%, respectively, by age 80 years. We estimated a C-index for the PRS₃₁₃, summarizing its discriminatory ability, of 0.563 in the European cohort.

One previous study has investigated the effect of a PRS, including 67 variants, and CBC risk¹⁰. This study found a risk ratio of 1.75 (95%CI=1.41-2.18) for women in the upper quartile of the PRS compared with women in the lowest quartile. To facilitate comparison, we performed a similar analysis in our case-case series, showing an OR of 1.98 (95%CI=1.79-2.18), adjusted for country and age at first diagnosis, for women in the upper quartile of the PRS₃₁₃. This indicates the PRS₃₁₃ improves stratification relative to PRSs including fewer variants. Moreover, in our European cohort, the C-index for the PRS alone improved from 0.547 (95%CI=0.536-0.575) for the previously reported PRS₇₇⁶ to 0.563 (95%CI=0.547-0.586) for the PRS₃₁₃.

We found no evidence that the association between the PRS₃₁₃ and CBC risk was confounded by family history, adjuvant therapy, morphology, age, or tumor receptor status of the first breast cancer, nor that there was effect modification by those factors. The absence of notable effect modification is in line with the abovementioned study of a 67-variant PRS and CBC risk; no heterogeneity in association was found by age, family history, morphology, ER-status, and adjuvant treatment¹⁰.

To provide an external validation of our findings, we examined data from UK Biobank, which includes many women diagnosed with breast cancer with data available on the PRS₃₁₃ (Supplemental Note). Unfortunately, UK Biobank has no information available on the laterality of the tumor, and it is, therefore, not possible to distinguish between contralateral and ipsilateral breast cancers. We therefore performed analyses using any second breast cancer as the endpoint. This secondary analysis did confirm the association between the PRS₃₁₃ and second breast cancer risk (HR per SD=1.13, 95%CI=1.01-1.27), but with a lower estimate than in our European cohort. The lower estimate may be explained by the inclusion of the ipsilateral breast cancers, which may be more likely to be recurrences than new primary breast cancers compared to CBCs. Indeed, when we used ipsilateral breast cancer as the outcome in our European cohort, we found no association with the PRS₃₁₃ (HR=1.02, 95%CI=0.90-1.15).

The association between the PRS₃₁₃ and CBC risk (OR per SD=1.30; 95%CI=1.26-1.35) in the BCAC database was weaker (expressed in terms of an OR) than was found for first breast cancer among independent prospective studies (OR per SD=1.61; 95%CI=1.57-1.65). Under a simple polygenic model, the relative risk would be expected to be similar for the second breast cancer. The attenuated estimate for CBC might however be explained by several factors. Some attenuation of the estimate might have been due to dilution in the end-point definition, *i.e.*, if some of the CBCs were metastases. Previous studies investigating the clonal relatedness of first breast cancers and CBCs using tumor sequencing have shown that 6-12% of CBCs represent metastases^{18; 19}. This hypothesis would be consistent with our finding of a slightly stronger association between the PRS₃₁₃ and late CBCs, diagnosed >five years after the first breast cancer, than for early CBCs, diagnosed ≤five years after the first cancer, since the latter are more likely to be metastases. In addition, 3-5% of the breast cancer patients will have a mutation in the *BRCA1* or *BRCA2* gene^{20; 21}, who have high CBC risks. It has been shown that

the relative risk associated with PRS is lower (for the first breast cancer) for women with a *BRCA1* and *BRCA2* mutation than in the general population²², diluting the overall relative risk for CBC. More generally, it is possible that the CBC association may be attenuated due to the effect of other, unmeasured, genetic or other risk factors. If the risks are high, cases with higher PRS₃₁₃ will have, on average, lower values of other risk factors, due to elimination of the highest risk individuals, again attenuating the CBC association. Finally, given the limited information on family history in our dataset, the estimate could have been biased due to a family history effect not detected in our data.

There was some suggestion that the relative risk associated with PRS₃₁₃ decreased with younger age, ($P_{\text{trend}}=.05$), and, specifically, was lower for women aged <40 years (HR per SD=1.13; 95%CI=0.98-1.31). Interestingly, Mavaddat et al⁷ also found a lower relative risk below age 40 for first breast cancer. This effect may reflect the different characteristics of breast cancers at young ages, both in terms of germline susceptibility and pathology^{23; 24}. For example, the proportion of ER-negative breast cancers is higher at young ages, and the PRS is less predictive for ER-negative disease^{6; 7; 24}.

In the logistic regression analyses in Asian women, the association between the PRS₃₁₃ and CBC risk was slightly weaker than in European women. This finding is consistent with a recent analysis investigating the association between a 287-variant PRS and first breast cancer risk in the Asian population²⁵, which showed an attenuated OR in Asian women (OR=1.52, 95%CI=1.49-1.56) compared to European women (OR=1.61, 95%CI=1.57-1.66). The lower estimate for Asian women might reflect the fact the PRS₃₁₃ was developed in European populations, and the different LD structure in Asians may attenuate the association since the variants in the PRS are likely to be surrogates for the causal variants. Other explanations for the attenuated estimate may be the slightly younger age at first breast cancer diagnosis and the

higher proportion ER-negative CBCs in Asian women compared to European women in our study. Finally, the imputation quality for variants was somewhat lower, on average, for the Asian than for the European dataset, with three variants on OncoArray and four variants on ICOGs with an imputation quality score <0.3 (Table S3). Nevertheless, we included those variants in the PRS for both European and Asian women, to keep the PRS comparable between ethnicities and studies. Future studies including larger numbers of Asian women, and women of other ethnicities, are needed to generate population-specific PRSs and to validate our findings in these groups.

A major strength of this study is the very large sample size in the BCAC dataset, including genotype information for ~150,000 women and a large number of CBC events. A limitation of this study is missing data on the patient, tumor, and treatment characteristics, which reduces the power of the multivariable Cox regression analyses and interaction analyses. In addition, registration of CBC was not complete; the 10-year cumulative CBC incidence was 2.2% in the BCAC dataset, compared to 3.8% using complete data from the Netherlands Cancer Registry¹. For this reason, we estimated relative risk estimates using the BCAC data and applied these to external registry data to obtain absolute risk estimates. The underreporting of CBC should not bias our HR estimates, given that the event rate is low and reporting of CBC is unlikely to be related to the PRS₃₁₃. Moreover, we reran the cohort analysis in the subset of countries with a 10-year cumulative CBC incidence $\geq 3.0\%$ in the BCAC dataset, and the estimates were very similar to the main analyses (HR per SD=1.23, 95%CI=1.14-1.33) (Figure S3).

In conclusion, the PRS₃₁₃ is predictive for the development of CBC. We found no evidence for confounding or effect modification by other previously established CBC risk factors. The PRS₃₁₃ is therefore likely to be an independent risk factor for CBC. Since the predictive ability of the PRS on its own is modest, it should be combined with other breast cancer risk factors to provide

more useful CBC risk prediction models. More accurate risk prediction will help identify women at high CBC risk who will benefit from additional surveillance and/or risk reducing mastectomy, and equally important, to identify those women at low risk in order to avoid unnecessary surgeries.

Supplemental Data

Supplemental data include four figures, eight tables, Supplemental Note and acknowledgements.

Data and Code Availability

Data used in this manuscript may be requested through the original providers. Data of the Breast Cancer Association Consortium may be requested for non-profit research through an application procedure with the Breast Cancer Association Consortium; more information: <http://bcac.ccge.medschl.cam.ac.uk/bcacdata/>. Data of the UK Biobank needs to be requested through UK Biobank; more information: <https://www.ukbiobank.ac.uk/researchers/>

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Declaration of Interests

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Figure 1. Estimates for contralateral breast cancer risk by percentile categories of the 313-variant PRS (PRS₃₁₃)

The figure shows the hazard ratios per SD and 95% confidence intervals for percentiles of the PRS₃₁₃ relative to the middle quintile (underlying table can be found in Table S5). The solid line denotes the estimates for contralateral breast cancer risk with the PRS₃₁₃ fitted as a continuous covariate. Coefficients to construct the PRS₃₁₃ are shown in Table S3. The PRS₃₁₃ was standardized by SD=0.61, in line with Mavaddat et al.⁷. The analyses were performed with attained age as time scale. PRS = polygenic risk score, SD = standard deviation

Figure 2. Predicted contralateral breast cancer risk by percentile of the 313-variant PRS (PRS₃₁₃) with death as competing risk

Coefficients to construct the PRS₃₁₃ are shown in Table S3. The PRS₃₁₃ was standardized by SD=0.61, in line with Mavaddat et al.⁷ The CBC incidences were calculated based on incidence data from the Netherlands Cancer Registry¹ and relative risks estimated as described in the Material and Methods. PRS = polygenic risk score, CBC = contralateral breast cancer

Figure 3. Distribution of the 313-variant PRS (PRS₃₁₃) in 62,830 control women without any diagnosis of breast cancer, 81,000 women with unilateral breast cancer, and 3,607 women with contralateral breast cancer

Coefficients to construct the PRS₃₁₃ are shown in Table S3. The PRS₃₁₃ was standardized by SD=0.61, in line with Mavaddat et al.⁷. PRS = polygenic risk score, BC = breast cancer, CBC = contralateral breast cancer, SD = standard deviation

Table 1. Association between PRSs and contralateral breast cancer risk in the European cohort (N=56,068)

Polygenic risk score (PRS)	No. of CBC	HR per unit SD ^a	95%CI	P-value
PRS₇₇^b				
All CBC	1,027	1.21	1.14-1.29	<.001
Invasive CBC	923	1.21	1.13-1.29	<.001
PRS₃₁₃^b				
All CBC	1,027	1.25	1.18-1.33	<.001
Invasive CBC	923	1.24	1.16-1.32	<.001
ER-positive invasive CBC ^d	275	1.38	1.23-1.55	<.001
ER-negative invasive CBC ^d	97	0.92	0.75-1.12	.39
ER-positive PRS₃₁₃^{b,c}				
All CBC	1,027	1.23	1.16-1.31	<.001
Invasive CBC	923	1.22	1.15-1.30	<.001
ER-positive invasive CBC ^d	275	1.37	1.22-1.54	<.001
ER-negative PRS₃₁₃^{b,c}				
All CBC	1,027	1.25	1.17-1.33	<.001
Invasive CBC	923	1.24	1.16-1.33	<.001
ER-negative invasive CBC ^d	97	1.06	0.86-1.30	.58

Abbreviations: PRS = polygenic risk score, No. = number, CBC = contralateral breast cancer, HR = hazard ratio, CI = confidence interval, ER = estrogen receptor, SD = standard deviation

^a All analyses were performed with attained age as time scale

^b Coefficients to construct the PRSs are shown in Table S3. All PRSs were standardized by the same SD as was used by Mavaddat et al.⁷. The SD was 0.45 for overall breast cancer PRS₇₇, 0.61 for overall breast cancer PRS₃₁₃, 0.65 for ER-positive PRS₃₁₃, and 0.59 for ER-negative PRS₃₁₃

^c ER-specific PRSs were constructed using a hybrid method, as described by Mavaddat et al.⁷

^d Patients with ER-unknown CBC (N=551) were censored in these analyses

Table 2. Association between the 313-variant PRS (PRS₃₁₃) and contralateral breast cancer risk for subgroups

Subgroups	No. of patients	No. of CBC	HR per unit SD ^{a,b}	95%CI	P-value	P _{heterogeneity} ^{c,d}	P _{trend} ^{c,e}
All patients	56,068	1,027	1.25	1.18-1.33	<.001	-	-
Age at first breast cancer diagnosis (years)						.26	.05
<40	5,877	171	1.13	0.98-1.31	.09		
40-49	11,928	265	1.25	1.11-1.41	<.001		
50-59	16,882	320	1.22	1.09-1.36	<.001		
60+	21,381	271	1.36	1.21-1.52	<.001		
Family history (first degree relative)						.63	-
no	33,623	618	1.26	1.16-1.36	<.001		
yes	10,369	302	1.22	1.09-1.36	<.001		
Morphology						.14	-
ductal	37,324	621	1.21	1.12-1.31	<.001		
lobular	5,878	118	1.32	1.10-1.59	.002		
mixed (ductal and lobular)	2,174	46	1.52	1.15-2.02	.004		
other	3,344	70	1.20	0.96-1.50	.11		
ER-status						.13	-
negative	9,527	194	1.13	0.98-1.30	.08		
positive	38,090	670	1.28	1.19-1.38	<.001		
PR-status						.26	-
negative	13,098	244	1.16	1.03-1.32	.02		
positive	27,044	554	1.27	1.17-1.38	<.001		
HER2-status						.42	-
negative	23,787	352	1.29	1.17-1.44	<.001		
positive	4,969	60	1.45	1.13-1.85	.004		
(Neo)adjuvant chemotherapy						.60	-
no	18,110	361	1.28	1.16-1.42	<.001		
yes	18,559	363	1.24	1.12-1.37	<.001		
(Neo)adjuvant endocrine therapy						.79	-
no	10,781	242	1.28	1.13-1.44	<.001		
yes	27,322	460	1.30	1.19-1.43	<.001		
Radiotherapy						.40	-
no	11,023	188	1.33	1.15-1.53	<.001		
yes	29,142	617	1.24	1.15-1.34	<.001		

Abbreviations: PRS = polygenic risk score, No. = number, CBC = contralateral breast cancer, HR = hazard ratio, CI = confidence interval, ER = estrogen receptor, PR = progesterone receptor, HER2 = human epidermal growth factor receptor 2

^a HR for CBC risk by unit SD of PRS₃₁₃. All analyses were performed with attained age as time scale

^b Coefficients to construct the PRS₃₁₃ are shown in Table S3. The PRS₃₁₃ was standardized by standard deviation=0.61, in line with Mavaddat et al.⁷

^c The interaction between the PRS₃₁₃ and each subgroup was tested in different models including the standardized PRS₃₁₃, modifier, and interaction. Patients with unknown values were excluded from these analyses. Since attained age was used as time scale in all models, the model with age at first breast cancer only included the PRS₃₁₃ and interaction

^d P for interaction based on test for heterogeneity across categories

^e P for interaction based on a trend test with age as continuous variable

Table 3. Discriminatory ability (C-index) of the 313-variant PRS (PRS₃₁₃) and other risk factors for contralateral breast cancer risk in the European cohort

	C-index (95%CI) ^{a,b}
<i>Model 1</i> PRS ₃₁₃ ^c alone	0.563 (0.547-0.586)
<i>Model 2</i> Other risk factors ^d	0.605 (0.591-0.629)
<i>Model 3</i> PRS ₃₁₃ ^c + other risk factors ^d	0.623 (0.608-0.645)

Abbreviations: PRS = polygenic risk score, CI = confidence interval

^a The Harrell's C-index was obtained by the STATA stcox postestimation command 'estat concordance', using time since first breast cancer on the time scale without taking delayed entry (prevalent cases) into account. We did not consider delayed-entry since no standard performance measures are currently available in the statistical literature to account for left-truncated follow-up time. The median of delayed entry was 0.4 years (standard deviation=2.7) in our study

^b The 95% CIs were obtained by use of the 'somersd' package in STATA

^c Coefficients to construct the PRS₃₁₃ are shown in Table S3. The PRS₃₁₃ was standardized by SD=0.61, in line with Mavaddat et al.⁷

^d Including age at first diagnosis, year of first diagnosis, family history for breast cancer in a first degree relative, and clinical characteristics of the first breast cancer (nodal status, tumor size, differentiation grade, morphology, estrogen receptor status, human epidermal growth factor receptor 2 status, chemotherapy, endocrine therapy, radiotherapy)

Table 4. Five- and ten-year cumulative risks of contralateral breast cancer by the 313-variant PRS (PRS₃₁₃) for different age groups with death as competing risk

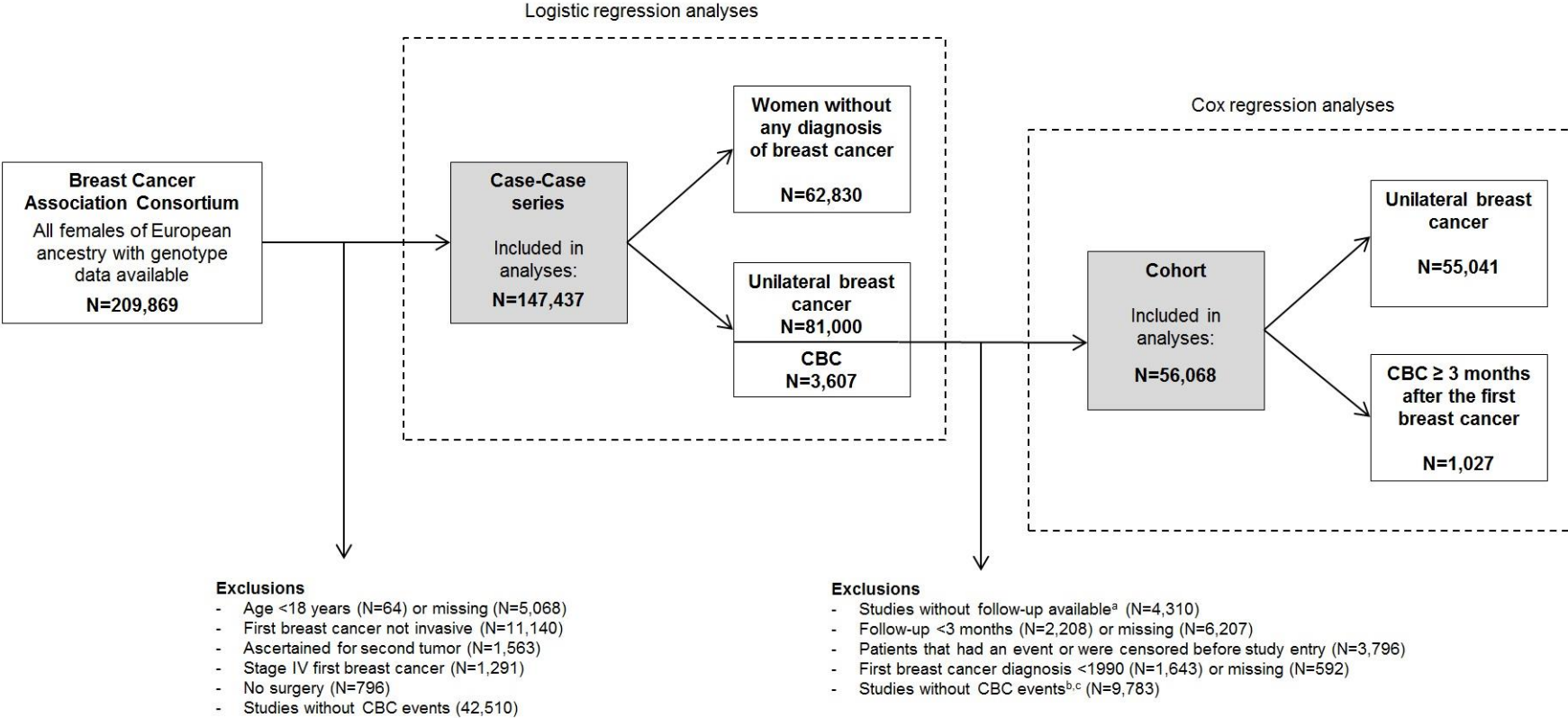
Age at first breast cancer diagnosis (years)	5-year cumulative CBC risks (%) range by age					10-year cumulative CBC risks (%) range by age				
	5 th percentile PRS ₃₁₃	10 th percentile PRS ₃₁₃	50 th percentile PRS ₃₁₃	90 th percentile PRS ₃₁₃	95 th percentile PRS ₃₁₃	5 th percentile PRS ₃₁₃	10 th percentile PRS ₃₁₃	50 th percentile PRS ₃₁₃	90 th percentile PRS ₃₁₃	95 th percentile PRS ₃₁₃
30-34	1.9-3.1	2.1-3.4	2.7-4.5	3.6-5.9	4.0-6.5	3.1-4.1	3.4-4.5	4.5-5.9	5.9-7.7	6.5-8.5
35-39	0.8-2.1	0.9-2.3	1.2-3.0	1.5-3.9	1.7-4.3	2.1-3.5	2.3-3.8	3.0-5.0	3.9-6.6	4.3-7.2
40-44	1.5-2.8	1.7-3.1	2.2-4.1	2.9-5.3	3.2-5.9	2.8-4.6	3.1-5.0	4.1-6.6	5.3-8.6	5.9-9.4
45-49	1.4-2.5	1.5-2.7	2.0-3.6	2.6-4.7	2.9-5.2	2.5-3.9	2.7-4.3	3.6-5.6	4.7-7.4	5.2-8.1
50-54	1.4-2.8	1.5-3.0	1.9-4.0	2.6-5.2	2.8-5.8	2.8-4.5	3.0-4.9	4.0-6.4	5.2-8.4	5.8-9.3
55-59	1.6-3.1	1.8-3.4	2.3-4.5	3.1-5.9	3.4-6.5	3.1-4.8	3.4-5.2	4.5-6.9	5.9-9.0	6.5-9.9
60-64	1.7-3.3	1.9-3.6	2.5-4.7	3.3-6.2	3.6-6.8	3.3-5.0	3.6-5.4	4.7-7.1	6.2-9.3	6.8-10.2
65-70	1.5-3.2	1.6-3.5	2.1-4.6	2.8-6.1	3.1-6.7	3.2-4.1	3.5-4.5	4.6-5.9	6.1-7.7	6.7-8.5

Abbreviations: PRS = polygenic risk score, CBC = contralateral breast cancer

Coefficients to construct the PRS₃₁₃ are shown in Table S3. The PRS₃₁₃ was standardized by SD=0.61, in line with Mavaddat et al⁷. The CBC incidences for each age group were calculated based on incidence data from the Netherlands Cancer Registry¹ and relative risks estimated as described in the Material and Methods. Death was taken into account as competing risk.

Supplemental Figures

Figure S1A. Overview of the selection of women with breast cancer and control women for the European series



Abbreviations: CBC = contralateral breast cancer

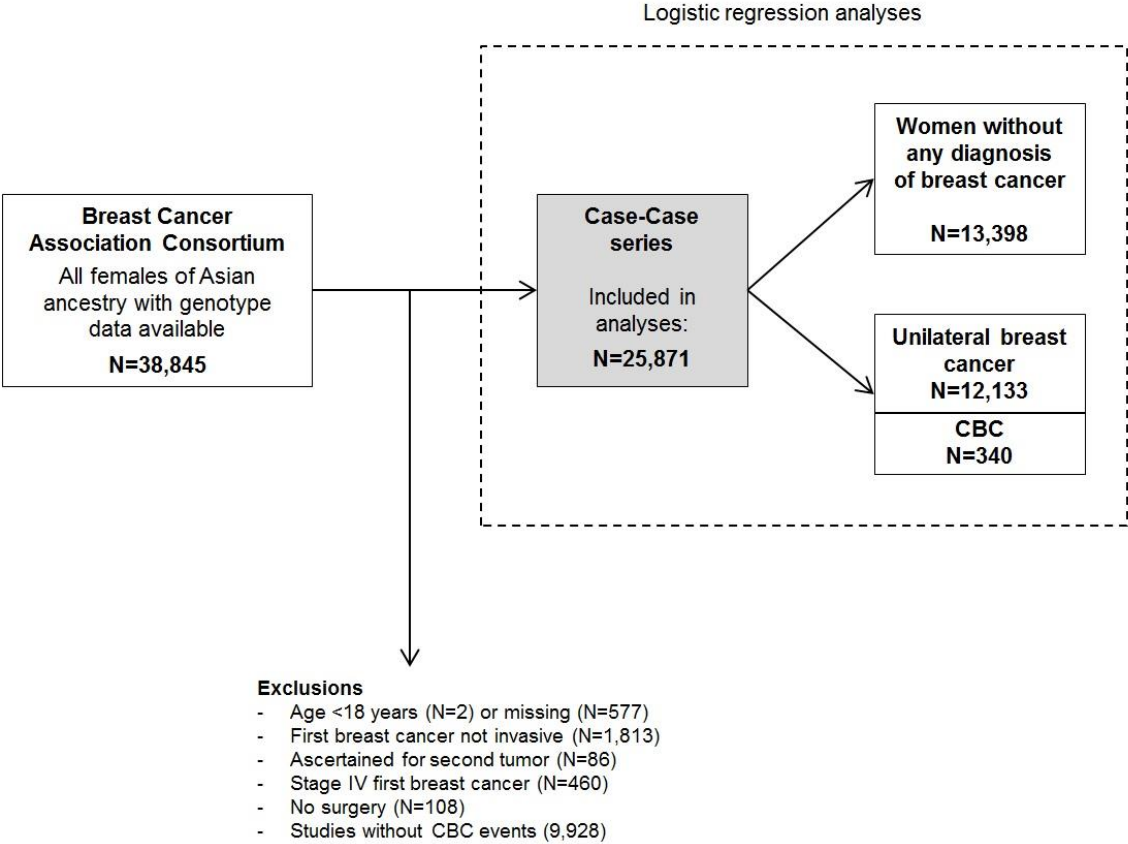
For a complete overview of all studies see Table S1

^a Excluded studies: CBCS, GLACIER, HMBCS, TNBCC

^b Excluded studies: BCFR-NY, BCFR-UTAH, CNIO-BCS, DIETCOMPLYF, FHRISK, GESBC, HABCS, HUBCS, ICICLE, KBCP, MCCS, MMHS, NCBCS, PREFACE, SUCCESSB, SUCCESSC

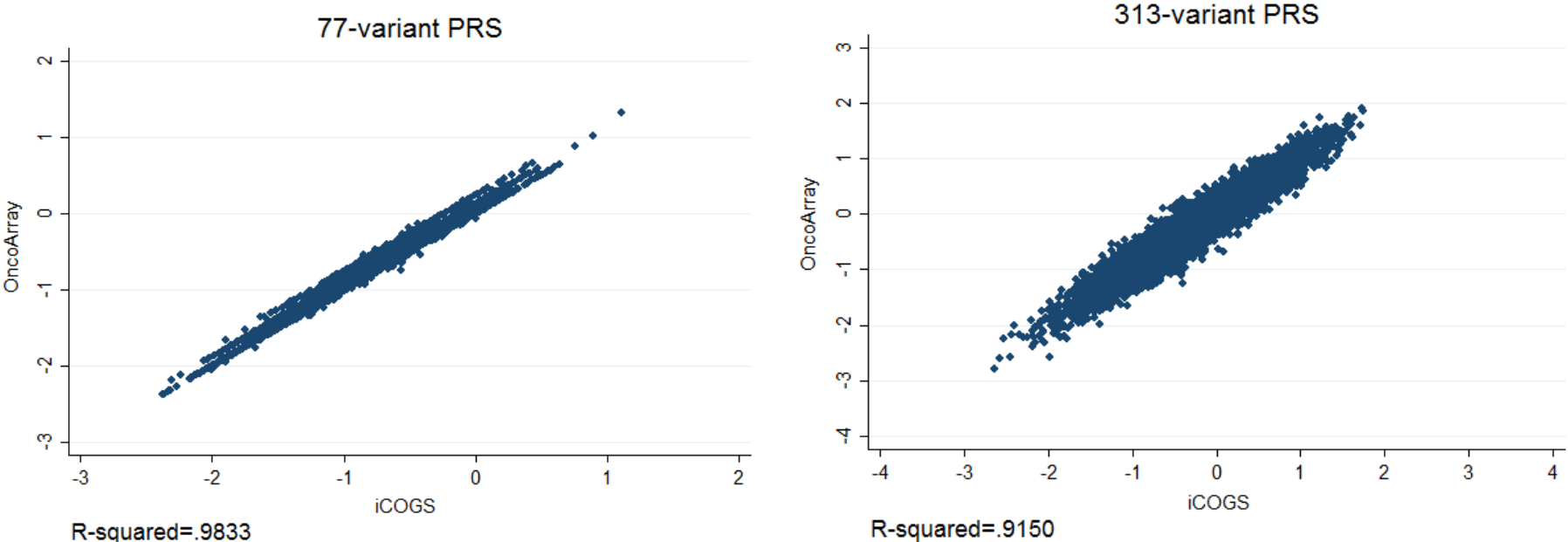
^c These studies dropped out because for these analyses the definition of CBC is based on the criteria that the CBC was diagnosed at least three months after the first breast cancer diagnosis

Figure S1B. Overview of the selection of women with breast cancer and control women for Asian series



Abbreviations: CBC = contralateral breast cancer

Figure S2. Correlation of total variant scores between the iCOGS array and OncoArray for the 77-variant PRS and the 313-variant PRS^{a,b}

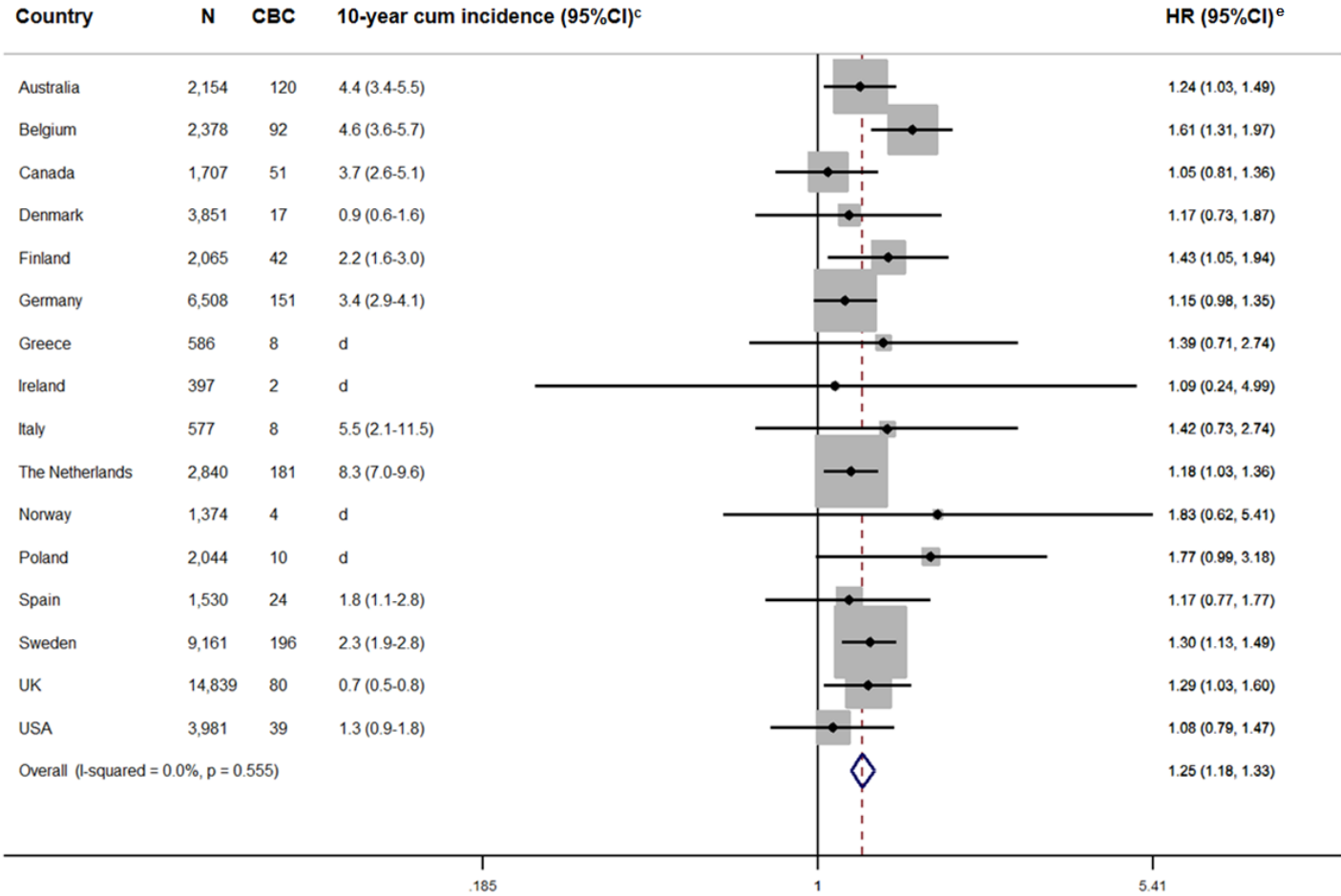


Abbreviations: PRS = polygenic risk score, SD = standard deviation

^aWe evaluated consistency between iCOGS and OncoArray using the intraclass correlation coefficient (ICC), showing a ICC of 0.99 (95%CI=0.99-0.99) for the PRS₇₇, and an ICC of 0.96 (95%CI=0.95-0.96) for the PRS₃₁₃, based on N=9,071 observations

^bCoefficients to construct the PRSs are shown in Table S3. The PRSs were standardized by the same SD as was used by Mavaddat et al.¹. The SD was 0.45 for overall breast cancer PRS₇₇, and 0.61 for overall breast cancer PRS₃₁₃

Figure S3. Forest plot of the association between the 313-variant PRS and contralateral breast cancer risk by country^{a,b}



Abbreviations: PRS = polygenic risk score, N = number of women, CBC = contralateral breast cancer, cum = cumulative, CI = confidence interval, HR = hazard ratio, SD = standard deviation

Fixed effect meta-analysis was used to calculate I-squared and P-value for heterogeneity

^a Republic of North Macedonia was left out this plot because of a too small sample size (N=76 women including N=2 CBC events)

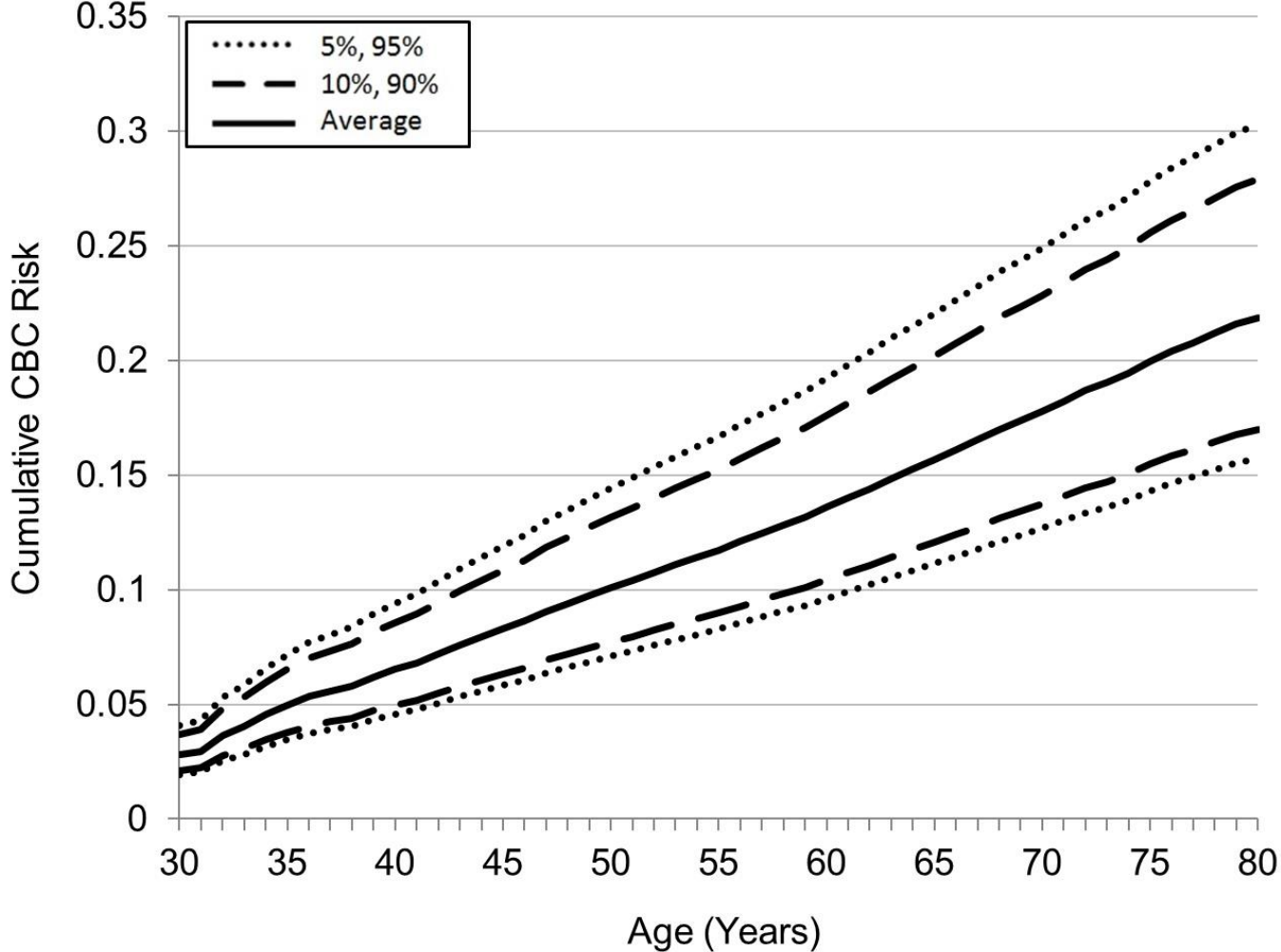
^b Coefficients to construct the PRS₃₁₃ are shown in Table S3. The PRS₃₁₃ was standardized by SD=0.61, in line with Mavaddat et al.¹

^c The 10-year cumulative incidence of CBC was estimated with time since first breast cancer as time scale, and distant metastases (where available) and death as competing risks

^d Follow-up too short for calculating 10-year cumulative incidence

^e HR per SD. The analyses were performed with attained age as the time scale

Figure S4. Predicted contralateral breast cancer risk by percentile of the 313-variant PRS (PRS₃₁₃)



Abbreviations: PRS = polygenic risk score, CBC = contralateral breast cancer
Coefficients to construct the PRS₃₁₃ are shown in Table S3. The PRS₃₁₃ was standardized by SD=0.61, in line with Mavaddat et al¹. The CBC incidences were calculated based on incidence data from the Netherlands Cancer Registry² and relative risks estimated as described in the Material and Methods. In contrast to Figure 2, death was not taken into account as competing risk.

Supplemental Tables

Table S1. Study characteristics of included studies of the Breast Cancer Association Consortium

Table S2. Studies and samples included in the analyses using the case-case series, cohort, and validation set

Studies	European									Asian		
	Case-case series N studies = 62			Cohort N studies = 42		Validation set N studies = 24			Case-case series N studies = 8			
	Control women ^a	Unilateral BC	CBC	Unilateral BC	CBC	Control women ^a	Unilateral BC	CBC	Control women ^a	Unilateral BC	CBC	
ABCFS	738	1,149	127	1,021	93	-	-	-	-	-	-	
ABCS	1,567	1,047	54	519	14	-	-	-	-	-	-	
ABCS-F	0	861	91	363	17	-	-	-	-	-	-	
ABCTB	375	900	17	708	1	74	180	8	-	-	-	
BBCC	711	845	58	766	6	49	56	5	-	-	-	
BBCS	1,768	1,266	80	466	1	-	-	-	-	-	-	
BCEES	-	-	-	-	-	166	133	0	-	-	-	
BCFR-NY	27	340	61	-	-	-	-	-	-	-	-	
BCFR-PA	0	104	14	69	4	-	-	-	-	-	-	
BCFR-UTAH	0	13	87	-	-	-	-	-	-	-	-	
BCINIS	-	-	-	-	-	144	262	0	-	-	-	
BIGGS	49	713	50	395	2	-	-	-	-	-	-	
BREOGAN	725	1,245	19	1,233	15	145	238	4	-	-	-	
BSUCH	1,122	900	36	727	3	-	-	-	-	-	-	
CBCS	817	530	21	-	-	163	105	4	170	238	10	
CCGP	321	598	19	578	8	66	125	7	-	-	-	
CGPS	5,250	4,135	60	3,834	17	142	227	3	-	-	-	
CNIO-BCS	829	742	5	-	-	-	-	-	-	-	-	
CTS	-	-	-	-	-	115	220	0	-	-	-	
DIETCOMPLYF	0	704	1	-	-	-	-	-	-	-	-	
FHRISK	0	119	2	-	-	-	-	-	-	-	-	
GC-HBOC	1,732	2,690	230	1,406	47	-	-	-	-	-	-	
GENICA	711	869	26	869	1	56	89	2	-	-	-	
GESBC	181	303	3	-	-	-	-	-	-	-	-	
GLACIER	0	1,733	230	-	-	-	-	-	-	-	-	
HABCS	863	774	84	-	-	173	141	6	-	-	-	
HCSC	0	362	13	273	9	-	-	-	-	-	-	
HEBCS	1,060	1,632	116	1,578	41	-	-	-	-	-	-	
HERPACC	-	-	-	-	-	-	-	-	1,659	756	18	
HKBCS	-	-	-	-	-	-	-	-	451	403	12	
HMBCS	345	729	28	-	-	-	-	-	-	-	-	
HUBCS	116	198	2	-	-	-	-	-	-	-	-	
ICICLE	1	138	12	-	-	-	-	-	-	-	-	
KARBAC	0	761	46	443	32	-	-	-	-	-	-	
KARMA	5,981	2,314	96	2,188	33	597	185	10	-	-	-	
KBCP	431	516	9	-	-	-	-	-	-	-	-	
KCONFAB/AOCS	898	397	83	305	26	-	-	-	-	-	-	
LMBC	1,821	3,016	208	2,286	92	87	142	14	-	-	-	
MABCS	88	80	9	74	2	-	-	-	-	-	-	

MARIE	2,066	1,540	115	1,535	53	-	-	-	-	-	-	
MBCSG	766	1,015	150	569	8	-	-	-	-	-	-	
MCBCS	2,093	1,999	59	1,903	6	35	96	3	-	-	-	
MCCS	1,207	1,034	2	-	-	142	86	0	-	-	-	
MEC	1,123	1,016	38	988	23	-	-	-	-	-	-	
MISS	1,529	582	6	563	3	304	83	0	-	-	-	
MMHS	1,635	273	4	-	-	320	48	4	-	-	-	
MYBRCA	-	-	-	-	-	-	-	-	4,197	3,652	105	
NBCS	212	2,334	31	1,370	4	-	-	-	-	-	-	
NBHS	-	-	-	-	-	122	79	0	-	-	-	
NC-BCFR	150	614	69	602	5	-	-	-	52	391	33	
NCBCS	1,006	1,988	42	-	-	-	-	-	-	-	-	
OBCS	414	467	10	445	1	-	-	-	-	-	-	
OFBCR	728	1,908	143	1,656	51	-	-	-	-	-	-	
ORIGO	0	1,090	89	1,053	69	132	134	15	-	-	-	
PBCS	2,082	1,719	40	1,625	9	331	215	2	-	-	-	
PKARMA	5,435	4,81	277	4,685	124	1	4	0	-	-	-	
POSH	0	1,069	19	1,063	16	-	-	-	-	-	-	
PREFACE	0	2,73	90	-	-	-	-	-	-	-	-	
PROCAS	1,647	488	9	422	3	-	-	-	-	-	-	
RBCS	0	873	152	724	81	-	-	-	-	-	-	
SASBAC	1,378	1,118	22	1,086	5	-	-	-	-	-	-	
SBCS	848	748	14	691	1	-	-	-	-	-	-	
SEARCH	9,056	12,423	118	12,117	59	197	628	0	-	-	-	
SEBCS	-	-	-	-	-	-	-	-	2,236	2,080	21	
SGBCC	-	-	-	-	-	-	-	-	4,141	1,250	124	
SKKDKFZS	29	1,084	71	1,054	41	-	-	-	-	-	-	
SMC	-	-	-	-	-	141	244	0	-	-	-	
SUCCESSB	0	438	2	-	-	-	-	-	-	-	-	
SUCCESSC	0	2,807	29	-	-	-	-	-	-	-	-	
SZBCS	489	676	6	409	1	-	-	-	-	-	-	
TNBCC	152	1,037	2	-	-	-	-	-	-	-	-	
TWBCS	-	-	-	-	-	-	-	-	492	1,250	17	
UCIBCS	258	397	1	380	1	51	61	7	-	-	-	
Total	62,830	81,000	3,607	55,041	1,027	3,753	3,781	94	13,398	12,133	340	
Characteristics												
Invasiveness	in situ	-	excluded	361	excluded	104	-	3 ^b	7	-	excluded	67
	invasive	-	79,876	2,200	54,675	670	-	3,777	60	-	11,929	209
	unknown	-	1,124	1,046	366	253	-	1	27	-	204	64
ER status	negative	-	13,828	446	9,333	105	-	766	8	-	3,457	54
	positive	-	52,238	2,048	37,420	289	-	3,001	47	-	7,826	163
	unknown	-	14,934	1,113	8,288	633	-	14	39	-	850	123

Abbreviations: BC = breast cancer, CBC = contralateral breast cancer, ER = estrogen receptor

^a Without any diagnosis of breast cancer

^b Due to the use of a new freeze of the BCAC data, N=3 breast cancers were now defined as in situ, which had previously been defined as invasive; the original validation dataset contained data of two additional studies¹

Table S3. Variant information and breast cancer risk coefficients for the 77-variant PRS, 313-variant PRS, and ER-specific PRSs; previously published in Mavaddat et al.^{1;3}

Table S4. Patient, tumor, and treatment characteristics of all women diagnosed with first invasive breast cancer since 1990 (European cohort)

Characteristics	Number of women (%)^a
Total	56,068 (100)
Median age at first diagnosis in years (range)	56 (18-98)
Year of diagnosis	
1990-1994	3,029 (5.4)
1995-1999	10,153 (18.1)
2000-2004	18,484 (33.0)
2005-2009	17,575 (31.3)
2010-2015	6,827 (12.2)
Family history (first degree relative)	
no	33,623 (76.4)
yes	10,369 (23.6)
unknown	12,076
Nodal status	
negative	29,070 (61.9)
positive	17,903 (38.1)
unknown	9,095
Tumor size, cm	
≤2	28,057 (63.8)
(2, 5]	14,138 (32.2)
>5	1,750 (4.0)
unknown	12,123
Differentiation grade	
I	8,721 (19.5)
II	21,621 (48.3)
III	14,454 (32.3)
unknown	11,272
Morphology	
ductal	37,324 (76.6)
lobular	5,878 (12.1)
mixed (ductal and lobular)	2,174 (4.5)
other	3,344 (6.9)
unknown	7,348

ER-status		
	negative	9,527 (20.0)
	positive	38,090 (80.0)
	unknown	8,451
PR-status		
	negative	13,098 (32.6)
	positive	27,044 (67.4)
	unknown	15,926
HER2-status		
	negative	23,787 (82.7)
	positive	4,969 (17.3)
	unknown	27,312
Surgery		
	yes, breast saving	16,468 (42.3)
	yes, mastectomy	11,315 (29.1)
	yes, type unknown	11,163 (28.7)
	unknown	17,122
(Neo)adjuvant chemotherapy		
	no	18,110 (49.4)
	yes	18,559 (50.6)
	unknown	19,399
(Neo)adjuvant endocrine therapy		
	no	10,781 (28.3)
	yes	27,322 (71.7)
	unknown	17,965
Radiotherapy		
	no	11,023 (27.4)
	yes	29,142 (72.6)
	unknown	15,903

Abbreviations: ER = estrogen receptor, PR = progesterone receptor, HER2 = human epidermal growth factor receptor 2

^a Total may not be 100% because of rounding

Table S5. Association between the 313-variant PRS (PRS₃₁₃) and contralateral breast cancer risk in the European cohort

Percentile categories of the PRS ₃₁₃	No. of women	No. of CBC	HR per unit SD ^a	95%CI	P-value
0 th to 10 th	5,607	65	0.59	0.45-0.78	<.001
10 th to 20 th	5,606	79	0.71	0.55-0.92	.01
20 th to 40 th	11,214	165	0.74	0.60-0.90	.003
40 th to 60 th	11,214	224	1.00	Ref.	-
60 th to 80 th	11,214	208	0.90	0.74-1.08	.25
80 th to 90 th	5,607	121	1.05	0.84-1.31	.69
90 th to 100 th	5,606	165	1.38	1.13-1.69	.002

Abbreviations: PRS = polygenic risk score, No = number, CBC = contralateral breast cancer, HR = hazard ratio, CI = confidence interval, SD = standard deviation

^aThe analysis was performed with attained age as time scale. Coefficients to construct the PRS₃₁₃ are shown in Table S3. The PRS₃₁₃ was standardized by SD=0.61, in line with Mavaddat et al.¹

Table S6. Multivariable Cox regression models of contralateral breast cancer risk by 313-variant PRS (PRS₃₁₃) in all women, all women excluding studies oversampling cases with family history, and those with complete covariate information

	All patients			All women excluding studies oversampling cases with family history			Complete case			
	N=56,068 (CBC=1,027)			N=51,883 (CBC=829)			N=12,065 (CBC=193)			
	HR per unit SD ^a	95%CI	P-value	HR per unit SD ^a	95%CI	P-value	HR per unit SD ^a	95%CI	P-value	
<i>Model 1</i> PRS ₃₁₃ ^b	1.25	1.18-1.33	<.001	1.26	1.17-1.34	<.001	1.35	1.17-1.56	<.001	
<i>Model 2</i> PRS ₃₁₃ ^b	1.23	1.16-1.31	<.001	1.25	1.17-1.34	<.001	1.33	1.15-1.54	<.001	
Family history	yes vs. no	1.43	1.24-1.64	<.001	1.34	1.13-1.59	.001	1.49	1.06-2.09	.02
	unknown vs. no	0.93	0.75-0.16	.54	0.92	0.73-1.16	.47	-	-	-
<i>Model 3</i> PRS ₃₁₃ ^b	1.25	1.18-1.33	<.001	1.26	1.17-1.34	<.001	1.35	1.17-1.56	<.001	
Nodal status	positive vs. negative	1.05	0.91-1.20	.50	1.07	0.92-1.25	.37	1.14	0.85-1.53	.37
	unknown vs. no	1.26	1.04-1.53	.02	1.29	1.04-1.60	.02	-	-	-
<i>Model 4</i> PRS ₃₁₃ ^b	1.25	1.18-1.33	<.001	1.26	1.18-1.35	<.001	1.35	1.17-1.56	<.001	
Tumor size,	(2-5] vs. ≤2	1.08	0.92-1.25	.34	1.12	0.95-1.32	.20	0.93	0.68-1.27	.66
	>5 vs. ≤2	1.37	0.99-1.89	.06	1.45	1.02-2.07	.04	1.63	0.93-2.85	.09
	unknown vs. ≤2	1.23	1.04-1.47	.02	1.14	0.94-1.39	.18	-	-	-
<i>Model 5</i> PRS ₃₁₃ ^b	1.25	1.17-1.33	<.001	1.25	1.17-1.34	<.001	1.35	1.17-1.57	<.001	
Differentiation grade	II vs. I	0.93	0.76-1.13	.45	0.99	0.80-1.24	.94	0.98	0.65-1.48	.93
	III vs. I	0.90	0.73-1.12	.35	0.97	0.76-1.24	.81	1.09	0.70-1.69	.69
	unknown vs. I	1.20	0.96-1.49	.11	1.45	1.13-1.86	.004	-	-	-
<i>Model 6</i> PRS ₃₁₃ ^b	1.25	1.17-1.33	<.001	1.25	1.17-1.34	<.001	1.33	1.16-1.54	<.001	
Morphology	lobular vs. ductal	1.26	1.03-1.53	.03	1.34	1.08-1.67	.008	1.48	0.99-2.21	.05
	mixed (ductal and lobular) vs. ductal	1.28	0.94-1.73	.11	1.36	0.98-1.88	.06	1.48	0.87-2.54	.15
	other vs. ductal	1.04	0.81-1.33	.75	0.91	0.66-1.24	.55	1.24	0.69-2.21	.47
	unknown vs. ductal	1.77	1.42-2.19	<.001	1.82	1.44-2.30	<.001	-	-	-
<i>Model 7</i> PRS ₃₁₃ ^b	1.25	1.18-1.33	<.001	1.26	1.18-1.35	<.001	1.35	1.17-1.56	<.001	
ER-status	positive vs. negative	0.88	0.75-1.04	.14	0.86	0.72-1.03	.11	0.90	0.62-1.32	.60
	unknown vs. negative	1.16	0.93-0.43	.19	1.11	0.86-1.43	.43	-	-	-
<i>Model 7</i> PRS ₃₁₃ ^b	1.25	1.18-1.33	<.001	1.26	1.18-1.35	<.001	1.35	1.17-1.56	<.001	

PR-status	positive vs. negative	0.95	0.81-1.11	.51	0.92	0.78-1.09	.32	0.91	0.66-1.25	.56
	unknown vs. negative	1.15	0.95-1.40	.14	1.10	0.88-1.37	.40	-	-	-
<i>Model 9</i>										
PRS₃₁₃^b		1.25	1.18-1.33	<.001	1.26	1.17-1.34	<.001	1.34	1.16-1.55	<.001
HER2-status	positive vs. negative	0.84	0.64-1.11	.22	0.76	0.56-1.05	.10	0.70	0.45-1.10	.12
	unknown vs. negative	1.29	1.11-1.50	.001	1.28	1.08-1.52	.004	-	-	-
<i>Model 10</i>										
PRS₃₁₃^b		1.25	1.17-1.32	<.001	1.26	1.17-1.34	<.001	1.35	1.16-1.56	<.001
Chemotherapy	yes vs. no	0.86	0.73-1.01	.06	0.99	0.83-1.19	.92	0.89	0.64-1.25	.51
	unknown vs. no	1.09	0.91-1.31	.34	1.20	0.97-1.47	.09	-	-	-
<i>Model 11</i>										
PRS₃₁₃^b		1.25	1.18-1.33	<.001	1.26	1.18-1.35	<.001	1.36	1.17-1.57	<.001
Endocrine therapy	yes vs. no	0.75	0.64-0.88	.001	0.92	0.75-1.12	.41	0.78	0.55-1.11	.17
	unknown vs. no	0.90	0.75-1.09	.28	1.11	0.87-1.41	.39	-	-	-
<i>Model 12</i>										
PRS₃₁₃^b		1.25	1.17-1.32	<.001	1.26	1.17-1.34	<.001	1.35	1.17-1.56	<.001
Radiotherapy	yes vs. no	1.00	0.85-1.18	1.00	0.98	0.82-1.18	.85	1.35	0.88-2.08	.17
	unknown vs. no	1.41	1.14-1.74	.001	1.18	0.93-1.50	.17	-	-	-
<i>Model 13</i>										
PRS₃₁₃^b		1.25	1.17-1.32	<.001	1.25	1.17-1.34	<.001	1.34	1.16-1.55	<.001
Year of first breast cancer diagnosis		0.95	0.94-0.96	<.001	0.95	0.93-0.96	<.001	0.90	0.86-0.95	<.001
<i>Model 14</i>										
PRS₃₁₃^b	full model ^c	1.23	1.16-1.31	<.001	1.25	1.16-1.33	<.001	1.33	1.15-1.53	<.001

Abbreviations: PRS = polygenic risk score, CBC = contralateral breast cancer, HR = hazard ratio, CI = confidence interval, SD = standard deviation, ER = estrogen receptor, PR = progesterone receptor, HER2 = human epidermal growth factor receptor 2

^a All analyses were performed with attained age as the time scale

^b Coefficients to construct the PRS₃₁₃ are shown in Table S3. The PRS₃₁₃ was standardized by SD=0.61, in line with Mavaddat et al.¹

^c Adjusted for family history, nodal status, tumor size, differentiation grade, morphology, ER status, HER2 status, chemotherapy, endocrine therapy, radiotherapy, and year of first breast cancer diagnosis

Table S7. Five- and ten-year cumulative risks of contralateral breast cancer by the 313-variant PRS (PRS₃₁₃) for different age groups

Age at first breast cancer diagnosis (years)	5-year cumulative CBC risks (%) range by age					10-year cumulative CBC risks (%) range by age				
	5 th percentile PRS ₃₁₃	10 th percentile PRS ₃₁₃	50 th percentile PRS ₃₁₃	90 th percentile PRS ₃₁₃	95 th percentile PRS ₃₁₃	5 th percentile PRS ₃₁₃	10 th percentile PRS ₃₁₃	50 th percentile PRS ₃₁₃	90 th percentile PRS ₃₁₃	95 th percentile PRS ₃₁₃
30-34	1.9-3.3	2.1-3.6	2.8-4.7	3.7-6.2	4.0-6.8	3.3-4.4	3.6-4.8	4.7-6.3	6.2-8.3	6.8-9.1
35-39	0.8-2.2	0.9-2.4	1.2-3.2	1.6-4.2	1.7-4.6	2.2-3.9	2.4-4.2	3.2-5.5	4.2-7.2	4.6-8.0
40-44	1.5-2.9	1.7-3.2	2.2-4.2	2.9-5.5	3.2-6.0	2.9-4.9	3.2-5.3	4.2-7.0	5.5-9.1	6.0-10.0
45-49	1.4-2.5	1.5-2.8	2.0-3.7	2.6-4.8	2.9-5.3	2.5-4.2	2.8-4.5	3.7-6.0	4.8-7.8	5.3-8.6
50-54	1.4-2.9	1.5-3.1	2.0-4.1	2.6-5.5	2.9-6.0	2.9-4.8	3.1-5.3	4.1-6.9	5.5-9.1	6.0-10.0
55-59	1.6-3.3	1.8-3.6	2.4-4.7	3.1-6.2	3.4-6.8	3.3-5.3	3.6-5.7	4.7-7.5	6.2-9.8	6.8-10.8
60-64	1.8-3.5	1.9-3.8	2.6-5.0	3.4-6.5	3.7-7.2	3.5-5.5	3.8-6.0	5.0-7.9	6.5-10.3	7.2-11.3
65-70	1.5-3.5	1.7-3.8	2.2-5.0	2.9-6.6	3.2-7.2	3.5-4.6	3.8-5.0	5.0-6.6	6.6-8.7	7.2-9.5

Abbreviations: PRS = polygenic risk score, CBC = contralateral breast cancer

Coefficients to construct the PRS₃₁₃ are shown in Table S3. The PRS₃₁₃ was standardized by SD=0.61, in line with Mavaddat et al¹. The CBC incidences for each age group were calculated based on incidence data from the Netherlands Cancer Registry² and relative risks estimated as described in the Material and Methods. In contrast to Table 4, death was not taken into account as competing risk

Table S8. Estimates of unilateral- and contralateral breast cancer risk by the 313-variant PRS (PRS₃₁₃) in the European case-case series and the Asian case-case series

	European						Asian		
	Case-case series ^a			Validation set ^b			Case-case series ^a		
PRS ₃₁₃ ^c	OR per unit SD	95%CI	P-value	OR per unit SD	95%CI	P-value	OR per unit SD	95%CI	P-value
Unilateral breast cancer versus control	1.82	1.80-1.84	<.001	1.67	1.59-1.76	<.001	1.56	1.52-1.60	<.001
CBC versus unilateral breast cancer	1.30	1.26-1.35	<.001	1.39	1.13-1.70	.002	1.15	1.02-1.29	.02

Abbreviations: PRS = polygenic risk score, CBC = contralateral breast cancer, OR = odds ratio, SD = standard deviation, CI = confidence interval

^a Adjusted for country and age. For all women with unilateral- and contralateral breast cancer we used age at first breast cancer diagnosis, and for control women without any diagnosis of breast cancer we used age at baseline questionnaire.

^b The validation set was previously used to develop the PRS₃₁₃; see details in materials and methods. For analyses in the current paper, this set is nested within the case-case series. These analyses were additionally adjusted for 10 principal components for comparability with the originally published PRS₃₁₃ overall estimates¹

^c Coefficients to construct the PRS₃₁₃ are shown in Table S3. The PRS₃₁₃ was standardized by SD=0.61, in line with Mavaddat et al.¹

Supplemental Note

Our initial aim was to externally validate our results using the UK Biobank, which seemed the most suitable cohort given the large number of women diagnosed with breast cancer with information available on the PRS₃₁₃. However, when we started the analyses, it turned out that the UK Biobank had no information available on the laterality of the second breast tumor. Therefore, we were unable to distinguish between ipsilateral and contralateral breast cancer, and had to define our endpoint in these analyses as ‘any second breast cancer’. In addition, in comparison to our analyses in the BCAC, we were unable to exclude patients diagnosed with stage IV invasive first breast cancer from the UK Biobank cohort, and had limited information on metastases developed during follow-up.

The association between the overall breast cancer PRS₃₁₃ and (any) second breast cancer was evaluated among women aged ≥ 18 years of European ancestry from the UK Biobank cohort who had had a diagnosis of invasive first breast cancer. UK Biobank samples were genotyped using Affymetrix UK BiLEVE Axiom array and Affymetrix UK Biobank Axiom® array and imputed to the combined 1000 Genome Project v3 and UK10K reference panels using SHAPEIT3 and IMPUTE3⁴. The lowest imputation info score for the variants used in these analyses was 0.86. Samples were included for this analysis of the UK BIOBANK study on the basis of female sex (genetic and self-reported) and ethnicity filter (Europeans/White British ancestry subset). Duplicates and individuals with high degree of relatedness (samples which have >10 putative third degree relatives) were removed, and we randomly excluded one of each related pair first-degree relatives. Samples were also excluded on standard quality control criteria. The PRS₃₁₃ was calculated as a weighted sum of the minor allele dosages; the variant selection and weights are as given by Mavaddat et al¹. The PRS₃₁₃ was standardized by SD=0.61, in line with our BCAC analyses and Mavaddat et al¹.

The final cohort included 10,567 women with invasive breast cancer among whom 302 registry-confirmed second breast cancers developed over 59,260 person-years of follow-up. A Cox proportional hazards model was used to assess the association between PRS_{313} and second breast cancer risk. Time at risk started three months after the age of first breast cancer diagnosis, where this was diagnosed after the baseline questionnaire date, or three months after the baseline questionnaire where first breast cancer was diagnosed before the baseline questionnaire date. Time at risk ended at the age of second breast cancer diagnosis (ipsilateral or contralateral), distant metastasis (where available), death or end of follow-up (at latest December 10, 2016). Potential effect modification of the PRS_{313} by age was evaluated by adding an interaction term ($PRS_{313} \times \text{age at first breast cancer diagnosis [continuous]}$) in the model. We performed a separate analysis for invasive second breast cancer (241 breast cancers), where we censored on in situ second breast cancer.

The HR for a second breast cancer (in situ or invasive) per SD of PRS_{313} in the UK Biobank cohort was 1.13 (95%CI=1.01-1.26). We found no indication for interaction with age at first breast cancer diagnosis ($HR_{\text{interaction}}=1.00$, 95%CI=0.99-1.01; $P=0.87$). When analyses were restricted to invasive second breast cancer, the HR per SD was 1.13 (95%CI=1.00-1.29).

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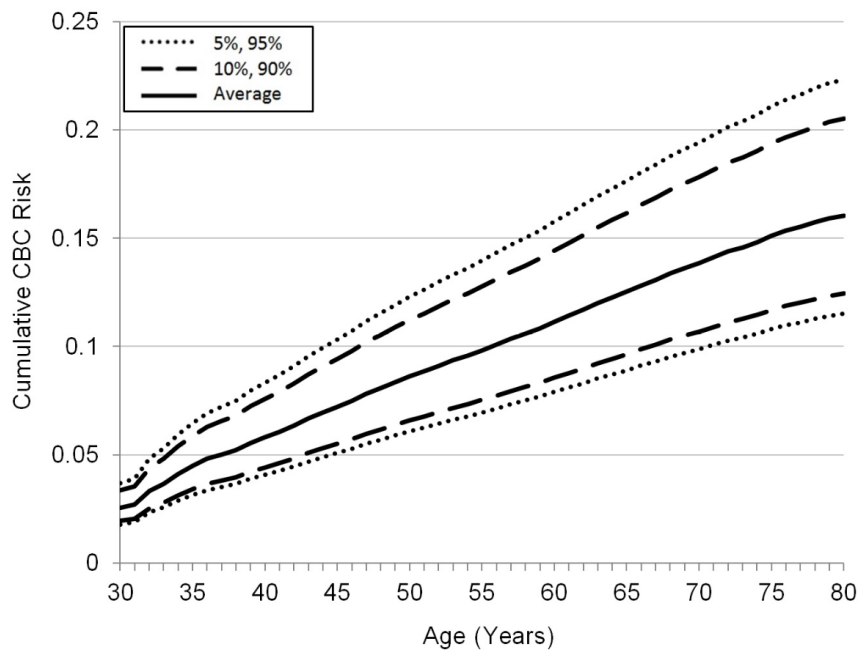
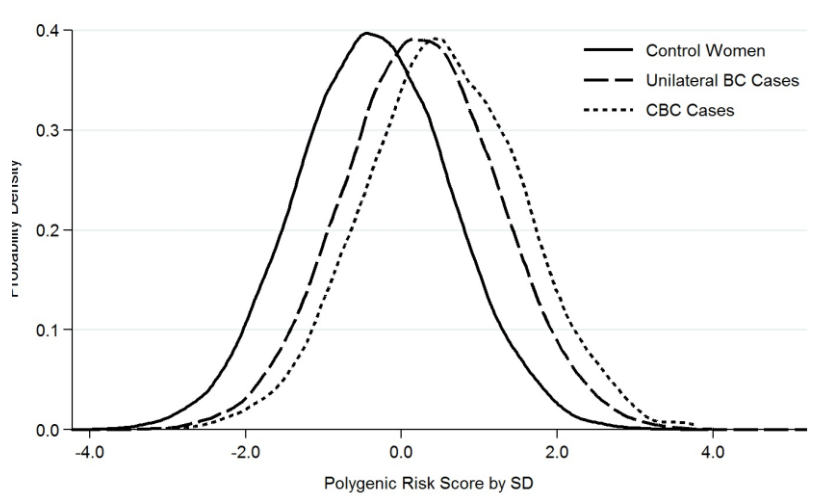
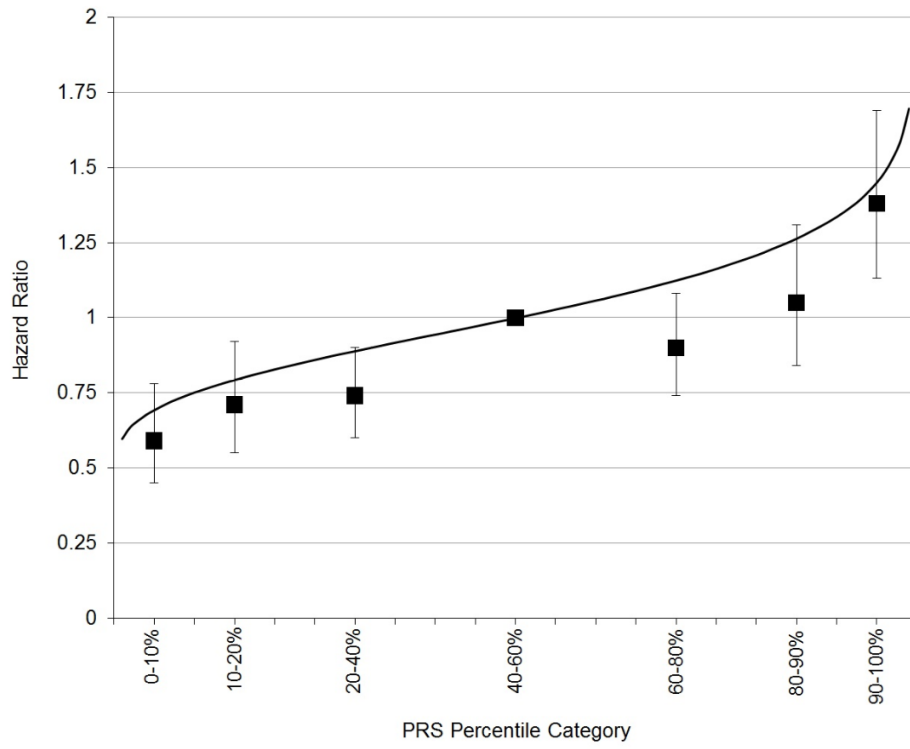
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Study	Abbreviation	Country	Studydesign	Case definition	Control definition	How was follow-up (including vital status) information obtained?	When was the most recent attempt to have complete follow-up?	References
Australian Breast Cancer Family Study	ABCFS	Australia	Population-based case-control study	All cases diagnosed < age 40 plus a random sample of those diagnosed ages 40-59 from cancer registries in Victoria and New South Wales, plus a limited number diagnosed aged 60-69; cases living in Melbourne recruited from 1992-99 and in Sydney from 1993-98	Identified from the electoral rolls in Melbourne from 1992-98 and Sydney from 1993-98. Frequency matched to cases by age 5 year categories	Systematic follow-ups by mail and telephone	2014	Dias, G.S. et al. Familial risks, early-onset breast cancer, and BRCA1 and BRCA2 germline mutations. <i>J. Natl. Cancer Inst.</i> 95, 448-57 (2003)
Amsterdam Breast Cancer Study	ABCS	Netherlands	Hospital-based consecutive cases; population-based controls (for COGS/Oncovary from blood bank)	Pre-COGS: All breast cancer patients (with operable, invasive mammary carcinoma) aged <50 years and diagnosed from 1970-1994 in four Dutch hospitals COGS/Oncovary: Breast cancer patients diagnosed before age 50 in 1995, 2011 at the Netherlands Cancer Institute - Antoni van Leeuwenhoek hospital (NKI-AVL)	No controls [Use controls of ABCS]	Hospital medical registry and linkage with municipality registry	January 2014	1) M. K. Schmidt, et al. Breast Cancer Survival and Tumor Characteristics in Premenopausal Women Carrying the CHEK2*1100delC Germline Mutation. <i>J. Clin. Oncol.</i> 25, 64-9 (2007) 2) K. Michalidis, et al. Large-scale genotyping identifies 41 new loci associated with breast cancer risk. <i>Nat. Genet.</i> 45, 353-61 (2013)
Amsterdam Breast Cancer Study - Familial	ABCS-F	Netherlands	Clinical Genetic Center-based cases	Pre-COGS: Only in BRCA Phase III familial non-BRCA1/2 cases <50 from the Clinical Genetic Centre of the Netherlands Cancer Institute were included COGS/Oncovary: All non-BRCA1/2 breast cancer cases from the family cancer clinic of the NKI-AVL, tested in the period 1995-2009; all ages and diagnosed with breast cancer in 1995-2012	No controls [Use controls of ABCS]	Hospital medical registry and linkage with municipality registry	April 2014	M.K. Schmidt, et al. Age- and Tumor Subtype-Specific Breast Cancer Risk Estimates for CHEK2*1100delC Carriers. <i>J. Clin. Oncol.</i> Jun 6; pii: JCO68844 (2016)
Australian Breast Cancer Tissue Bank	ABCTB	Australia	Hospital-based multi site newly diagnosed breast cancer case	Newly diagnosed unselected cases from 32 hospitals in New South Wales from 2006	Female controls from the Hunter Community Study (HCS) invited by a newspaper advertisement in Northern NSW, and recruited during 1999-2013	Multiple (medical records at surgeon's rooms, medical records and databases at hospital clinics and GP Clinics)	Follow-up ongoing: Follow-up is attempted at 1 and 5 years after ABCTB, and 5 year intervals after the	McEwen, J. et al. Cohort profile: The Hunter Community Study. <i>Int. J. Epidemiol.</i> 2010 Dec;39(6):1452-63
Bavarian Breast Cancer Cases and Controls	BICC	Germany	Hospital-based cases; population based controls	Consecutive, unselected cases with invasive breast cancer recruited at the University Breast Centre, Franconia in Northern Bavaria during 1999-2013	Healthy women with no diagnosis of cancer aged 50 or older, invited by a newspaper advertisement in Northern Bavaria, and recruited during 1999-2013	Cancer registry and Medical records	12/1/2013	1) Fackenthal PA, et al. Single nucleotide polymorphisms of the aromatase-like gene (CYP19A1), HER2/neu status, and prognosis in breast cancer patients. <i>Breast Cancer Res Treat.</i> DOI: 10.1007/s10548-007-9822-2 (2007) 2) Schrader M, et al. Single nucleotide polymorphism 185SN of the ATM gene may alter the risk for breast cancer. <i>J. Cancer Res. Clin. Oncol.</i> 134, 873-82 (2008)
British Breast Cancer Study	BBCS	UK	Cancer registry and National Cancer Research network (NCRN) based cases; population-based controls	1) English & Scottish Cancer Registries: all breast cancer cases who developed a first primary before age 65 in 1971 or later and who subsequently developed a second primary cancer. 2) Unilateral breast cancer cases diagnosed before age 70 in 1971 or later	1) A friend, sister-in-law, daughter-in-law or other non-blood relative of cases. Recruitment of cases and controls began in January 2001	Cases are not followed up	We have not attempted to follow up	1) Johnson, N. et al. Interaction between CHEK2*1100delC and other low-penetrance breast-cancer susceptibility genes: a familial study. <i>Lancet Oncol.</i> 2006, 7(7):507-13. 2) Fletcher, O. et al. Inconsistent association between the CTR1-F11 genetic polymorphism and breast cancer risk. <i>J. Natl. Cancer Inst.</i> 98, 1014-8 (2006)
Breast Cancer Employment and Environment Study	BCEES	Australia	Population-based case-control study	First incident invasive breast cancer diagnosed between May 2009 and January 2011, residing in Western Australia and reported to the state wide mandatory Cancer Registry	Randomly selected from Western Australia electoral roll (registration is compulsory for Australian citizens)	No follow-up has been completed	No follow-up has been completed	Fitzhugh L, Erren TC, Glass DC, Girschik J, Thomson AK, Saunders C, et al. The association between different high shiftwork factors and breast cancer: a case-control study. <i>Br J Cancer.</i> 2015;109:2472-80
New York Breast Cancer Family Registry	BCFRNY	USA	Clinic-based recruitment of families; family-based cohort	Recruitment took place from Jan 1996 to Dec 2012. Eligibility was based on one or more of the following criteria: 2) or more relatives with a personal history of breast or ovarian cancer; a woman diagnosed with breast or ovarian cancer at a young age; a woman with a history of both breast and ovarian cancer; an affected male, or known BRCA1 or BRCA2 mutation carriers	Unaffected family members also enrolled Jan 1996 to Dec 2012	Systematic follow-up every five years for questionnaire data and annual update of cancer history and vital records by at least one family member	Ongoing	1) John, EM et al. The Breast Cancer Family Registry: an infrastructure for cooperative multinational, interdisciplinary and translational studies of the genetic epidemiology of breast cancer. <i>Breast Cancer Res.</i> 6, R275-R288 (2004) 2) Wu, H. et al. Repetitive element DNA methylation levels in white blood cell DNA from sisters discordant for breast cancer from the New York site of the Breast Cancer Family Registry. <i>Carcinogenesis.</i> 2012 Oct;33(10):1948-52 3) Quanté et al. Practical problems with clinical guidelines for breast cancer prevention based on remaining lifetime risk. <i>J. Natl. Cancer Inst.</i> 2015 May 8;107(17)
Philadelphia Breast Cancer Family Registry	BCFR-PA	USA	Clinic-based recruitment of families; family-based cohort	Recruitment took place from 1996 to 2011. Eligibility was based on one or more relatives with a personal history of breast or ovarian cancer; a woman diagnosed with breast or ovarian cancer at a young age; a woman with a history of both breast and ovarian cancer; an affected male, or known BRCA1 or BRCA2 mutation carriers	Unaffected family members also enrolled 1996 to 2011	Self-report on questionnaires or clinical database	2015 questionnaire	1) John, EM et al. The Breast Cancer Family Registry: an infrastructure for cooperative multinational, interdisciplinary and translational studies of the genetic epidemiology of breast cancer. <i>Breast Cancer Res.</i> 6, R275-R288 (2004) 2) Terry M, Phillips K, Daly M, et al. Cohort Profile: The Breast Cancer Prospective Family Study Cohort (ProF-SC). <i>International Journal of Epidemiology</i> 2015;44:118
Utah Breast Cancer Family Registry	BCFR-UT	USA	Clinic-based recruitment of non-BRCA1/2 familial breast cancer cases; unaffected BRCA1/2 carriers as controls	Index cases from families tested negative for BRCA1/2 mutations. Recruited in Utah during 1995-2008	Unaffected BRCA1/2 carriers. Recruited in Utah during 1995-2008	Follow-up questionnaire	15 years after enrollment	1) John, EM et al. The Breast Cancer Family Registry: an infrastructure for cooperative multinational, interdisciplinary and translational studies of the genetic epidemiology of breast cancer. <i>Breast Cancer Res.</i> 6, R275-R288 (2004) 2) Terry M, Phillips K, Daly M, et al. Cohort Profile: The Breast Cancer Prospective Family Study Cohort (ProF-SC). <i>International Journal of Epidemiology</i> 2015;44:118
Breast Cancer in Northern Israel Study	BOINIS	Israel	Population-based case-control study	All consecutive cases of invasive BC and DCIS diagnosed in a geographically defined area of Northern Israel since 1990. On-going	Age-sex-residence-ethnicity (Jews/Arabs)-matched controls, randomly sampled from the population using population registries. Recruitment since 1990 and on-going. A matched control will be interviewed within 1 month of the interview of its matched case.	Oncology files, medical records, population database	June 2016	1) Rennett G, Pinchev M, Rennett HS. Use of Postmenopausal Breast Cancer. <i>J. Clin. Oncol.</i> 2014; 32(26):2823-2829. 2) Rennett G, Leibkowitz F, Cohen I, Pinchev M, Rennett HS, Ramot-Grinberg O. Multi-trait mutation analysis increased breast cancer risk. <i>Cancer.</i> 2011 Sep 22; 117(10):2929-2936. [Epub ahead of print]
Breast Cancer in Galway Genetic Study	BIGGS	Ireland	Hospital-based cases; population based controls	Unselected cases recruited from West of Ireland since 2001. Cases were recruited from University College Hospital Galway and surrounding hospitals	Women > 60 years with no personal history of any cancer and no family history of breast or ovarian cancer were identified from internet groups in the West of Ireland (same catchment area as cases) during the period 2001-2008	Local clinical database	01/08/2013	1) Collier G, McInerney N, Rowan A, Bartsley E, Jones AM, Curran C, Miller N, Keim M, Tomlinson I, Sawyer E. The GGBR1>A>A polymorphism is not associated with differential risk of breast cancer. <i>Breast Cancer Res Treat.</i> 2009 Apr 24; 114(1-2):169-74. 2) Niall McInerney, Gabrielle Collier, Andrew Rowan, Aislinn Bartsley, Eilish Breen, Sarah Annette M. Jones, Stephen Tuohy, Catherine Curran, Nicola Miller, Michael Keim, Ian Tomlinson, Eileen J. Sawyer. Low penetrance SNPs are site specific. <i>Breast Cancer Research Treatment</i> 2008 Nov 13 online
Breast Cancer Galicia Network	BREGAN	Spain	Population-based case-control	A population-based study conducted since 1997 in two cities in Galicia, Spain (Vigo and Santiago) covering approximately 700,000 inhabitants. The study currently includes over 1000 incident breast cancer cases diagnosed from 1997-2014 in two Galician hospitals with blood, tumor tissue and risk factor questionnaire	Controls were frequency-matched to cases according to 5-year age group, inclusion in the universal Galician Public Health Service (SERGAS) registry database, and place of residence. They were healthy, unrelated female individuals from the same base population as cases randomly selected from SERGAS primary healthcare centers in the health areas of Santiago and Vigo. Recruitment began in 1997	From the computerized electronic individual medical history program unique for each individual in the state. Each individual is identified by his/her social security number which is unique to each person during his/her lifetime, his/her first name and his/her last names, his/her national ID number which is unique to each individual during his/her lifetime, date of birth, etc. All information is 100% computerized under the larus program. This program includes information from all services and departments in all state hospitals, continuing update from all hospitals and family clinics, as well as electronic prescriptions and medical imaging	Every six months	1) Castaño JE, Jiang XJ, Chavez-Urbe E, Fernandez Rodriguez B, Celeste Muñoz C, Redondo Marey C, Peña Fernandez M, Nova Domercq A, Dorso Perez A, Dorso Pereira C, Torre MA, Garcia-Caballero T, Martinez ME, Rago Rodriguez M, Carrascosa A, Fontana J, Gago-Dominguez M. Family History and Breast Cancer Subtypes in a Spanish Cohort. <i>PLoS One.</i> 2012; 7(2):e34592. 2) Redondo, C. M., M. Gago-Dominguez et al. (2015). Breast feeding, parity and alcohol consumption in a Spanish cohort. <i>PLoS One</i> 10(10):e0140163. 3) Redondo, C. M., M. Gago-Dominguez et al. (2015). Epidemiological Biomarkers Prev 2014; 17(4): 424-431. 4) M. E. Mattarini, et al. (2012). Hypothetical role of pregnancy hormones on HER2+ breast tumor development. <i>Breast Cancer Res Treat.</i> 0. 5) Gago-Dominguez et al. Biomarkers 2016; 18(3): 204-210. 1186940064-015-1630-2 RESEARCH Alcohol and breast cancer subtypes in a Spanish Cohort
Breast Cancer Study of the University of Heidelberg	BSUJH	Germany	Hospital-based cases; healthy blood donor controls	Cases diagnosed with breast cancer/breast cancer metastases in 2008-2011 at the University Women's Clinic Heidelberg	Healthy, unrelated, ethnically matched female blood donors recruited in 2007, 2009 & 2012 by German Red Cross Blood Service of Baden-Württemberg-Hessen, Institute of Transfusion Medicine & Immunology, Mannheim	Individual clinical investigation of patients, medical records	January 2017	Yang, R. et al. Genetic variants within miR-126 and miR-335 are not associated with breast cancer risk. <i>Breast Cancer Res Treat.</i> 127, 549-554 (2011)
Canadian Breast Cancer Study	CBCS	Canada	Population-based case-control study	British Columbia (BC): Incident cases diagnosed 2005-2009, resident in Vancouver area ascertained from the population cancer registry, Ontario - recruited from the Hot Spot Breast Assessment Program in Kingston, Ontario, 2005-2009	BC - cancer-free women who consented to participate in research studies through routine screening mammography Available to women in BC aged 40-79 years through the Screening Mammography Program of BC in 2007-2008, resident in the same geographic areas and frequency-matched to cases by 5-year age group, Ontario - women with either normal mammogram results or a diagnosis of benign breast disease, frequency matched by 5-year age group recruited from the Hot Spot Breast Assessment Program in Kingston, Ontario, 2005-2009	Annual GP letter follow-up for patients discharged from BCCA (only for those who attended a BCCA clinic) - 65% of incident breast cancer cases; vital status info primarily obtained through the British Columbia Vital Statistics Agency	Ongoing	1) Grundy A, Schetz JM, Lai AS, Janoo-Gilani R, Leach S, Burstin H, Richardson H, Brooks-Wilson A, Spinelli JJ, Kronen KJ, Shiit wok, circadian gene variants and risk of breast cancer. <i>Cancer Epidemiol.</i> 2013 Oct;37(10):606-12. doi: 10.1016/j.cane.2013.04.008. Epub 2013 May 28. PMID: 23725643 2) Kobayashi LC, Janssen I, Richardson H, Lai AS, Spinelli JJ, Aronson KL. Moderate-to-intensive intensity physical activity across the life course and risk of pre- and post-menopausal breast cancer. <i>Breast Cancer Res Treat.</i> 2013 Jun;138(3):1851-60. doi: 10.1007/s10548-013-2596-8. Epub 2013 Jun 15. PMID: 23717136, 32161. 3) Grundy A, Richardson H, Burstin J, Lohrich G, Sengupta SK, Lai AS, Lee D, Spinelli JJ, Aronson KL. Increased risk of breast cancer associated with long-term shift work in Canada. <i>Occup Environ Med.</i> [with commentary] 2013 Dec;70(12):831-8. doi: 10.1136/occup-2013-101482. Epub 2013 Jul 1. PMID 23878141, 4) Ziv, Kobayashi LC, Janssen I, Richardson H, Lai AS, Spinelli JJ, Aronson KL. A case-control study of lifetime light intensity physical activity and breast cancer risk. <i>Cancer Causes Control</i> 2014 Jan 25; 25:133-140. DOI: 10.1007/s10552-013-0312-z. Epub 2013 Oct 25. PMID: 24158779
Crete Cancer Genetics Program	CCGP	Greece	Hospital-based case-control study	Incident breast cancer cases treated between 2004 and 2013 at the University Hospital of Heraklion on Crete; all enrolled within 6 months of diagnosis	Healthy, unrelated, ethnically matched female blood donors recruited in 2014 by the laboratory of Hemostasis at the General Hospital of Heraklion "Venizelos"	Individual patient medical records	August 2014 for all patients included in the database	No references yet
Copenhagen General Population Study	CGPS	Denmark	Population-based case-control study	Consecutive, incident cases from 1 hospital with centralized care for a population of 400,000 women from 2001 to the present	Community controls residing in the same region as cases and with no history of breast cancer were identified from the Copenhagen General Population Study recruited 2003-2007. All controls were known to still be breast cancer-free at the end of 2007	Vital status: from the citizen registry	01/12/2017	Weischer M, Bøgesen S.E., Tybjaerg-Hansen A, Nordestgaard B.G. Increased risk of breast cancer associated with CHEK2*1100delC. <i>J. Clin. Oncol.</i> 25, 97-83 (2007)
Spanish National Cancer Centre Breast Cancer Study	ONCO-BCS	Spain	Case-control study	Two groups of cases: 1) 574 consecutive breast cancer patients, unselected for family history, from 3 public hospitals, 2 in Madrid and one in Oviedo, from 2000 to 2005; 2) 291 cases with at least one first degree relative also affected with breast cancer, recruited through the ONCO family cancer clinic in Madrid from 2000 to 2004	Women attending the Menopause Research Centre between 2000 and 2004 and female members of the College of Lawyers attending a free, targeted medical check-up in 2005, all free of breast cancer and all in Madrid	Only available for a subset of cases (using clinical records?)	2007/2008	Mirne, RL et al. EROCA associated with breast cancer risk: a two-stage case-control study using high throughput genotyping. <i>Cancer Res.</i> 66, 9425-7 (2006)
California Teachers Study	CTS*	USA	Prospective cohort study; nested case-control	This is a nested case-control study conducted within a cohort of California teachers (113,590) who were under age 80 years at baseline, had no prior history of invasive or in situ breast cancer. Cases are women newly diagnosed with a histologically confirmed invasive primary adenocarcinoma of the breast at age 80 years or younger from 1998 to 2008	Controls are a probability sample of at-risk cohort members, frequency matched to cases on age at baseline (5-year age groups), self-reported race/ethnicity (white, African American, Latina, Asian, other), and broad geographic region within California. Controls were recruited during 1968 to 2008 and selected without replacement, using an assigned reference date	The vital status and follow-up date are standard items of the California Cancer Registry	Completed information is as of 10/30/2014	Bernstein L, Allen M, Anton-Culver H, Deegan D, Horn-Ross PL, Peil D, Preder R, Reynolds P, Sullivan-Halley J, West D, et al. High breast cancer incidence rates among California teachers: results from the California Teachers Study (United States). <i>Cancer Causes Control</i> 2002, 13(7):625-635
DiCoMPLy Breast Cancer Survival Study	DIETCOMPLY F	UK	Multi-centre prospective cohort study	Invasive primary breast cancer grade I-III, patients recruited < 10 months after diagnosis, < age 75, Recruitment throughout UK. Patient first recruited on 18/07/97. Study joined NCRN in July 2004. Recruitment finished on 31/10/10	No controls	From the patients at the last time they visited the hospital	26th March 2014	Brewer R, Pentre M, Kallakuri LS, Gira C, Dutton SJ, Milligan AA, Woodside JV, Cartmill MM, Leatham AJ, Robertson CE, Dwek MW. The DiCoMPLy study: A prospective cohort study of breast cancer survival and phytoestrogen consumption. <i>Maturitas</i> (2013) 75, 232-240
Family History Risk Study	FRISIK	UK	Clinic-based cohort study with a nested case-control study	Women diagnosed with breast cancer and attending the Family History Clinic in Manchester for increased risk of breast cancer. Recruitment period 2009-2012	Women attending the same Family History Clinic as the cases but without a breast cancer diagnosis. Recruitment period is the same as for the cases	Follow up continues until the participant has been discharged from the Family History Clinic	Ongoing	1) Evans DG, Astley S, Stavrinou P, Harkeess E, Donnelly LS, Dawe S, Jacob F, Harvey M, Cuzick J, Brentnall A, Wilson M, Harrison F, Payne K, Howell A. Improvement in risk prediction, early detection and prevention of breast cancer in the NHS Breast Screening Programme and family history clinics: a dual cohort study. <i>Southern (UK): NIHR Journals Library.</i> 2016 Aug. 2) Ingalls SL, Warwick J, Butcher I, Sains S, O'Hara C, Moore A, Horgan S, Evans DG. Ovarian cancer among 8005 women from a breast cancer family history clinic: no increased risk of invasive ovarian cancer in families testing negative for BRCA1 and BRCA2. <i>J. Med Genet</i> 2013;50(6):368-72

Study Name	Country	Study Design	Study Description	Recruitment	Follow-up	Year	Notes
German Consortium for Hereditary Breast and Ovarian Cancer	Germany	Clinic-based case study and prospective cohort study	Women diagnosed with breast cancer in one of the GC-HBOC centres (Cologne, Munich, Kiel, Heidelberg, Düsseldorf, Ulm, Würzburg, Münster and Hannover). Recruitment period: 1996-present	Healthy, unrelated, ethnically and age-matched female control individuals (LIFE study, Leipzig, Germany)	Medical records or personal visit for women under intensified surveillance	Updated at least annually for women under intensified surveillance	1) Kato K, et al. Prevalence of BRCA1/2 germline mutations in 21,401 families with breast and ovarian cancer. <i>J Med Genet</i> 2016; 53(7):465-71. 2) Rahn K, et al. <i>Breast Cancer Res</i> 2012; 14(4):R106. 3) Graer MK, et al. <i>Chin Oncol</i> 2009; 18(12):1587-92. 4) Eneci C, et al. <i>BMC Cancer</i> 2016; 16:7181-1285
Gene Environment Interaction and Breast Cancer in Germany	Germany	Population-based case-control study	Incident breast cancer cases enrolled between 2000 and 2004 from the Greater Bonn area (by of the hospitals within the study region), all enrolled within 6 months of diagnosis	Selected from population registries from 31 communities in the greater Bonn area; matched to cases in 5-year age classes between 2001 and 2004	Through telephone interview with the patient or patient's relative, as well as information from the registration office and clinical records	Year 2012	1) Pesch, C, et al. Factors modifying the association between hormone replacement therapy and breast cancer risk. <i>Eur J Epidemiol</i> , 2006; 21(1):20-27. 2) Jostrombom C, et al. The CYP1B1_1508_G/G genotype is associated with estrogen receptor-negative breast cancer. <i>Breast Cancer Res</i> 2011; 13(1):177
Genetic Epidemiology Study of Breast Cancer by Age 50	Germany	Population-based study of women <50 years	All incident cases diagnosed <50 years of age in 1992-5 in two regions: Rhein-Neckar-Odenwald and Freiburg, by surveying the 38 clinics serving these regions	Selected from random lists of residents of the study regions supplied by population registries; two controls were selected for each case, matched by age and study region. Recruitment was carried out 1992-1998	Vital status was obtained by requesting the central tumour registry at the population registry	2009, but only for vital status and cause of death	Chang-Claude J, Eby N, Kueche M, Bastert G, & Becher H. Breastfeeding and breast cancer risk by age 50 among women in Germany. <i>Cancer Causes Control</i> 11; 687-695 (2000)
Study to Investigate the Genetics of Lobular Carcinoma in Europe	UK	Hospital-based case-control study	Cases aged 60 or younger with LCIS (pure or associated with invasive cancer of any subtype) or invasive lobular breast cancer from 96 hospitals throughout the UK. Recruitment period was from Jan 2007 to Sep 2012	Women with healthy women of any age with no history of LCIS, DCIS, breast disease or invasive breast cancer and who had no close relative (mother, sister, daughter or aunt) who had been so affected. Controls were recruited by asking non-blood relatives (generally sisters-in-law or friends) of affected individuals to act as a control. Recruitment period was from Jan 2007 to Jun 2013	Self reported by patient at time of recruitment	Not reported	Sawyer E et al. Genetic predisposition to in situ and invasive lobular carcinoma of the breast. <i>PLoS Genet</i> 2014 Apr 17;10(4)
Hannover Breast Cancer Study	Germany	Hospital-based case-control study	Cases who received radiotherapy for breast cancer at Hannover Medical School between 1996-2003 (HABC5 I), or were diagnosed with breast cancer at a certified Breast Cancer Clinics in the Hannover region between 2012-2016 (HABC5 II), unselected for age or family history	Anonymous female blood bank donors at Hannover Medical School, collected from 8/2005-12/2006, with known age and ethnic background	Follow-up information was obtained through the central tumour registry at MHH; rarely through telephone contact with clinicians	Summer 2016	Dork, T. et al. Spectrum of ATM gene mutations in a hospital-based series of unselected breast cancer patients. <i>Cancer Res</i> , 61, 7608-7615 (2001)
Hospital Clinico San Carlos	Spain	Population-based study of sporadic breast cancer cases	This is a cohort of a priori sporadic breast cancer patients which includes a cohort of 200 patients that were enrolled in a re-evaluation trial in which women were randomized to neoadjuvant docetaxel vs neoadjuvant epirubicin. Recruitment period was from 2000 to 2013. Most patients were treated in the Clinico San Carlos hospital (Madrid) and most are and being followed in different hospitals of Madrid	No controls	Medical records	February 2018	1) Romero A, Garcia-Saenz JA, Fuentes-Ferre M, Lopez Garcia-Arenjo JA, Furió V, Román JM, Moreno A, de la Hoya M, Diaz-Rubio E, Martín M, Castán L. Correlation between response to neoadjuvant chemotherapy and survival in locally advanced breast cancer patients. <i>Ann Oncol</i> 2013; 24(3):654-61. 2) Martín M, Romero A, Cheang MC, Lopez Garcia-Arenjo JA, Garcia-Saenz JA, Oliva B, Román JM, de la Hoya M, Castán L, Furió V, Puente J, Castán L, Vidari JA, Lopez-Tamulela S, Diaz-Rubio E, Pesus CM. Genomic predictors of response to docetaxel versus docetaxel in primary breast cancer. <i>Breast Cancer Res Treat</i> 2011 Jul;126(1):127-33
Heelsinki Breast Cancer Study	Finland	Hospital-based case-control study, plus additional familial cases	1) Consecutive cases (883) from the Department of Oncology, Helsinki University Central Hospital 1997-8 and 2000. 2) Consecutive cases (986) from the Department of Surgery, Helsinki University Central Hospital 2001 - 2004. 3) Familial breast cancer patients (536) from the Helsinki University Central Hospital, Departments of Oncology and Clinical Genetics (1995-)	Healthy females from the same geographical region in Southern Finland in 2003	Hospital medical records, Cancer registry, population registry	2011 breast cancer, spec 2015 overall	1) Syrjävälik, K. et al. Population-based study of BRCA1 and BRCA2 mutations in 1035 unselected Finnish breast cancer patients. <i>J Natl Cancer Inst</i> , 92, 1529-31 (2000). 2) Olgovaara, O. et al. Correlation of CHEK2 protein expression and c.1100delC mutation status with tumor characteristics among unselected breast cancer patients. <i>Int J Cancer</i> , 113, 575-80 (2005). 3) Faghehri, R. et al. MADD/Hsp90ome oxidoreductase 1 NOD1*2 genotype (P163) is a strong prognostic and predictive factor in breast cancer. <i>Nat Genet</i> 40, 844-853 (2008)
Hospital-based Epidemiologic Research Program at Aichi Cancer Center	Japan	Hospital-based case-control study	Incident breast cancer cases who first visited Aichi Cancer Center between 2001 and 2013 and were diagnosed within 1 year from the first visit. No previous history of any type of cancer	Controls were selected from pool of non-cancer patients who first visited Aichi Cancer Center between 2001-2011. Non-cancer status is defined as "having no positive finding on any of clinical/laboratory/genographic examination within 1 year from their first visit. No previous history of cancer is allowed"	Checking of medical records	End of 2014	Kawase T et al. FGFR2 intronic polymorphisms interact with reproductive risk factors of breast cancer: results of a case control study in Japan. <i>Int J Cancer</i> 2009; 125:1946-1952
Hong Kong Breast Cancer Study	Hong Kong	Hospital-based case-control study	Genetic screening of high risk breast cancer patients from all Hong Kong hospitals. Incident cases classified as high risk group: 1) first degree relative with breast and/or ovarian cancer, 2) cases where age is less than or equal to 45 years, 3) bilateral breast cancer, 4) high negative breast cancer cases, 5) family history of breast and/or ovarian cancer. Cases were recruited 2006-2014.	Controls were selected from pool of non-cancer patients who visited Hong Kong hospitals. Same period of recruitment as cases	Not reported	Not reported	1) Kwong A et al. Novel BRCA1 and BRCA2 genomic rearrangements in Southern Chinese breast/ovarian cancer patients. <i>Breast Cancer Res Treat</i> 2012; 136(3):593-3. 2) Kwong A et al. Identification of BRCA1/2 founder mutations in Southern Chinese breast cancer patients using family sequencing and high resolution DNA melting analysis. <i>PLoS One</i> 2012; 7(9):e43964
Hannover-Minsk Breast Cancer Study	Belarus	Hospital-based cases; population based controls	Ascertainment at the Byelorussian Institute for Oncology and Medical Radiology Aleksandrov N.N. in Minsk or at one of 6 regional oncology centers in Gomel, Mogilev, Grodno, Brest or Vitebsk through the years 2002-2008	Controls from the same population aged 18-72 years. Healthy (without personally history of cancer) female probands recruited from the same geographical region as cases during the years 2002-2008. About 75% of controls were women invited for general medical examination at the regional gynecology clinics in Gomel, Mogilev, Grodno, Brest or Vitebsk) and cancer-free volunteers ascertained at the Institute for Inherited Diseases in Minsk. 20% were cancer-free female blood bank donors recruited at Republican Blood Bank, Minsk, Belarus. Finally 5% of controls were healthy cancer-free relatives of some breast cancer patients	No data provided	No data provided	Bogdanov, N. et al. A nonsense mutation (E1978X) in the ATM gene is associated with breast cancer. <i>Breast Cancer Res</i> 11(8):207-211 (2009)
Hannover-Ufa Breast Cancer Study	Russia	Hospital-based cases; population based controls	Consecutive Russian breast cancer patients aged 24-86 years ascertained at one of the two participating oncology centers in Bashkortostan and Siberia through the years 2006-2008	Population controls aged 18-84 years recruited from a population study of different populations of Russia. Healthy volunteers (without any malignancy) were selected from the same geographical regions during the years 2002-2008	Medical records	Varies by participant	Bogdanov, N. et al. A nonsense mutation (E1978X) in the ATM gene is associated with breast cancer. <i>Breast Cancer Res</i> 11(8):207-211 (2009)
Study to Investigate the Genetics of In Situ Carcinoma of the Ductal Subtype	UK	Hospital-based case-control study	Cases aged 60 or younger with pure DCIS (no associated invasive cancer of any subtype) from 96 hospitals throughout the UK. Recruitment period was from Jul 2008 to Nov 2012	Controls were selected from pool of non-cancer patients who had no close relative (mother, sister, daughter or aunt) who had been so affected. Controls were recruited by asking non-blood relatives (generally sisters-in-law or friends) of affected individuals to act as a control. Recruitment period was from Jun 2007 to Jun 2013	Self reported by patient at time of recruitment	Not reported	No references yet
Karolinska Breast Cancer Study	Sweden	Population and hospital-based cases, geographically matched controls	1. Familial cases from Department of Clinical Genetics, Karolinska University Hospital, Stockholm. 2. Consecutive cases from Department of Oncology, Hudvinge & Söder Hospital, Stockholm 1998-2000	Blood donors of mixed gender from same geographical region. Excess material was received from all blood donors over a 3 month period in 2004 (approximately 2000) and DNA was extracted from a random sample of 1500	Medical records	2016	1) Wernöf, C. et al. Tumor spectrum in non-BRCA hereditary breast cancer families in Sweden. <i>Heredit</i> 2008; 110:163-171. 2) Margolin S, et al. BRCA1 mutations in a population-based study of breast cancer in Stockholm County, Genet. Test., 8, 127-32 (2004)
Karolinska Mammography Project for Risk Prediction of Breast Cancer - Cohort Study	Sweden	Cohort study	Inclusion of 70,877 women Oct 2010 - March 2013. 3000 women had BC at cohort entry. In all, 800 women have been diagnosed with breast cancer since study entry (Oct 2015). Approximately 250 women are diagnosed with BC annually	Non-BC cases in the Karolinska Cohort	Through the Swedish Cause of Death register; Clinical Breast Cancer register and the Inpatient register	We match the data sets to all registers twice a year	Submitted
Kuopio Breast Cancer Project	Finland	Population-based prospective cohort study	1. Women seen at Kuopio University Hospital between 1990 and 1995 because of breast lump, mammography abnormality, or other breast symptom who were found to have breast cancer. 2. Consecutive malignant breast cancer cases diagnosed at KUUM from 2011 onwards	Age and long-term area-of-residence matched controls selected from the National Population Register and interviewed in parallel with the cases	Follow-up is done by an oncologist	2016	1) Hartikainen, J.M. et al. An autosomal-wide scan for linkage disequilibrium-based association in sporadic breast cancer cases in eastern Finland: three candidate regions found. <i>Cancer Epidemiol Biomarkers Prev</i> 2008; 17(12):2211-22. 2) Hartikainen, J.M. et al. Refinement of the 22q12.3 (13) breast cancer-associated region: evidence of THRSPO as a candidate gene in an eastern Finnish population. <i>Clin Cancer Res</i> 12, 1454-1462 (2006)
Kathleen Cunningham Foundation Consortium for research into Familial Breast Cancer/Australian Ovarian Cancer Study	Australia and New Zealand	Clinic-based recruitment of familial breast cancer patients (cases); population-based case-control study of ovarian cancer (controls only)	Cases were from multiple case breast and breast-ovarian families recruited through family cancer clinics from across Australia and New Zealand from 1998 to the present. Cases were selected for inclusion in BRCA studies if (i) family was negative for mutations in BRCA1 and BRCA2 (ii) case was the index for the family, defined as youngest breast cancer affected family member	Female controls were ascertained by the Australian Ovarian Cancer Study identified from the electoral rolls from all over Australia from 2002-2006	Patent self and family reports, medical records	Ongoing, varies by participant	1) Mann, G.J. et al. Analysis of cancer risk and BRCA1 and BRCA2 mutation prevalence in the KCForFab familial breast cancer resource. <i>Breast Cancer Res</i> , 11(12):2008. 2) Beesley, J. et al. Association between single nucleotide polymorphisms in hormone metabolism and DNA repair genes and epithelial ovarian cancer: Results from two Australian studies and an additional validation set. <i>Cancer Epidemiol Biomarkers Prev</i> , 12, 2297-48 (2003)
Leuven Multidisciplinary Breast Centre	Belgium	Hospital-based case-control study	All patients diagnosed with breast cancer and seen in the Multidisciplinary Breast Centre in Leuven (Gasthuisberg) between 2007 plus retrospective collection of cases diagnosed since 2003.	Healthy controls (blood donors) collected at the Red Cross and located in Gasthuisberg hospital (Oct-2007-March 2008)	KWIS; the latest data covers all departments, not only when they come for the treatment	2013 in the database from BRCA. Normally fits in the KWIS when the patient has been there	1) Neven P, Brauwaert O, Van den V, Vandens Bempt Hendrickx W, Cho H, Derardt K, Van Calster B, Van Huffel S, Moerman P, Amant F, Leunen K, Smeets A, Wilders H, Christens R, Vergote I, Christens MB. In early-stage breast cancer, the estrogen receptor interacts with corepressors between human epidermal growth factor receptor 2 status and age at diagnosis, tumor grade, and lymph node involvement. <i>J Clin Oncol</i> 2008 Apr 15;26(15):1768-71. 2) De Maeyer, V, Van Limbergen E, De Hys K, Moerman P, Pochet N, Hendrickx W, Wilders H, Christens R, Smeets A, Christens MB, Vergote I, Leunen K, Amant F, Neven P. Does estrogen receptor negative/progesterone receptor positive breast carcinoma exist. <i>J Clin Oncol</i> 2008; 26:1592-3. 3)38
Macedonian Breast Cancer Study	Republic of North Macedonia	Hospital-based case-control study	Prospectively ascertained cases of breast cancer in two hospitals in Skopje, Macedonia from 2012 to 2014. Ethnic origin: Macedonians (~62.8%) and Albanians (~17.2%). Age of the cases: 29 to 86, mean: 53.8	Pregnant women without breast cancer undergoing prenatal screening for chromosomal aneuploidy from 2013-2014. Recruited in three hospitals in Skopje, two of which are the same as those for recruitment of cases. Controls were matched for ethnic origin with the cases. Age of the controls: 18-45, mean 31.1	No information available	Not reported	No References
Mammary Carcinoma Risk Factor Investigation	Germany	Population-based case-control study	Incident cases diagnosed from 2001-2005 in the study region Hamburg in Northern Germany, and from 2002-2005 in the study region Rhein-Neckar-Karlsruhe in Southern Germany	2 controls per case were randomly drawn from population registries and frequency matched by birth year and study region to the case. Controls were recruited from 2002 to 2006	Follow-up information was obtained through follow-up interviews/questionnaires and new events through medical records to verify clinical events either reported by treating physicians or self-reported during follow-up interviews. Vital status was obtained by requesting this information from the population registry	May 2016	Flisch-Jaays, D et al. Role of different histological types of postmenopausal breast cancer by type and regimen of menopausal hormone therapy. <i>Int J Cancer</i> 2008 Aug 15;123(4):933-41
Milan Breast Cancer Study Group	Italy	Clinic-based recruitment of familiarly onset breast cancer patients (cases); population-based controls	Familial and/or early onset breast cancer patients (aged <22-87) negative for mutations in BRCA genes, ascertained in two large cancer centres in Milan from 1990 to 2008	Healthy blood donors aged 18-71 years, recruited at two blood centres in Milan from 2004 (centre 1) and 2007 (centre 2) to 2009	80% Medical records; 5% Phone contact; 15% Referral by patients/family members	Informations on follow-up not routinely collected	1) De Vecchi et al. Evidence for association of the CASP9 452 G del promoter polymorphism with age at diagnosis in familial breast cancer cases (letter). <i>Breast Cancer Res Treat</i> 133(6):778-2009. 2) Calabuig et al. Letter to the editor: SNPs in ultraconserved elements and familial breast cancer risk. <i>Carcinogenesis</i> 30:544-545, 2009
Mayo Clinic Breast Cancer Study	USA	Hospital-based case-control study	Incident cases residing in 6 states (MN, WI, IA, IL, ND, SD) seen at the Mayo Clinic in Rochester, MN from 2002-5	Women without cancer presenting for general medical examination at the Mayo Clinic. Controls were recruited concurrently with cases and were frequency matched to cases on age, ethnicity and county/state	Not reported	Not reported	Olson, JE. et al. A comprehensive examination of CYP19 mutation and breast cancer. <i>Cancer Epidemiol Biomarkers Prev</i> , 16, 623-5 (2007)
Melbourne Collaborative Cohort Study	Australia	Prospective cohort study; nested case-control study	Incident cases diagnosed between baseline (1990-1994) and last follow-up (2012) among the 24469 women participating in the cohort	For each case a control was randomly selected from women from the cohort who did not develop breast cancer before the age at diagnosis of the case and matched the case by age at birth and county of birth	Record linkage to the national and state cancer and death registries	Record linkages are carried out at least annually	Giles GG, et al. The Melbourne Collaborative Cohort Study. <i>Lancet</i> 1991; 337:129-31
Multieethnic Cohort	USA	Prospective cohort study; nested case-control	Incident cases identified from SEER cancer registries in Los Angeles County & State registries in California & Hawaii, USA from 1993-2002. Grouped by self-reported ethnicity	Women without cancer from the same States, recruited concurrently with cases & frequency matched by age at blood-draw & self-reported ethnicity	Linkage to SEER registries, state vital statistics and National Death Index	Linkages are performed annually	Kolonel, L. N. et al. A multi-ethnic cohort in Hawaii and Los Angeles: Baseline characteristics. <i>Am J Epidemiol</i> , 151, 348-357 (2000)
Melanoma Inquiry of Southern Sweden	Sweden	Population-based prospective cohort study	Population based cohort of off women aged 25-65 in southern Sweden, born in Sweden, no cancer diagnosis before, interviewed about cancer risk factors 1960-2000/010, saliva sampled 2011, cancer incidence/mortality followed through registries	2 matched controls within the cohort	Cause of death registry, records	2016	1) Olsson HE, Hyger C, Baidarim A. Hormone replacement therapy containing progestins and given continuously increases breast carcinoma risk in Sweden. <i>Cancer</i> 2003 Mar;91(6):1387-92. 2) Nielsen K, Mørkbak A, Olesen H, Ingvar C. A prospective study of 40000 women regarding risk factors, UV exposure and sunbed use in relation to risk and anatomic site of malignant melanoma. <i>Br J Cancer</i> 2012; 117(10):1706-15
Mayo Mammography Health Study	USA	Prospective Cohort Study (2003-2005) of women age 35+ receiving screening mammography at Mayo Clinic and living in MN, IA, WI; nested case-control	Incident cases (invasive or in situ) diagnosed at least 3 months after enrollment	Two sets of controls. One set frequency matched to cases on age. Second set of premenopausal women with density measures	Multiple sources: linkage to registration/marriages	2014	Olson, JE, Selman, TA, Scott, CG, Schuler, BA, Brand, KR, Serie, DJ, Jensen, MR, Wu, FF, Morton, MJ, Heine, JF, Couch, FJ, Parkins, VS, Vachon, CM. The influence of mammographic acquisition on the mammographic density and breast cancer association in the Mayo Mammography Health Study cohort. <i>Breast Cancer Res</i> 2012; Nov 15;14(6):R147
Malaysian Breast Cancer Genetic Study	Malaysia	Hospital-based case-control study	Breast cancer cases identified at the Breast Cancer Clinic in University Malaya Medical Centre, Jan 2003-July 2014, and Subang Jaya Medical Centre Sep 2012-Sept 2014; cases are a mixture of prevalent and incident cases	Controls are cancer-free individuals (37-74 years) selected from women attending mammography screening at the same hospitals	From National Registry of Births and Deaths	Annual exercise, in January or February	Tan MM, Ho W-K, Yoon S-Y, Marapan S, Hasan SN, Lee BS-C, et al. (2018) A case-control study of genetic risk factors in 7863 women in Malaysia. <i>PLoS One</i> 13(9): e0202849. https://doi.org/10.1371/journal.pone.0202849

Norwegian Breast Cancer Study	NBCS*	Norway	Hospital-based case-control study	Incidence cases from three different hospitals: 1) Cases (114) mean age 64 (20-82) at Ullevål University Hospital 1990-94, 2) cases (182) mean age 59 (26-76) referred to Norwegian Radium Hospital 1975-1986, 3) cases (124) mean age 56 (24-82) with stage I or II disease, in the Oslo micro-metastases study at Norwegian Radium Hospital between 1995-1998, 4) Breast cancer cases referred to the Norwegian hospitals Akerhus University Hospital in Lørenskog, Ullevål university hospital in Oslo and Rikshospitalet-Radiumhospitalet in Oslo from 2007-2010. Mean age is 63 years. Consecutive series. 5) Breast cancer cases referred to the Norwegian Radium Hospital hospitalised 2010-2013. Neoadjuvantly treated with Avastin (Bevacizumab). 6) Consecutive series of Breast cancer incidents referred to Akerhus University Hospital 2004-2014.	Control subjects were healthy women, age 55-71, residing in Tromsø (400), and Bergen (109) attending the Norwegian Breast Cancer Screening Program. Healthy issue from mammographic reduction surgery at a private clinic in Oslo	Medical records	Every 5 years, follow up 5-20 years, different for all sub-cohorts	1) Aune et al. <i>Genome Med.</i> 2015 Feb 7;11(2): 2) Fleischer et al. <i>2014 Genome Biol.</i> 2014;15(8):435. 3) Fleischer et al. <i>2014 Int J Cancer.</i> 2014 Jun 11;134(11):2615-26. 4) Duggan et al. <i>2014 Mol Oncol.</i> 2014;14(12):1273-74
Nashville Breast Health Study	NBHS*	USA	Population-based case-control study	Through a rapid case-ascertainment system, we identified newly-diagnosed breast cancer cases through the Tennessee State Cancer Registry and five major hospitals in the city that provide medical care for breast cancer patients. Eligible cases were women diagnosed with invasive breast cancer or ductal carcinoma in situ, who were between the ages of 25 and 75, had no prior history of cancer other than non-melanoma skin cancer, had a resident telephone, spoke English, and who were able to provide consent to the study. Recruitment period was from 2001 to 2011. The recruitment for European Americans ended in 2008.	Controls were identified via random digit dialing (RDD) of households in the same geographic area as cases during 2001-2011. Eligibility criteria for controls were the same as cases with the exception that controls did not have a prior cancer diagnosis other than simple skin cancer. Controls were frequency matched to cases on 5-year age group, race, and county of residence	Not reported	Not reported	Zheng W, Long J, Gao Y, Li C, Zhang Y, Xiang YB, Wen W, Levy S, Dennis SL, Haines JL, Gu K, Fan AM, Cai Q, Lu W, Shu XO. Genome-wide association study identifies a new breast cancer susceptibility locus at 6q25.1. <i>Nature Genetics</i> 41(3):324-8, 2009. PMC
Northern California Breast Cancer Family Registry	NC-BCFR	USA	Population-based recruitment of families, family-based cohort; population-based controls for subset of cases	Incident breast cancer cases included women aged <65 years diagnosed from 1995-2009, identified through the SEER cancer registry of the Greater San Francisco Bay Area. All cases with indicators of increased genetic risk (i.e. age <35 yrs, personal history of ovarian or childhood cancer, bilateral breast cancer or cancer in first-degree relatives). Cases not meeting these criteria were randomly sampled (5% of non-Hispanic whites, 25% of other races/ethnicities). Incident cases also included men aged <80 years diagnosed from 1995-1996	1) Unaffiliated family members enrolled from 1995-2011. 2) Unaffiliated unrelated population controls identified through random digit dialing conducted from 1999-2001 in the San Francisco Bay Area. Controls were frequency matched to cases diagnosed from 1995-1998 on 5-year age group and race/ethnicity, at a ratio of 1 control per 2 cases.	Active follow-up by questionnaire and linkage with the California Cancer Registry. Annual phone follow-up from 1999-2012 to obtain updates on vital status and new cancers in the family. Updates on risk factors, vital status, and new cancers in family in 2007-2011, 2012-2014, and 2015-2017	2015-2017	1) John, E.M. et al. The Breast Cancer Family Registry: an infrastructure for cooperative multinational, interdisciplinary and translational studies of the genetic epidemiology of breast cancer. <i>Breast Cancer Res.</i> 6(3):R389-R399 (2004). 2) Terry M, Phillips K, Daly M, et al. Cohort Profile: The Breast Cancer Prospective Family Study Cohort (Pfam-SC). <i>International Journal of Epidemiology</i> 2015;44:1111-1118
North Carolina Breast Cancer Study	NBCBS	USA	NBCBS Phases 1 & 2: population-based case-control study NBCBS Phase 3: population-based case-only study	NBCBS Phase 1: women aged of 20-74 residing in the 24 North Carolina county area and diagnosed with a first primary invasive breast cancer from 1993-1996 NBCBS Phase 2: women aged of 20-74 residing in the same study area and diagnosed with a first primary invasive breast cancer from 1996-2000. All women diagnosed with DCIS, DCIS with microinvasion to a depth of at least 1.0mm, and mixed of DCIS & LCIS from 1998-2000 were also eligible NBCBS Phase 3: Study area was expanded to 44 NC counties. Women aged 20-74 residing in this area and diagnosed with a first primary invasive breast cancer from May, 2008 to July, 2013	NBCBS Phase 1 & 2: Invasive: controls were from the same county area and frequency matched to cases by race (African American vs. non-AA) and five-year age group (20-24, 25-29, 70-74) DCIS: controls from the same study area were frequency matched to cases by race and age groups (20-24, 25-29, 45-54, 65-74) NBCBS Phase 3: no controls	Vital status was obtained through linkage to the US National Death Index	NA	1) Newman, B., Mooman, P. G., Milikan, R., Qaqish, B. F., Gerards, J., Altrich, T. E., and Lu, E. T. The Carolina Breast Cancer Study: integrating population-based epidemiology and molecular biology. <i>Breast Cancer Res.</i> 2005; 7(5):R56-60. 2) Milikan, R., Eaton, A., Worely, K., Brocho, L., Hodgson, E., Huang, W., Yu, Gerards, J., Iacocca, M., Cowan, D., Conway, K., and Dresler, L. HER2 in 456 polymorphisms and risk of breast cancer in African Americans and whites. <i>Breast Cancer Res. Treat.</i> 79: 355-364, 2003
Oulu Breast Cancer Study	OBCS	Finland	Hospital-based case-control study	Consecutive incident cases diagnosed at the Oulu University Hospital between 2003 and 2004	Healthy, consecutive, anonymous, female Finnish Red-Cross blood donors recruited in 2002 from the same geographical region in Northern Finland	Hospital medical records, Cancer registry, population registry	2012 breast cancer spec/2010 overall	Ekin, H. et al. A recurrent mutation in PALB2 in Finnish cancer families. <i>Nature</i> 446, 316-319 (2007)
Ontario Familial Breast Cancer Registry	OFBCR	Canada	Population-based family case-control study	Cases diagnosed between 1 Jan 1996-31 Dec 1998 were identified from the Ontario Cancer Registry which registers 99% of all cases residing in the province at the time of diagnosis. All women with invasive breast cancer aged 20-84 years who met the OFBCR definition for high genetic risk (family history of specific cancer, bilateral breast and ovarian, early onset disease, Ashkenazi ethnicity or a diagnosis of multiple breast cancer) were invited to participate by completing risk factor questionnaires and providing a blood sample. A 25% random sample of individuals in this age category who did not meet the OFBCR definition, 30% of those aged 55-69 at high risk and 8.75% aged 50-59 at low risk were also asked to participate. Individuals diagnosed in 2001 and 2002 were also included if they met high-risk criteria	Unrelated, unaffiliated population controls were recruited between 2003-2005 by calling randomly selected residential telephone numbers throughout the same geographical region. Eligible controls were women with no history of breast cancer and were frequency-matched by 5-year age group to the expected age distribution of cases. Approximately 65% of identified eligible women returned questionnaires, and 63% of these donated a blood specimen	Follow-up data including vital status were collected through annual family history follow-up questionnaires and through personal history follow-up questionnaires collected at years 10, 15 and 20 since baseline. Vital status of cases was ascertained also through linkage to cancer registry	15-year personal history follow-up information were collected from April 2012 to April 2015	John, E.M. et al. The Breast Cancer Family Registry: an infrastructure for cooperative multinational, interdisciplinary and translational studies of the genetic epidemiology of breast cancer. <i>Breast Cancer Res.</i> 6(3):R389-R399 (2004)
Leiden University Medical Centre Breast Cancer Study	ORIGO	Netherlands	Hospital-based prospective cohort study	Consecutive cases diagnosed 1996-2006 in 2 hospitals of South-West Netherlands (Leiden & Rotterdam). No selection for family history. Rotterdam cases selected for diagnosis aged <70. Cases with in situ carcinomas eligible	Three groups of controls: (1) Blood bank healthy donors from Southwest Netherlands recruited in 1996, 2000 or 2007; (2) People who married a person who was part of a family with high breast cancer risk (BRCA1/2); (3) Females tested at the local clinical genetics department for familial diseases, including familial cancer syndromes (no mutation found in gene(s) related to the disease being tested), recruited 1995-2007	Linkage to Municipal Population Register; National Pathology Registry; Hospital Information System; General Practitioner	2015 for cases from LUMC (N=650); 2007 for cases from GGDW (N=860)	1) de Boek, G.H. et al. Tumour characteristics and prognosis of breast cancer patients carrying the germline CHEK2*1100del variant. <i>Journal of Medical Genetics</i> 41, 731-735 (2004). 2) Huijts PE et al. Clinical correlates of low-risk variants in FGFRL2, TNRC9, MAP3K1, LSP1 and BR2 in a Dutch cohort of incident breast cancer cases. <i>Breast Cancer Res.</i> 9, R78 (2007)
NCI Polish Breast Cancer Study	PBCS	Poland	Population-based case-control study	Incident cases from 2000-2003 identified through a rapid identification system in participating hospitals covering ~90% of all eligible cases, and cancer registries in Warsaw and Lodz covering 100% of all eligible cases	Randomly selected from population lists of all residents of Poland, stratified and frequency matched to cases by case city and age in 5-year categories. Recruited 2002-2003	Complete follow-up for Warsaw cases. Every vital status for Lodz cases. We are reviewing medical records plus Cancer Registry data and base plus Death Certificates data base	Every 5 years, last data from 2012, currently running follow up after 15 yrs	Barca-Closas, M. et al. Polymorphisms in DNA double-strand break repair genes and risk of breast cancer: two population based studies in USA and Poland, and meta-analyses. <i>Hum. Genet.</i> 119, 376-88 (2006)
Karolinska Mammography Project for Risk Prediction of Breast Cancer - Case-Control Study	pKARMA	Sweden	Case-control study	Incident cases from Jan 2001 - Dec 2008 from the Stockholm/Gotland area. Identified through the Stockholm breast cancer registry	Unmatched participants of the KARMA mammography screening study recruited between 2010 and 2011 from Helsingborg and Stockholm	Through the Swedish Cause of Death register; Clinical Breast Cancer register and the patient register	We match the data sets to all registers twice a year	Unpublished
Prospective Study of Outcomes in Sporadic Versus Hereditary Breast Cancer	POSH	UK	Prospective cohort	Cases aged 40 or younger at breast cancer diagnosis. Recruited from breast cancer centre oncology clinics across 126 UK hospitals and diagnosed between January 2000 to December 2007	No in-house controls	National data	Vital status updated every 6 months	1) Eccles et al. (2007) Prospective study of Outcomes in Sporadic versus Hereditary breast cancer (POSH): Study Protocol. <i>BMC Cancer</i> 7, 169. 2) Tepper et al. (2008). Association between common genetic variants and prognosis of early onset breast cancer in 30 candidate genes. <i>Breast Cancer Research</i> 10(108). 3) Coppson E et al. (2013) Prospective observational study of breast cancer treatment outcomes for UK women aged 18-40 years at diagnosis: The POSH study. <i>J Natl Cancer Inst</i> 105, 978-988. 4) Coppson, E., et al. (2014). Ethnicity and outcome of young breast cancer patients in the United Kingdom: The POSH study. <i>Br J Cancer</i> 110, 220-241. 5) Coppson, E. R., et al. (2015). Obesity and the outcome of young breast cancer patients in the UK: the POSH study. <i>Ann Oncol</i> 26, 101-112. 6) Gough T et al. (2015). Inherited mutations in 17 breast cancer susceptibility genes among a large triple-negative breast cancer cohort unselected for family history of breast cancer. <i>J Clin Oncol</i> 33, 304-311
Evaluation of Predictive Factors regarding the Effectivity of Aromatase Inhibitor Therapy	PREFACE	Germany	Multicenter, prospective, randomized, open-label phase IV study	Postmenopausal, steroid hormone receptor positive breast cancer patients who are treated with letrozole. Recruitment at multicenters in Germany between 2009-2011	No controls	Medical records	12/1/2014	No References
Predicting the Risk of Cancer At Screening Study	PROCAS	UK	Population based study	Women diagnosed with breast cancer since joining the study of women attending the Breast Screening Programme (NHSBSPP) in Greater Manchester. Recruitment period Oct 2009-May 2014	Women attending routine NHS breast screening in Greater Manchester without a breast cancer diagnosis. Recruited during the same period as for the cases	Cancer report updated monthly from NHS systems. Deaths report updated weekly from local GP records	Expected to be in follow up until 2020	Evans DG, Ashley S, Stavrinou P, Hartness E, Donnelly LS, Dawes S, et al. Improvement in risk prediction, early detection and prevention of breast cancer in the NHS Breast Screening Programme and family history clinics: a dual cohort study. <i>Programme Grants Appl Res</i> 2016:4111
Rotterdam Breast Cancer Study	RBCS	Netherlands	Hospital-based case-control study, Rotterdam area	Familial breast cancer patients selected from the Clinical Genetics Central Erasmus MC Cancer Institute, recruited 1994 - 2005 (RBCS1) and 1995 - 2009 (RBCS2; for Ductal) and 2005 - 2009 (RBCS2)	Spouses or mutation-negative siblings of heterozygous Cystic Fibrosis mutation carriers selected from the Clinical Genetics Center at Erasmus MC Cancer Institute, recruited 1996 - 2006 (RBCS1) and 2005 - 2009 (RBCS2)	From medical file/inform from GP or other hospital. Vital status also from Municipal registry	October 2013	1) Krijger, A. H. et al. Survival and contralateral breast cancer in CHEK2*1100del breast cancer patients: impact of adjuvant chemotherapy. <i>Br J Cancer</i> 2014; 111(6):1094-13
Singapore and Sweden Breast Cancer Study	SASBAC	Sweden	Population-based case-control study	Incident cases from October 1993 to March 1995 identified via the 6 regional cancer registries in Sweden, to which reporting is mandatory	Controls were randomly selected from the total population registry in 5-year age groups to match the expected age-frequency distribution among cases. Patients and controls were recruited from Oct 1993 through April 1995	Through medical records the first 6-8 years and the Swedish Cause of Death register, Clinical Breast Cancer register and the patient register thereafter	We match the data sets to registers when needed	Weiden, S. et al. Oestrogen receptor alpha gene haplotype and postmenopausal breast cancer risk: a case control study. <i>Breast Cancer Res.</i> 6, R437-49 (2004)
Sheffield Breast Cancer Study	SBCS	UK	Hospital-based case-control study	Women with pathologically confirmed breast cancer recruited from surgical outpatient clinics at the Royal Hallamshire Hospital, Sheffield, 1998 - 2005; cases are a mixture of prevalent and incident disease	Unselected women attending the Sheffield Mammography Screening Service between Sep 2000 - Aug 2004, of their mammograms showed no evidence of a breast lesion	Trent Cancer Registry	9th Sept 2013 (death data)	1) MacPherson, G. et al. Association of a common variant of the CASP8 gene with reduced risk of breast cancer. <i>Journal of the National Cancer Institute</i> 96, 1806-1809 (2004). 2) Palli, S. et al. A potential role for the XRCC2 R188H polymorphic site in DNA-damage repair and breast cancer. <i>Human Molecular Genetics</i> 11, 1433-1438 (2002)
Study of Epidemiology and Risk factors in Cancer Heredity	SEARCH	UK	Population-based case-control study	2 groups of cases identified through East Anglian Cancer Registry: 1) prevalent cases diagnosed 1991-1996 under 55 years of age at diagnosis, recruited 1996-2002; 2) incident cases diagnosed since 1996 under 70 years of age at diagnosis, recruited 1996-present	Two groups of controls: (1) selected from the EPIC-Norfolk cohort study of 25,000 individuals age 45-74 recruited between 1992 and 1994, based in the same geographic region as cases; (2) selected from GP practices from March 2000 to present, frequency matched to cases by age and geographic region	The National Cancer Registration and Analysis Service	8/1/2014	Lesueur, F. et al. Allelic association of the human homologue of the mouse modifier Plopi with breast cancer. <i>Hum. Mol. Genet.</i> 14, 2349-56 (2005)
Seoul Breast Cancer Study	SEBCS	Korea	Hospital-based case-control study	Consecutive, incident, cases from 2 hospitals in Seoul recruited 2001-2005	Healthy community controls from same catchment area and participating in annual health check-up, 2001-2005	Follow-up information was obtained irregularly as occasion demands	January 2013	1) Lee, K.M. et al. Genetic polymorphisms of ataxia telangiectasia mutated and breast cancer risk. <i>Cancer Epidemiol Biomarkers Prev.</i> 14, 821-5 (2005). 2) Han, S. et al. CASP8 polymorphisms, estrogen and progesterone receptor status, and breast cancer risk. <i>Breast Cancer Res. Treat.</i> 110, 387-393 (2008)
Singapore Breast Cancer Cohort	SGBCO	Singapore	Hospital-based breast cancer cohort and population-based controls	Living breast cancer patients diagnosed with primary in situ or invasive breast cancer at National University Hospital between 2006-2013. Cases are a mixture of prevalent and incident cases	All community-dwelling individuals who are Singaporeans or Singaporean Permanent Residents, 21 years and older. Participants were recruited between 2006 and 2010 through word-of-mouth and personal recommendations. In some cases, recruiters also sought participants through "cold-calling" or through door-to-door invitations. Exclusion criteria were a medical history of cancer, acute myocardial infarction or stroke, or major psychiatric morbidity including schizophrenia, psychosis, depression, and advanced Alzheimer's Disease	I visit only, no follow-up required	I visit only, no follow-up required	No References
Städtisches Klinikum Karlsruhe Deutsches Krebsforschungszentrum Study	SKDKDFZ*	Germany	Hospital-based breast cancer cohort	Women diagnosed with primary in situ or invasive breast cancer at the Städtisches Klinikum Karlsruhe from March 1993 to July 2005	No controls	Follow-up (including vital status) information was obtained from medical records, pathology reports or population registries. Risk factor data were collected from about 10% of patients by questionnaire; from the remaining patients risk factor data were obtained from the medical records	2012 - 2014	Stevens, K.N. et al. 9p13.1 is a triple-negative-specific breast cancer susceptibility locus. <i>Cancer Res.</i> 2012;72(7):1795-803
Swedish Mammography Cohort	SMC	Sweden	Nested case control study from a population-based cohort	All breast cancer cases in the cohort (information from the Swedish Cancer Register) from 1987-2011 for women who gave saliva in 2005-2008 or a blood specimen in 2003-2009 are included	Controls were randomly selected from cancer free women in the cohort who also gave saliva in 2005-2008 or a blood specimen in 2003-2009. Controls were matched to cases on birth year	Swedish Death Register & Swedish Cancer Register	End of 2014 for mortality, end of 2013 for cancer incidence, end of 2011 for ERPR receptors	Suzuki R1, Ye W, Rylander-Rudqvist T, Saij S, Colditz GA, Wolk A. Heritability and postmenopausal breast cancer risk defined by estrogen and progesterone receptor status: a prospective cohort study. <i>J Natl Cancer Inst.</i> 2006 Nov 2;97(11):1601-8

Simultaneous Study of Gemcitabine-Docetaxel Combination Adjuvant Treatment	SUCCESSB	Germany	Multicenter, prospective, randomized, open-label phase III study	Patients with primary Her2-positive and high risk breast cancer (pN+ or \geq T3b or \geq G1 or \geq 4ly or H5.) Recruitment at multicenters in Germany during 2008-2011	No controls	Medical records	5/1/2014	1) Andersen U, Neugebauer J, Jans W, Hepp P, Ortmann U, Semmer H, Raab B, Gips SS. (2011) Simultaneous study of gemcitabine-docetaxel combination adjuvant treatment, as well as biological targeted treatment: the SUCCESS B Trial. <i>Breast</i> 20:266-266. 2) Jaeger BAS, et al. (2012) HER2 expression on circulating tumor cells (CTCs) in patients with early HER2-positive breast cancer: Results of the German SUCCESS B trial. <i>J Clin Oncol</i> 30 (15). 3) Neugebauer JK, et al. (2013) Persistence of HER2 overexpression on circulating tumor cells in patients after systemic treatment for HER2-positive breast cancer: Follow-up results of the German Success B trial. <i>J Clin Oncol</i> 31 (15).
Simultaneous Study of Docetaxel Based Anthracycline Free Adjuvant Treatment Evaluation	SUCCESSC*	Germany	Multicenter, prospective, randomized, open-label phase III study	Patients with primary Her2-negative and high risk breast cancer (pN+ or \geq T3b or \geq G1 or \geq 4ly or H5.) Recruitment at multicenters in Germany during 2009-2011	No controls	Medical records	01/05/2014	1) Hepp P, et al. (2009) Simultaneous Study of Docetaxel Based Anthracycline Free Adjuvant Treatment Evaluation, as Well as Lifestyle Intervention Strategies Success C-Subst. <i>Anticancer Res</i> 29 (5):1567-1568. 2) Jaeger BA, et al. (2015) Persistence of circulating tumor cells immediately after and two years after systemic adjuvant chemotherapy in patients with early breast cancer - Results of the German SUCCESS trials. <i>Cancer Res</i> 75 (9). 3) Ortmann U, Salmen J, Hepp PGM, Beckmann MW, Fehm TN, Hindenburg H, Lichtenegger W, Raab BK, Schwenewe A, Jans W. (2011) The SUCCESS-C trial: Interim analysis of toxicity evaluating the role of an anthracycline-free chemotherapy regimen in the adjuvant treatment of HER2/neu-negative breast cancer. <i>J Clin Oncol</i> 29 (15). 4) Raab B, et al. (2010) The German SUCCESS-C Study - The First European Lifestyle Study on Breast Cancer. <i>Breast Care</i> 5 (6):395-400. doi:10.1159/00032877
HCC-Szczecin Breast Cancer Study	SZBCS	Poland	Hospital-based case-control study	Prospectively ascertained cases of invasive breast cancer patients diagnosed at the Regional Oncology Hospital (Szczecin) in the years 2002, 2003, 2008 and 2007 or at the University Hospital from 2002 to 2007 in Szczecin, West Pomerania, Poland. Patients with pure intraductal or in situ cancer were excluded (DCIS or LCIS) but patients with DCIS with micro-invasion were included	Unaffected, matched to cases for year of birth, sex and region; from families with negative cancer family history; controls were part of a population-based study of the 1.3 million inhabitants of West Pomerania performed in 2003 and 2004 designed to identify familial aggregations of cancer by sex centre	Questionnaire sent by post		About 10 years ago; currently we are sending questionnaires only to all mutation carriers (BRCA1/2, CHEK2, etc)
Triple Negative Breast Cancer Consortium Study	TNBCC	Various	See studies marked * and 0 studies below for details of individual studies in TNBCC					
Demokritos	DEMOKRITOS	Greece	Hospital-based case-control study	Triple negative breast cancer cases enrolled from 1997-2010 in hospitals serving geographical areas of Greece, including Athens metropolitan area, Thessaloniki, Ioannina, Patras, and Crete (Chania), in collaboration with the Hellenic Cooperative Oncology Group (HECOG)	Regional controls, identified between 2010-2011 from Athens and Thessaloniki, were population-based unaffected women of the same age range		Not reported	Not reported
A randomized phase III trial comparing nanoparticle-based paclitaxel with solvent-based paclitaxel as part of neoadjuvant chemotherapy for patients with early breast cancer	GEPARSEPT O	Germany	Multicenter, prospective, randomized, open-label phase III study	Patients with early primary breast cancer who are eligible for neoadjuvant chemotherapy. Recruitment at multicenters in Germany during 2011-2013.	No controls		Not reported	Not reported
University of Kansas Medical Center	KUMC	USA					Not reported	Not reported
Ohio State University	OSU	USA	Hospital-based case-control study	Incident triple negative invasive breast cancer cases from a collection of incident breast cancer cases diagnosed in Columbus, Ohio (2006-2011).	Population-matched controls accrued through primary care clinics in the OSU medical center system (2006-2011).		Not reported	Not reported
Roswell Park Cancer Institute	RPCI	USA	Hospital-based case-control study	Triple negative invasive breast cancer cases from incident cases recruited to the RPCI Data Bank and Biorepository from 2006-2010.	Healthy controls identified from employee volunteers, and women recruited from community events from 2006-2010.		Not reported	Not reported
University of Texas MD Anderson Cancer Center	UTMDACC	USA					Not reported	Not reported
Taiwanese Breast Cancer Study	TWBCS	Taiwan	Hospital-based case-control study	Incident cases diagnosed & treated at 2 major teaching hospitals in Taiwan (between March 2002 and August 2005)	Controls cancer-free individuals, randomly selected from women attending health exam. at same hospital during study period. Underwent 1-day health examination - any showing evidence cancer excluded	The information was from the cooperation of the hospitals	2013	1) Hsu, HM et al. Breast cancer risk is associated with genes encoding the DNA double-strand break repair Mre11/Rad50/Nbs1 complex. <i>Cancer Epidemiol Biomarkers Prev</i> 16: 2024-22 (2007). 2) Ding, SL et al. Genetic variants of BLM interact with RAD51 to increase breast cancer susceptibility. <i>Carcinogenesis</i> 30, 43-9 (2009)
UCI Breast Cancer Study	JCIBCS*	USA	Population-based case-control study	All cases diagnosed in Orange County, California, during one-year period beginning March 1, 1994. Ascertained through the population-based Cancer Surveillance Program of Orange County California (CSPOC)	Female controls under age 75 years without history of cancer recruited using random digit dialing among Orange County residents & frequency matched to cases by age & race/ethnicity. Recruited from 1998-2003.	The vital status and follow-up date are standard items of the California Cancer Registry	Completed information is as of 10/30/2013	1) Anton-Culver, H et al. Characteristics of BRCA1 mutations in a population-based case series of breast and ovarian cancer. <i>Eur J Cancer</i> 36, 1200-8 (2000). 2) Friggs, A et al. Cancer risk estimates for family members of a population-based family registry for breast and ovarian cancer. <i>Cancer Epidemiol Biomarkers Prev</i> 9, 1033-11 (2000).
			*Subset of samples are part of the TNBCC					

Variant (n=313) ^a	Chromosome	Position	Reference Allele	Effect Allele	Overall Breast Cancer (coefficient)	ER positive (coefficient) ^b	ER negative (coefficient) ^b	Imputation quality ^c	Imputation quality ^c	Imputation quality ^c	Imputation quality ^c
					(coefficient)	(coefficient) ^b	(coefficient) ^b	OncoArray (European) ^d	iCOGS (European) ^d	OncoArray (Asian) ^d	iCOGS (Asian) ^d
1_100880328_A_T	1	100880328	A	T	0.0373	0.0373	0.0373	1.0000	0.9200	1.0000	0.9060
1_10566215_A_G	1	10566215	A	G	-0.0586	-0.0407	-0.1109	1.0000	1.0000	1.0000	1.0000
1_110198125_CAAA_C	1	110198125	CAAA	C	0.0458	0.0458	0.0458	0.9980	0.9560	0.9980	0.9560
1_114445880_G_A	1	114445880	G	A	0.0621	0.0621	0.0621	0.9980	0.9980	0.9910	0.9980
1_118141492_A_C	1	118141492	A	C	0.0452	0.0452	0.0452	0.9970	0.9780	0.9920	0.9580
1_120257110_T_C	1	120257110	T	C	0.0385	0.0430	0.0226	0.9960	1.0000	0.9980	1.0000
1_121280613_A_G	1	121280613	A	G	0.0881	0.1052	0.0209	1.0000	1.0000	1.0000	1.0000
1_121287994_A_G	1	121287994	A	G	-0.0673	-0.0514	-0.1114	0.9920	0.9560	0.9970	0.9880
1_145604302_C_CT	1	145604302	C	CT	-0.0399	-0.0469	-0.0126	0.9380	0.9080	0.9360	0.8800
1_149906413_T_C	1	149906413	T	C	0.0548	0.0548	0.0548	1.0000	0.9720	1.0000	0.9790
1_155556971_G_A	1	155556971	G	A	0.0499	0.0499	0.0499	0.9950	0.9800	0.9430	0.9330
1_168171052_CA_C	1	168171052	CA	C	-0.0680	-0.0680	-0.0680	0.9000	0.7070	0.5120	0.4880
1_172238767_T_TA	1	172238767	T	TA	-0.0435	-0.0435	-0.0435	0.9200	0.8660	0.8450	0.8150
1_18807339_T_C	1	18807339	T	C	-0.0564	-0.0649	-0.0248	0.9990	0.8670	0.9960	0.8380
1_201437832_C_T	1	201437832	C	T	0.0917	0.0917	0.0917	1.0000	0.8200	1.0000	0.4860
1_202184600_C_T	1	202184600	C	T	-0.0065	0.0133	-0.0822	1.0000	0.9810	1.0000	0.9670
1_203770448_T_A	1	203770448	T	A	0.0498	0.0498	0.0498	0.9980	0.9930	0.9990	0.9920
1_204525214_T_TTCTGAAA	1	204525214	T	TTCTGAAA	-0.0321	-0.0024	-0.1345	0.9720	0.9430	0.8770	0.5350
1_208076291_G_A	1	208076291	G	A	-0.0366	-0.0366	-0.0366	0.9950	0.9640	0.9710	0.9390
1_217053815_T_G	1	217053815	T	G	0.0417	0.0417	0.0417	0.9030	0.7020	0.9180	0.5890
1_217220574_G_A	1	217220574	G	A	-0.0440	-0.0459	0.0029	0.9960	0.9990	0.9030	0.9570
1_217220574_G_A	1	220671050	G	A	0.0418	0.0418	0.0418	0.9520	0.8820	0.9550	0.8980
1_242032683_A_G	1	242032683	A	G	0.0428	0.0428	0.0428	1.0000	0.9420	1.0000	0.9150
1_41380440_C_T	1	41380440	C	T	0.0426	0.0426	0.0426	0.9840	0.7510	0.9760	0.5460
1_41389220_T_C	1	41389220	T	C	0.1550	0.1550	0.1550	0.9620	0.8450	0.9170	0.7310
1_46670206_TC_C	1	46670206	TC	C	0.0447	0.0595	0.0216	1.0000	0.9540	1.0000	0.9120
1_51467096_CT_C	1	51467096	CT	C	0.0374	0.0374	0.0374	0.9620	0.8890	0.8070	0.8080
1_7917076_G_A	1	7917076	G	A	-0.0427	-0.0409	-0.0409	0.9920	0.9300	0.9710	0.9100
1_88156923_G_A	1	88156923	G	A	0.0494	0.0580	0.0183	1.0000	1.0000	1.0000	1.0000
1_88428199_C_A	1	88428199	C	A	-0.0387	-0.0387	-0.0387	1.0000	1.0000	1.0000	1.0000
2_10138983_T_C	2	10138983	T	C	0.0603	0.0603	0.0603	0.9580	0.9340	0.7470	0.7480
2_121058254_A_G	2	121058254	A	G	-0.0334	-0.0232	-0.0882	0.9960	0.9990	0.9980	1.0000
2_121089931_T_C	2	121089931	T	C	0.0316	0.0316	0.0316	1.0000	0.5300	1.0000	0.4850
2_121159205_G_A	2	121159205	G	A	-0.0440	-0.0440	-0.0440	1.0000	0.8630	1.0000	0.7270
2_121246568_T_C	2	121246568	T	C	0.0992	0.0992	0.0992	1.0000	0.9970	1.0000	0.9930
2_172974566_C_G	2	172974566	C	G	-0.0473	-0.0611	-0.0601	1.0000	1.0000	1.0000	1.0000
2_174212810_A_G	2	174212810	A	G	0.0563	0.0561	0.0175	1.0000	0.9960	1.0000	0.9790
2_192381934_C_G	2	192381934	C	G	0.0316	0.0316	0.0316	1.0000	0.5300	1.0000	0.4850
2_19315675_T_A	2	19315675	T	A	-0.0331	-0.0229	-0.0570	1.0000	0.9960	1.0000	0.9910
2_202204741_T_C	2	202204741	T	C	-0.0492	-0.0492	-0.0492	1.0000	1.0000	1.0000	1.0000
2_217920769_G_T	2	217920769	G	T	-0.1318	-0.1532	-0.0589	1.0000	1.0000	1.0000	1.0000
2_217955986_GA_G	2	217955986	GA	G	-0.2016	-0.2062	-0.0960	0.9840	0.9040	0.9680	0.7980
2_218292158_C_G	2	218292158	C	G	-0.0257	-0.0757	0.0730	0.9770	0.9730	0.9870	0.9800
2_218714845_G_A	2	218714845	G	A	-0.0431	-0.0463	-0.0184	1.0000	0.6770	1.0000	0.7350
2_241388857_C_A	2	241388857	C	A	-0.1232	-0.1232	-0.1232	1.0000	0.7360	1.0000	0.4340
2_25129473_A_G	2	25129473	A	G	-0.0427	-0.0427	-0.0427	1.0000	0.7080	0.9990	0.7750
2_29179452_G_C	2	29179452	G	C	0.0066	0.0066	0.0066	0.9860	0.9690	1.0000	0.9000
2_29615233_T_C	2	29615233	T	C	-0.0427	-0.0427	-0.0427	0.9840	0.7710	0.9100	0.5720
2_39699510_C_CT	2	39699510	C	CT	-0.0402	-0.0402	-0.0402	0.9020	0.7580	0.6360	0.5350
2_70172587_G_A	2	70172587	G	A	-0.0412	-0.0412	-0.0412	0.9660	0.8050	0.9490	0.7930
2_88358825_G_C	2	88358825	G	C	0.0473	0.0473	0.0473	0.9430	0.5970	0.9490	0.5520
3_141112859_CTT_C	3	141112859	CTT	C	0.0551	0.0551	0.0551	0.9670	0.9670	0.9670	0.9670
3_17228237_G_A	3	17228237	G	A	0.0422	0.0501	-0.0133	0.9970	0.9970	0.9820	0.9850
3_189774456_C_T	3	189774456	C	T	-0.0478	-0.0478	-0.0478	0.9960	0.7060	0.9610	0.5980
3_27353716_C_A	3	27353716	C	A	0.0748	0.0822	0.0310	1.0000	1.0000	1.0000	1.0000
3_2738664_C_G	3	2738664	C	G	0.0502	0.0502	0.0502	1.0000	1.0000	1.0000	1.0000
3_29294945_C_T	3	29294945	C	T	-0.0288	-0.0288	-0.0288	0.9280	0.8480	0.9450	0.7350
3_30684907_C_T	3	30684907	C	T	0.0592	0.0557	0.0170	1.0000	1.0000	1.0000	1.0000
3_4688198_T_C	3	4688198	T	C	-0.0806	-0.0806	-0.0806	0.9930	0.9280	0.9940	0.8990
3_4742251_A_G	3	4742251	A	G	0.0616	0.0616	0.0616	1.0000	0.9990	1.0000	0.9970
3_49709912_C_CT	3	49709912	C	CT	-0.0387	-0.0355	-0.0721	0.9470	0.9370	0.8670	0.8190
3_5591077_A_CT	3	5591077	A	CT	-0.1195	-0.1195	-0.1195	0.9670	0.8480	0.9670	0.7540
3_59373745_C_T	3	59373745	C	T	-0.0394	-0.0394	-0.0394	0.9970	0.6400	0.9740	0.5500
3_63887449_T_TTG	3	63887449	T	TTG	0.0648	0.0648	0.0648	0.9880	0.9820	0.9980	0.9960
3_71620370_T_G	3	71620370	T	G	-0.0374	-0.0374	-0.0374	0.9860	0.8860	0.9880	0.8570
3_87037543_A_G	3	87037543	A	G	-0.0723	-0.0723	-0.0723	0.9430	1.0000	0.6670	1.0000
3_8940387_G_A	3	8940387	G	A	-0.0466	-0.0466	-0.0466	0.9570	0.9760	0.9760	0.9760
4_106069013_G_T	4	106069013	G	T	0.0471	0.0594	0.0097	1.0000	0.9980	1.0000	0.9930
4_126752992_A_AAT	4	126752992	A	AAT	-0.0377	-0.0377	-0.0377	0.9630	0.9640	0.9500	0.9310
4_143467195_C_T	4	143467195	C	T	-0.0569	-0.0569	-0.0569	0.9990	0.8140	0.9970	0.5800
4_151218296_CATATTT_C	4	151218296	CATATTT	C	0.0388	0.0388	0.0388	0.9910	0.9830	0.9830	0.9820
4_175842495_G_A	4	175842495	G	A	-0.0389	-0.0389	-0.0389	1.0000	0.9860	1.0000	0.9860
4_175847436_C_A	4	175847436	C	A	-0.0099	-0.0099	-0.0099	1.0000	0.9830	1.0000	0.9700
4_187503758_A_T	4	187503758	A	T	0.0357	0.0357	0.0357	0.9970	0.9700	0.9940	0.9730
4_38784633_G_T	4	38784633	G	T	0.0489	0.0489	0.0489	0.9990	1.0000	0.9980	1.0000
4_84370124_TAA_TA	4	84370124	TAA	TA	-0.0464	-0.0464	-0.0464	0.9440	0.9390	0.9560	0.9560
4_8920476_G_A	4	8920476	G	A	0.0352	0.0352	0.0352	1.0000	0.9860	1.0000	0.9860
4_92594859_TTCTTT_C	4	92594859	TTCTTT	C	-0.0407	-0.0407	-0.0407	0.9400	0.7990	0.9320	0.8320
5_104300273_G_T	5	104300273	G	T	-0.0487	-0.0487	-0.0487	0.9940	0.8240	0.9860	0.7320
5_122478676_C_A	5	122478676	C	A	-0.0386	-0.0386	-0.0386	0.9990	0.9790	0.9990	0.9910
5_122705244_C_T	5	122705244	C	T	0.0944	0.0944	0.0944	0.9970	1.0000	0.9870	0.9940
5_1279790_C_T	5	1279790	C	T	0.0817	0.0817	0.0817	1.0000	1.0000	1.0000	1.0000
5_1296255_A_AG	5	1296255	A	AG	-0.0549	-0.0417	-0.1056	1.0000	0.9960	1.0000	0.9880
5_131640536_G_G	5	131640536	G	G	0.0392	0.0467	0.0099	0.9810	0.9830	0.9550	0.9550
5_132407058_C_T	5	132407058	C	T	-0.0388	-0.0561	-0.0214	0.9990	0.7640	0.9980	0.7980
5_1335077_T_C	5	1335077	T	C	0.1552	0.1552	0.1552	1.0000	0.9030	1.0000	0.2850
5_158244083_C_T	5	158244083	C	T	-0.0677	-0.0677	-0.0677	1.0000	0.9850	1.0000	0.9700
5_16231194_G_C	5	16231194	G	C	-0.0426	-0.0426	-0.0426	1.0000	0.9990	1.0000	0.9980
5_169591460_T_C	5	169591460	T	C	0.0412	0.0501	0.0182	0.9970	0.9150	0.9980	0.9550
5_173358154_G_A	5	173358154	G	A	0.0365	0.0365	0.0365	0.9870	0.9920	0.9900	0.9920
5_176134882_T_C	5	176134882	T	C	0.0363	0.0363	0.0363	0.9950	0.7080	0.9970	0.6640
5_2177029_G_A	5	2177029	G	A	0.0391	0.0391	0.0391	0.9940	0.9850	0.9870	0.9870
5_32579616_TCA_T	5	32579616	TCA	T	0.0363	0.0363	0.0363	1.0000	0.9190	0.9980	0.9590
5_345109_T_C	5										

7_91459189_A_ATT	7	91459189	A	ATT	0.0452	0.0452	0.0452	0.9200	0.9310	0.9560	0.9650
7_94113799_T_C	7	94113799	T	C	0.0449	0.0449	0.0449	0.9950	0.9940	0.9990	0.9950
7_98025255_G_A	7	98025255	G	A	-0.0467	-0.0467	-0.0467	1.0000	1.0000	1.0000	1.0000
7_99948655_T_G	7	99948655	T	G	0.0420	0.0420	0.0420	0.9840	0.8900	0.9720	0.7640
8_102483100_T_C	8	102483100	T	C	0.0593	0.0736	0.0137	0.9500	0.9440	0.6000	0.8270
8_106358620_A_T	8	106358620	A	T	-0.0745	-0.0895	-0.1000	0.9790	0.7560	0.6000	0.6720
8_117209548_A_G	8	117209548	A	G	-0.0417	-0.0417	-0.0417	1.0000	0.9820	1.0000	0.9910
8_12082186_A_G	8	12082186	A	G	0.0267	0.0267	0.0267	0.9520	0.9680	0.9680	0.9670
8_124563705_T_C	8	124563705	T	C	0.0477	0.0477	0.0477	0.9900	0.9210	0.9840	0.8650
8_124571581_G_A	8	124571581	G	A	0.0340	0.0340	0.0340	0.9910	0.9550	0.9870	0.9680
8_124739913_T_G	8	124739913	T	G	0.0466	0.0466	0.0466	0.9810	0.9840	0.9670	0.9740
8_128213561_C_CA	8	128213561	C	CA	-0.0430	-0.0430	-0.0430	1.0000	0.8550	1.0000	0.7370
8_128370949_C_G	8	128370949	C	G	0.0472	0.0320	0.0372	0.9990	0.9990	0.9990	0.9990
8_128372172_A_G	8	128372172	A	G	0.0597	0.0597	0.0597	1.0000	1.0000	0.9980	0.9980
8_129199566_G_A	8	129199566	G	A	0.0615	0.0615	0.0615	1.0000	0.9970	1.0000	0.9980
8_143669254_A_G	8	143669254	A	G	-0.0346	-0.0346	-0.0346	0.9600	0.7790	0.9040	0.7900
8_170892_T_C	8	170892	T	C	0.0477	0.0348	0.1040	0.9230	0.7450	0.9590	0.7950
8_17761942_C_T	8	17761942	C	T	-0.0377	-0.0377	-0.0377	0.9320	0.8840	0.8370	0.7520
8_23447496_A_G	8	23447496	A	G	-0.0389	-0.0389	-0.0389	1.0000	1.0000	1.0000	1.0000
8_23663653_C_A	8	23663653	C	A	0.0335	0.0451	0.0059	0.9990	0.9990	0.9990	0.9980
8_29509616_A_C	8	29509616	A	C	-0.0601	-0.0601	-0.0601	1.0000	1.0000	1.0000	1.0000
8_36858483_A_G	8	36858483	A	G	-0.0760	-0.0760	-0.0760	1.0000	0.9910	1.0000	0.9890
8_76230943_A_G	8	76230943	A	G	0.0755	0.0755	0.0755	1.0000	1.0000	0.9970	0.9950
8_76333056_C_T	8	76333056	C	T	0.1129	0.1129	0.1129	1.0000	0.9970	1.0000	0.9800
8_76378165_G_T	8	76378165	G	T	-0.0391	-0.0391	-0.0391	1.0000	0.9400	1.0000	0.9170
9_110303808_TAA_T	9	110303808	TAA	T	0.0797	0.1007	0.1030	0.9920	0.9120	0.9950	0.9400
9_110837073_A_G	9	110837073	A	G	0.1158	0.1315	0.0289	1.0000	1.0000	1.0000	1.0000
9_110837176_C_T	9	110837176	C	T	0.0859	0.0859	0.0859	0.9990	0.9990	0.9990	0.9990
9_110849525_G_T	9	110849525	G	T	0.0153	0.0153	0.0153	1.0000	0.8920	1.0000	0.6520
9_110885479_C_T	9	110885479	C	T	0.0877	0.1110	0.0019	1.0000	1.0000	1.0000	0.9830
9_119313486_A_G	9	119313486	A	G	-0.0462	-0.0462	-0.0462	0.9880	0.9820	0.9830	0.9700
9_129424719_A_G	9	129424719	A	G	-0.0382	-0.0382	-0.0382	0.9950	0.9030	0.9080	0.8690
9_136146597_C_T	9	136146597	C	T	0.0400	0.0400	0.0400	0.9980	0.9860	0.9860	0.9860
9_21964882_CAAA_C	9	21964882	CAAAA	C	0.0550	0.0550	0.0550	0.9710	0.9650	0.8960	0.9020
9_22041998_C_G	9	22041998	C	G	0.0289	0.0168	0.0906	1.0000	1.0000	1.0000	1.0000
9_36928288_T_C	9	36928288	T	C	0.0249	0.0249	0.0249	0.9940	0.9800	0.9880	0.9720
9_6880263_A_G	9	6880263	A	G	0.0348	0.0499	-0.0078	1.0000	0.8470	1.0000	0.8880
9_87782211_T_C	9	87782211	T	C	0.0381	0.0381	0.0381	1.0000	1.0000	1.0000	1.0000
9_98362587_T_C	9	98362587	T	C	0.0576	0.0576	0.0576	0.9920	0.9790	0.9810	0.9860
10_114777670_C_T	10	114777670	C	T	0.0472	0.0472	0.0472	1.0000	0.9990	1.0000	0.9990
10_115128491_T_C	10	115128491	T	C	-0.0592	-0.0592	-0.0592	1.0000	1.0000	1.0000	1.0000
10_123095209_G_A	10	123095209	G	A	-0.0538	-0.0702	0.0048	0.9970	0.9640	0.9940	0.9500
10_123340107_A_G	10	123340107	A	G	0.0509	0.0509	0.0509	0.9990	0.9990	0.9990	0.9980
10_123340431_GC_G	10	123340431	GC	G	-0.2408	-0.2913	-0.0326	0.9990	0.9980	0.9990	0.9950
10_123349324_A_T	10	123349324	A	T	-0.2609	-0.3270	-0.0137	0.9660	0.9380	0.9880	0.6880
10_13892298_G_A	10	13892298	G	A	0.0371	0.0371	0.0371	1.0000	0.9740	1.0000	0.9390
10_22032942_A_G	10	22032942	A	G	-0.0580	-0.0719	0.0344	1.0000	1.0000	1.0000	1.0000
10_22477776_ACC_A	10	22477776	ACC	A	0.0472	0.1687	0.0363	0.9980	0.9930	0.9930	0.9910
10_22861490_A_C	10	22861490	A	C	0.0875	0.0960	0.0201	0.9770	0.9740	0.9440	0.9490
10_38523626_C_A	10	38523626	C	A	0.0404	0.0404	0.0404	0.9510	0.9410	0.9940	0.9890
10_5794652_A_G	10	5794652	A	G	0.0470	0.0470	0.0470	1.0000	1.0000	1.0000	1.0000
10_64299890_A_G	10	64299890	A	G	-0.1345	-0.1428	-0.1030	0.9830	0.9820	0.9940	0.9920
10_64819996_G_T	10	64819996	G	T	0.0472	0.0472	0.0472	0.9990	0.9990	0.9990	0.9990
10_71335574_C_T	10	71335574	C	T	-0.0404	-0.0404	-0.0404	0.9580	0.7060	0.9060	0.5620
10_80851257_G_T	10	80851257	G	T	-0.0805	-0.0898	-0.0443	1.0000	1.0000	1.0000	1.0000
10_80886726_A_G	10	80886726	A	G	0.0762	0.0762	0.0762	0.9970	0.9970	1.0000	0.9890
10_95292187_CAA_C	10	95292187	CAA	C	-0.0512	-0.0512	-0.0512	0.9430	0.8730	0.9000	0.8380
11_103614438_T_G	11	103614438	T	G	0.0147	0.0147	0.0147	0.9990	0.7650	0.9990	0.7650
11_108267402_C_CA	11	108267402	C	CA	-0.0022	0.0141	-0.0629	0.9980	0.9940	1.0000	0.9960
11_111696440_T_C	11	111696440	T	C	0.0396	-0.0396	-0.0396	0.9980	0.9980	0.9980	0.9980
11_116727936_A_T	11	116727936	A	T	-0.0423	-0.0423	-0.0423	0.9980	0.9970	0.9990	0.9980
11_122966626_A_G	11	122966626	A	G	-0.0383	-0.0383	-0.0383	0.9980	0.9080	0.9970	0.8500
11_129243417_T_G	11	129243417	T	G	-0.0543	-0.0543	-0.0543	0.9990	0.9990	0.9990	0.9990
11_129461016_A_G	11	129461016	A	G	0.0453	0.0453	0.0453	1.0000	0.9990	1.0000	0.9930
11_18664241_T_G	11	18664241	T	G	0.0461	0.0461	0.0461	0.9550	0.7170	0.9400	0.7390
11_1895708_C_A	11	1895708	C	A	-0.0782	-0.0782	-0.0782	0.9870	1.0000	0.9880	0.9880
11_42644441_C_T	11	42644441	C	T	-0.0336	-0.0336	-0.0336	0.9800	0.8070	1.0000	0.7670
11_4338117_T_G	11	4338117	T	G	-0.0437	-0.0437	-0.0437	0.9900	0.9400	0.9940	0.9340
11_44368892_G_A	11	44368892	G	A	0.0374	0.0374	0.0374	0.9930	0.9890	0.9940	0.9930
11_46318032_C_G	11	46318032	C	G	-0.0748	-0.0748	-0.0748	0.9280	0.8000	0.9380	0.4530
11_65553492_C_A	11	65553492	C	A	0.0425	0.0425	0.0425	0.9970	0.9870	0.9980	0.9770
11_65572431_G_A	11	65572431	G	A	-0.0347	-0.0347	-0.0347	0.9990	0.9990	0.9990	0.9990
11_69328130_A_T	11	69328130	A	T	-0.0423	-0.0538	-0.0067	0.9540	0.9610	0.9300	0.9360
11_69330983_G_A	11	69330983	G	A	0.1022	0.1240	0.0174	1.0000	0.9990	0.9990	0.9990
11_69331418_C_T	11	69331418	C	T	0.1782	0.2018	0.0066	0.9910	0.9910	0.9990	0.9990
11_803017_A_G	11	803017	A	G	0.0457	0.0457	0.0457	0.9940	0.9930	0.9970	1.0000
12_10309787_C_T	12	10309787	C	T	0.0565	0.0565	0.0565	1.0000	0.9740	0.9740	0.9190
12_111601034_T_G	12	111601034	T	G	-0.0442	-0.0442	-0.0442	1.0000	0.8830	1.0000	0.7300
12_115108136_T_C	12	115108136	T	C	0.0465	0.0465	0.0465	0.9700	0.9700	1.0000	0.9310
12_115796577_A_G	12	115796577	A	G	-0.0428	-0.0643	-0.0148	1.0000	0.9990	0.9990	0.9950
12_115835836_T_C	12	115835836	T	C	-0.0813	-0.0977	-0.0153	1.0000	1.0000	1.0000	1.0000
12_120832146_C_T	12	120832146	C	T	0.0516	0.0516	0.0516	1.0000	0.9740	0.9740	0.9140
12_14413931_G_C	12	14413931	G	C	0.0484	0.0484	0.0484	1.0000	1.0000	1.0000	1.0000
12_28149568_C_T	12	28149568	C	T	-0.0620	-0.0620	-0.0620	1.0000	1.0000	1.0000	1.0000
12_28174817_C_T	12	28174817	C	T	-0.0856	-0.0856	-0.0856	1.0000	1.0000	1.0000	1.0000
12_2847382_C_T	12	2847382	C	T	-0.0521	-0.0521	-0.0521	0.9760	0.9810	0.8190	0.9360
12_29140260_G_A	12	29140260	G	A	0.0647	0.0647	0.0647	0.9980	0.9740	0.9920	0.9170
12_293626_A_G	12	293626	A	G	0.0401	0.0401	0.0401	0.9950	0.8100	0.9950	0.8350
12_57146069_T_G	12	57146069	T	G	-0.0579	-0.0579	-0.0579	1.0000	1.0000	1.0000	1.0000
12_70798355_A_T	12	70798355	A	T	0.0469	0.0469	0.0469	0.9960	0.9330	0.9870	0.9910
12_83064195_G_GA	12	83064195	G	GA	0.0671	0.0671	0.0671	0.9700	0.4990	0.9390	0.4820
12_85034551_C_T	12	85034551	C	T	0.0348	0.0348	0.0348	0.9870	0.9970	0.9970	0.9990
12_96027759_A_G	12	96027759	A	G	-0.0867	-0.0867	-0.0867	1.0000	1.0000	1.0000	1.0000
13_32839990_G_A	13	32839990	G	A	0.0424	0.0424	0.0424	1.0000	0.7550	1.0000	0.4080
13_32972626_A_T	13	32972626	A	T	0.2887	0.2308	0.4284	1.0000	1.0000	1.0000	1.0000
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19_13249921_G_T	19	13249921	G	T	0.0956	0.0956	0.0956	0.9820	0.8510	0.7220	0.5750
19_17393925_C_A	19	17393925	C	A	0.0378	0.0036	0.1692	1.0000	1.0000	1.0000	0.9990
19_18589492_C_T	19	18589492	C	T	-0.0719	-0.0719	-0.0719	1.0000	0.9960	1.0000	0.9990
19_19517054_C_CGGGCG	19	19517054	C	CGGGCG	0.0437	0.0437	0.0437	0.9990	0.9470	0.9980	0.9440
19_44283031_T_C	19	44283031	T	C	0.0619	0.0619	0.0619	1.0000	0.8500	1.0000	0.8770
19_46166073_T_C	19	46166073	T	C	-0.0360	-0.0447	-0.0117	0.9500	0.8260	0.9430	0.7760
19_55816678_C_T	19	55816678	C	T	-0.0359	-0.0359	-0.0359	0.9910	0.9890	0.9860	0.9870
20_11379842_T_C	20	11379842	T	C	0.0844	0.0844	0.0844	0.9500	0.7690	0.9010	0.5330
20_41613706_C_G	20	41613706	C	G	0.0315	0.0266	0.0784	0.9490	0.9150	0.9260	0.8390
20_52296849_G_A	20	52296849	G	A	0.0440	0.0539	0.0144	0.9520	1.0000	0.9150	1.0000
20_5948227_G_A	20	5948227	G	A	0.0760	0.0760	0.0760	1.0000	1.0000	1.0000	1.0000
21_16364756_T_G	21	16364756	T	G	0.0646	0.0646	0.0646	1.0000	0.6290	1.0000	0.6540
21_16566350_A_G	21	16566350	A	G	0.0595	0.0678	0.0172	1.0000	0.9710	1.0000	0.8950
21_16574455_C_A	21	16574455	C	A	-0.0707	-0.0808	-0.0329	0.9980	0.9920	0.9940	0.9880
21_47762932_G_A	21	47762932	G	A	0.0946	0.0946	0.0946	0.9770	0.9370	0.9240	0.6980
22_19766137_C_T	22	19766137	C	T	-0.0367	-0.0367	-0.0367	0.9850	0.9370	0.9710	0.9130
22_29121087_A_G	22	29121087	A	G	0.1839	0.2812	-0.1566	1.0000	1.0000	1.0000	0.0030
22_29135543_G_A	22	29135543	G	A	0.0654	0.0654	0.0654	0.9970	0.9980	0.9950	0.9900
22_29203724_C_T	22	29203724	C	T	0.1405	0.1793	0.0191	1.0000	0.9480	0.9670	0.8150
22_29551872_A_G	22	29551872	A	G	-0.1716	-0.1716	-0.1716	0.9140	0.9080	0.8540	0.6280
22_38583315_AAAAG_AAA	22	38583315	AAAAG	AAAAGAAAG	-0.0471	-0.0608	0.0079	0.9590	0.9360	0.9800	0.9640
22_39343916_T_A	22	39343916	T	A	0.0407	0.0407	0.0407	1.0000	0.8980	1.0000	0.7970
22_40904707_CT_C	22	40904707	CT	C	0.1148	0.1148	0.1148	0.9680	0.9630	0.9960	0.9830
22_43433100_C_T	22	43433100	C	T	-0.0600	-0.0600	-0.0600	0.9940	0.9960	0.9890	0.9920
22_45319953_G_A	22	45319953	G	A	-0.0134	-0.0060	-0.0611	0.9960	0.9200	0.9980	0.9200
22_46283297_G_A	22	46283297	G	A	0.0736	0.0736	0.0736	0.9730	0.6320	0.7520	0.4790

^a Previously published by Mavaddat et al. (2019)

^b ER-specific PRS was constructed using a hybrid method, as described by Mavaddat et al. (2015)

^c If $r^2 = 1$ the variant was genotype