**Prediction of contralateral breast cancer:   
External validation of risk calculators in 20 international cohorts**

Daniele Giardiello1, 2, Michael Hauptmann3, 4, Ewout W. Steyerberg2, 5, Muriel A. Adank6, Delal Akdeniz7, Jannet C. Blom7, Carl Blomqvist8, 9, Stig E. Bojesen10-12, Manjeet K. Bolla13, Mariël Brinkhuis14, Jenny Chang-Claude15, 16, Kamila Czene17, Peter Devilee18, 19, Alison M. Dunning20, Douglas F. Easton13, 20, Diana M. Eccles21, Peter A. Fasching22, 23, Jonine Figueroa24-26, Henrik Flyger27, Montserrat García-Closas26, 28, Lothar Haeberle23, Christopher A. Haiman29, Per Hall17, 30, Ute Hamann31, John L. Hopper32, Agnes Jager33, Anna Jakubowska34, 35, Audrey Jung15, Renske Keeman1, Linetta B. Koppert36, Iris Kramer1, Diether Lambrechts37, 38, Loic Le Marchand39, Annika Lindblom40,41,, Jan Lubiński34, Mehdi Manoochehri31, Luigi Mariani42, Heli Nevanlinna43, Hester S.A. Oldenburg44, Saskia Pelders7, Paul D.P. Pharoah13,20, Mitul Shah20, Sabine Siesling45, Vincent T.H.B.M. Smit18, Melissa C. Southey46, 47, William J. Tapper48, Rob A.E.M. Tollenaar49, Alexandra J. van den Broek1, Carolien H.M. van Deurzen50, Flora E. van Leeuwen51, Chantal van Ongeval52, Laura J. Van't Veer1, Qin Wang13, Camilla Wendt53, Pieter J. Westenend54, Maartje J. Hooning7, Marjanka K. Schmidt1, 51

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

2 Leiden University Medical Center, Department of Biomedical Data Sciences, Leiden, The Netherlands.

3 Brandenburg Medical School, Institute of Biostatistics and Registry Research, Neuruppin, Germany.

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

5 Erasmus MC Cancer Institute, Department of Public Health, Rotterdam, The Netherlands.

6 The Netherlands Cancer Institute - Antoni van Leeuwenhoek hospital, Family Cancer Clinic, Amsterdam, The Netherlands.

7 Erasmus MC Cancer Institute, Department of Medical Oncology, Family Cancer Clinic, Rotterdam, The Netherlands.

8 University of Helsinki, Department of Oncology, Helsinki University Hospital, Helsinki, Finland.

9 Örebro University Hospital, Department of Oncology, Örebro, Sweden.

10 Copenhagen University Hospital, Copenhagen General Population Study, Herlev and Gentofte Hospital, Herlev, Denmark.

11 Copenhagen University Hospital, Department of Clinical Biochemistry, Herlev and Gentofte Hospital, Herlev, Denmark.

12 University of Copenhagen, Faculty of Health and Medical Sciences, Copenhagen, Denmark.

13 University of Cambridge, Centre for Cancer Genetic Epidemiology, Department of Public Health and Primary Care, Cambridge, UK.

14 Laboratory for Pathology, East-Netherlands, Hengelo, The Netherlands.

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

16 University Medical Center Hamburg-Eppendorf, Cancer Epidemiology Group, University Cancer Center Hamburg (UCCH), Hamburg, Germany.

17 Karolinska Institutet, Department of Medical Epidemiology and Biostatistics, Stockholm, Sweden.

18 Leiden University Medical Center, Department of Pathology, Leiden, The Netherlands.

19 Leiden University Medical Center, Department of Human Genetics, Leiden, The Netherlands.

20 University of Cambridge, Centre for Cancer Genetic Epidemiology, Department of Oncology, Cambridge, UK.

21 University of Southampton, Cancer Sciences Academic Unit, Faculty of Medicine, Southampton, UK.

22 University of California at Los Angeles, David Geffen School of Medicine, Department of Medicine Division of Hematology and Oncology, Los Angeles, CA, USA.

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

24 The University of Edinburgh Medical School, Usher Institute of Population Health Sciences and Informatics, Edinburgh, UK.

25 Cancer Research UK Edinburgh Centre, Edinburgh, UK.

26 National Cancer Institute, National Institutes of Health, Department of Health and Human Services, Division of Cancer Epidemiology and Genetics, Bethesda, MD, USA.

27 Copenhagen University Hospital, Department of Breast Surgery, Herlev and Gentofte Hospital, Herlev, Denmark.

28 Institute of Cancer Research, Division of Genetics and Epidemiology, London, UK.

29 University of Southern California, Department of Preventive Medicine, Keck School of Medicine, Los Angeles, CA, USA.

30 Södersjukhuset, Department of Oncology, Stockholm, Sweden.

31 German Cancer Research Center (DKFZ), Molecular Genetics of Breast Cancer, Heidelberg, Germany.

32 The University of Melbourne, Centre for Epidemiology and Biostatistics, Melbourne School of Population and Global Health, Melbourne, Victoria, Australia.

33 Erasmus MC Cancer Institute, Department of Medical Oncology, Rotterdam, The Netherlands.

34 Pomeranian Medical University, Department of Genetics and Pathology, Szczecin, Poland.

35 Pomeranian Medical University, Independent Laboratory of Molecular Biology and Genetic Diagnostics, Szczecin, Poland.

36 Erasmus MC Cancer Institute, Department of Surgical Oncology, Rotterdam, The Netherlands.

37 VIB, VIB Center for Cancer Biology, Leuven, Belgium.

38 University of Leuven, Laboratory for Translational Genetics, Department of Human Genetics, Leuven, Belgium.

39 University of Hawaii Cancer Center, Epidemiology Program, Honolulu, HI, USA.

40 Karolinska Institutet, Department of Molecular Medicine and Surgery, Stockholm, Sweden.

41 Karolinska University Hospital, Department of Clinical Genetics, Stockholm, Sweden.

42 Fondazione IRCCS Istituto Nazionale dei Tumori, Unit of Clinical Epidemiology and Trial Organization, Milan, Italy.

43 University of Helsinki, Department of Obstetrics and Gynecology, Helsinki University Hospital, Helsinki, Finland.

44 The Netherlands Cancer Institute - Antoni van Leeuwenhoek Hospital, Department of Surgical Oncology, Amsterdam, The Netherlands.

45 Netherlands Comprehensive Cancer Organisation, Department of Research, Utrecht, The Netherlands.

46 Monash University, Precision Medicine, School of Clinical Sciences at Monash Health, Clayton, Victoria, Australia.

47 The University of Melbourne, Department of Clinical Pathology, Melbourne, Victoria, Australia.

48 University of Southampton, Faculty of Medicine, Southampton, UK.

49 Leiden University Medical Center, Department of Surgery, Leiden, The Netherlands.

50 Erasmus MC Cancer Institute, Department of Pathology, Rotterdam, The Netherlands.

51 The Netherlands Cancer Institute - Antoni van Leeuwenhoek Hospital, Division of Psychosocial Research and Epidemiology, Amsterdam, The Netherlands.

52 Leuven Cancer Institute, University Hospitals Leuven, Leuven Multidisciplinary Breast Center, Department of Oncology, Leuven, Belgium.

53 Karolinska Institutet, Department of Clinical Science and Education, Södersjukhuset, Stockholm, Sweden.

54 Laboratory for Pathology, Dordrecht, The Netherlands.

**Corresponding author:** Marjanka K Schmidt, Netherlands Cancer Institute, Plesmanlaan 121, 1066 CX Amsterdam, The Netherlands; +31205122767; [mk.schmidt@nki.nl](mailto:mk.schmidt@nki.nl)

**Abstract**

**Background:**

Three tools are currently available to predict the risk of contralateral breast cancer (CBC). We aim to compare the performance of the Manchester formula, CBCrisk, and PredictCBC in patients with invasive breast cancer (BC).

**Methods:**

We analyzed data of 132,756 patients (4,682 CBC) from 20 international studies with a median follow-up of 8.8 years. Prediction performance included discrimination, quantified as a time-dependent Area-Under-the-Curve (AUC) at 5 and 10 years after diagnosis of primary BC, and calibration, quantified as the expected-observed (E/O) ratio at 5 and 10 years and the calibration slope.

**Results:**

The AUC at 10 years was around 0.6: 0.58 (95% confidence intervals [CI]:0.57–0.59) for CBCrisk; 0.60 (95%CI:0.59–0.61) for the Manchester formula; 0.63 (95%CI:0.59–0.66) and 0.59 (95%CI:0.56–0.62) for PredictCBC-1A (for settings where *BRCA1/2* mutation status is available) and PredictCBC-1B (for the general population), respectively. The E/O at 10 years was close to 1 for all models: 0.82 (95%CI: 0.51–1.32) for CBCrisk; 1.53 (95%CI:0.63–3.73) for the Manchester formula; 1.28 (95%CI:0.63–2.58) for PredictCBC-1A and 1.35 (95%CI:0.65–2.77) for PredictCBC-1B. The calibration slope was close to 1 for CBCrisk (1.26, 95%CI:1.01–1.50) and PredictCBC-1A and 1B (0.90, 95%CI:0.79–1.02 and 0.81, 95%CI:0.63–0.99), while the Manchester formula overestimated the CBC risk (slope 0.39, 95%CI:0.34–0.43).

**Conclusions:**

Current CBC risk prediction tools provide only moderate discrimination. Better predictors and re-calibration are needed to improve CBC prediction and to identify low and high CBC risk patients for clinical decision making.

**Keywords:** contralateral breast cancer, risk prediction, validation, clinical decision making

**Introduction**

A rising number of women with breast cancer (BC) are at risk to develop a new primary tumor in the contralateral breast (CBC) with consequently another cancer treatment and potentially less favorable prognosis[1]. Although CBC incidence is low (~0.4% per year) in the general BC population, contralateral preventive mastectomy (CPM) is increasing, also among women with low CBC risk[2-5].

Three tools are tools currently available to predict the risk of CBC, although probably none are widely used: 1) the Manchester formula; 2) CBCrisk, and 3) PredictCBC[6-8]. The Manchester group in the United Kingdom (UK) proposed a set of guidelines for counseling women about CPM[8]. Based on a systematic review of the literature, they devised a formula to estimate lifetime CBC risk based on age at first primary BC, family history of BC, estrogen-receptor (ER) status, diagnosis of ductal carcinoma in situ (DCIS), and oophorectomy.

The second tool, CBCrisk, was developed using data on 1,921 CBC cases and 5,763 matched controls with primary BC[7]. The model uses data on age at first BC diagnosis, age at first birth, first degree family history of BC, high-risk pre-neoplasia, breast density (obtained using the BI-RADS system), ER status, first BC type (pure invasive, pure DCIS, a mix of the two, unknown), and adjuvant endocrine therapy. External validation was performed using two independent studies in the United States (US) of 5,185 and 6,035 patients with 111 and 117 CBC events[7,9]. A web-based application provides individualized prediction of CBC risk[10].

Third, PredictCBC was developed, cross-validated and evaluated using data from 132,756 patients with first BC and 4,672 CBC events, as part of an international collaboration[5]. PredictCBC predicts CBC risk as a function of family history (first degree) of primary BC, and information of primary BC diagnosis: age, nodal status, size, grade, morphology, ER status, human epidermal growth factor receptor 2 (HER2) status, administration of adjuvant or neoadjuvant chemotherapy, adjuvant endocrine therapy, adjuvant trastuzumab therapy, and radiotherapy. Two versions were developed: PredictCBC version 1A includes presence or absence of a mutation in the *BRCA1* or *BRCA2* genes, an important determinant of CBC[5,11,12], while PredictCBC version 1B was developed for untested patients.

External validation in different studies is relevant to assess the prediction performance of prediction models[13]. Our aim was to perform a head-to-head comparison between CBCrisk, PredictCBC and the Manchester formula. We hereto used several large population- and hospital-based studies used to develop and cross-validate the PredictCBC models.

**Material and Methods**

External validation of CBCrisk and the Manchester formula was performed in 20 studies: four with individual patient data from the Netherlands (the Amsterdam Breast Cancer Study (ABCS), the Breast Cancer Outcome Study of Mutation carriers (BOSOM), the Erasmus MC Breast Cancer Registry (EMC), the Netherlands Cancer Registry (NCR)); and 16 other studies of the Breast Cancer Association Consortium (BCAC). The latter is an international consortium of 102 studies comprising 182,898 patients (data version: January 2017) with a primary BC diagnosed between 1939 and 2016[14]. In particular, 16 non-familiar BC BCAC studies with at least 10 CBC events were selected including invasive non-metastatic European-descent female patients with first primary invasive BC diagnosed from 1990. Details about studies and patient selection, and data imputation were described previously[5].

The outcome wasin situ or invasive metachronous CBC. Follow-up started 3 months after invasive first primary BC diagnosis, to exclude synchronous CBCs, and ended at date of CBC, distant metastasis (but not at loco-regional relapse), CPM or last date of follow-up (due to death, being lost to follow-up, or end of study), whichever occurred first. In the BCAC, 27,155 patients were recruited more than 3 months after diagnosis of the first primary BC (prevalent cases); for these patients, follow-up started at date of recruitment (left truncation). Distant metastasis and death due to any cause were competing events.

The Manchester formula provides an estimate of a woman’s individual life-time CBC risk. To assess the prediction performance, we translated the life-time CBC risk to 5- and 10-year CBC risks (see **Supplementary Material**). The predictors included in the CBC risk estimation in the Manchester formula, CBCrisk and PredictCBC models are provided in **Table 1**. Predictors that were sporadically missing were multiply imputed as described elsewhere[5].

**Statistical analysis**

Discrimination, the ability of the model to differentiate between patients who experienced CBC and those who did not, was calculated by time-dependent Area-Under-the-Curve (AUCs) based on Inverse Censoring Probability Weighting at 5 and 10 years[15,16]. Values of AUCs close to 1 indicate good discrimination while values close to 0.5 indicate poor discrimination (a coin flip). Calibration is the agreement between observed and predicted risk and is commonly characterized by calibration-in-the-large and slope statistic. Calibration-in-the-large characterizes the overall difference between the observed and predicted risks. It was calculated using the expected/observed (E/O) ratio. An E/O less than 1 indicates that the model systematically underestimates CBC risk, while an E/O above 1 indicates that the model systematically overestimates CBC risk. The expected number of cases was calculated by summing the individual predicted probabilities at 5 and 10 years, based on the patient-specific covariate values[17]. The observed number of cases was estimated by the non-parametric CBC cumulative incidence at 5 and 10 years. The calibration slope was estimated using a Fine and Gray regression model using the linear predictor of the prediction tools. The linear predictor was vs constructed as the sum of the factors included in each model weighted by the corresponding regression coefficients (or parameters), and then computed in the validation dataset exactly as reported for the development set. The calibration slope is determined as the regression coefficient for this linear predictor when fitted as a single covariate in a regression model of disease outcome in the validation dataset. A well-calibrated model should have a calibration slope of 1; slopes < 1 indicate that coefficients were too optimistic for the validation setting[18]. Calibration results were graphically displayed.

Analyses were stratified by geographic groups of studies, since stratification by individual studies would provide too few events in some strata[13,19,5]. To allow for heterogeneity across multiple studies, random-effect meta-analyses were performed. We calculated 95% confidence intervals (CI) and 95% prediction intervals (PI), which indicate the likely range for prediction accuracy of the model in a new dataset, for discrimination and calibration measures. A sensitivity analysis was performed to check the consistency of CBCrisk performance measures when metachronous CBC was defined as an event after 6 instead of 3 months since the first BC diagnosis. More details are provided in the **Supplementary Material**. All analyses were implemented using SAS (SAS Institute Inc., NC, USA) and R software[20].

**Results**

We included 132,756 patients from 20 studies who experienced 4,862 CBC events during a median follow-up of 8.8 years. The main patient and clinical characteristics across studies and geographic areas are shown in **Table 2**.

The AUCs at 5 and 10 years was around 0.6: 0.59 (95% CI: 0.57–0.61; 95% PI: 0.54–0.64) and 0.58 (95% CI: 0.57–0.59; 95% PI: 0.55–0.61) for CBCrisk (**Figure 1**); 0.61 (95% CI: 0.60–0.62; 95% PI: 0.59–0.63) and 0.60 (95% CI: 0.59–0.61; 95% PI: 0.58–0.62) for the Manchester formula (**Figure 2**). The E/O ratio at 5 and 10 years was close to 1 for all models: 0.86 (95% CI: 0.50–1.46; 95% PI: 0.20–3.75) and 0.82 (95% CI: 0.51–1.32; 95% PI: 0.21–3.14) for CBCrisk **(Table 3)**; 1.54 (95% CI: 0.61–3.92; 95% PI: 0.11–20.72, **Table 4**), and 1.53 (95% CI: 0.63–3.73; 95% PI: 0.13–18.52) for the Manchester formula (**Table 4**); 1.26 (95% CI: 0.57–2.77; 95% PI: 0.14–11.34), and 1.28 (95% CI: 0.63–2.58; 95% PI: 0.18–9.18) for PredictCBC-1A (**Table 5**); 1.33 (95% CI: 0.59–2.99, 95% PI: 0.14–12.76), 1.35 (95% CI: 0.65–2.77; 95% PI: 0.19–10.24) for PredictCBC-1B (**Table 5**)[5]. The calibration slope was close to 1 for CBCrisk (1.26, 95% CI: 1.01–1.50 and 95% PI: 1.01–1.50, **Table 3-5**), and PredictCBC-1A and 1B 0.90 (95% CI: 0.79–1.02; 95% PI: 0.73–1.08), and 0.81 (95% CI: 0.63–0.99; 95% PI: 0.50–1.12) (**Table 5**), while prognostic effects were far too large for the Manchester formula (slope: 0.39, 95% CI: 0.34–0.43, 95% PI: 0.34–0.43, **Table 4-5**). Calibration plots of CBCrisk at 5 and 10 years are shown in **Supplementary Figure 1** and **Supplementary Figure 2.** As reported previously[5], the AUCs at 5 and 10 years for PredictCBC-1A were 0.63 (95% CI: 0.58–0.67, 95% PI: 0.52–0.74), and 0.63 (95% CI: 0.59–0.66, 95% PI: 0.53–0.72), respectively; for PredictCBC-1B 0.59 (CI: 0.54–0.63, 95% PI: 0.46–0.71, **Table 5**), and 0.59 (95% CI: 0.56–0.62, 95% PI: 0.52–0.66, **Table 5**), respectively.

Sensitivity analysis showed that the performance measures of CBCrisk did not change when metachronous CBC was defined after 6 months since first BC diagnosis (see **Supplementary Materials, Supplementary Table 1-2 and Supplementary Figure 3**).

**Discussion**

Accurate CBC risk predictions are essential in clinical decision making around CPM or tailored surveillance among patients with first primary BC. In particular, overestimation of risk can lead to recommending CPM among BC patients with low risks. Underestimation can lead to suboptimal surveillance or hesitance about recommending CPM for patients with substantial risk. Using individual patient data from multiple studies with long follow-up, we externally evaluated the prediction performance accuracy of CBCrisk, a tool developed and validated to provide individualized CBC risk prediction, and the Manchester formula, a heuristically derived calculation of CBC lifetime risk[6,8,7,9]. In addition, the availability of different European-descendent studies allowed heterogeneity in the performance by geographic area to be assessed.

CBCrisk under-predicted the risk of CBC and had moderate discrimination ability with considerable heterogeneity between studies. The Manchester formula was empirically derived from a systematic review, and its discrimination accuracy was higher than CBCrisk. This may be explained by the inclusion of *BRCA1/2* mutation carrier information, an important determinant of CBC risk[21]. With the same large individual patient data sets, PredictCBC models had been developed and validated[5]. In particular, PredictCBC version 1A includes information of *BRCA1/2* mutation carriers and extensive information about the primary BC including treatments. The discrimination of all three prediction models was moderate, with AUC values around 0.6.

CBCrisk was previously externally validated using two independent clinical studies from Johns Hopkins University (JH) and MD Anderson Cancer Center (MDA) in the US[9]. Discrimination ability was 0.61 and 0.65 at 3 years, and 0.62 and 0.61 at 5 years for JH and MDA, respectively. The risk of CBC was overestimated in JH with E/O ratios of 2.02 and 1.56 at 3 and 5 years, while underestimated in MDA with E/O ratios of 0.61 and 0.62, respectively.

The considerable heterogeneity in all CBC risk calculators, especially in the CBCrisk and the Manchester formula, reflects the different CBC incidences in every study[13]. Another potential source of heterogeneity is the carrier frequency of germline mutations associated with CBC that may vary among studies, especially in the CBC calculators not including information of *BRCA1/2* mutation as CBCrisk and the PredictCBC-1B[22]. In addition, heterogeneity may be due to the different proportions of the use of (neo)adjuvant systemic therapies explained by the different distribution of tumor subtypes among studies[4]. Besides, inter-observer variation in pathological examination of BC among studies may deceive adjuvant systemic therapy advice and, consequently, prediction of CBC risk[23]. Variation in prediction performance and limited generalizability of CBC risk calculators can also be partially explained by differences in how predictors are measured among studies[24,25]. For example, lack of family history knowledge may lead to uncertainty in risk prediction and varies according to demographics of the patients[26]. In particular, if in some studies BC patients misreported information about family history, the CBC risk would be over(under)estimated causing inappropriate decision-making regarding CPM or tailored surveillance. Some limitations of our study must be recognized. Firstly, our dataset, while large, had missing data for three covariates that were used in the CBCrisk model: breast density, age at first birth, and high risk preneoplasia. The authors of CBCrisk estimated the relative risks for patients with the unknown characteristics, but the use of the missing indicator variable is suboptimal compared to having the prognostic information available. It may lead to over or under-estimation of absolute CBC risk[27]. For this reason, we suggest that it is preferable to use multiple imputation of missing data, as is done in the PredictCBC models[28,29]. In addition, investigation of the potential source of model misspecification due to possible different definitions or measurement error was not possible[30-32].

In conclusion, current statistical risk prediction models and heuristic formulas provided moderate CBC individualized prediction performance. Careful re-calibration is required before considering these models for clinical decision making. A more direct comparison between the current CBC risk prediction models using a large external dataset with complete information on all factors included in all CBC prediction models would be ideal, but is currently unavailable. There is an ongoing debate about improvements of clinical prediction performance using machine learning approaches compared to standard regression approaches for risk prediction[33,34]. However, irrespective of the methodology, better predictors are needed to predict CBC more accurately. Deeper biological insights and potential inclusion of other genetic markers such as *CHEK2* c.1100del mutation status and polygenic risk scores based on common genetic variants may improve CBC risk prediction, although rare mutations are unlikely to contribute substantially to CBC risk in the general population[35,36]. Life-style factors such as body mass index, alcohol consumption, and smoking also may help to better stratify high and low CBC risk patients even though these factors are difficult to measure accurately. Moreover, breast density may be important. More detailed information about adjuvant systemic therapies may better identify patients with low and high CBC risk since chemotherapy and especially endocrine therapy reduce CBC risk and influent CBC receptor subtype[4]. After extension and further external validation of prediction models for CBC risk, investigation of their potential clinical utility is an important future step.

**Acknowledgements**

We thank all individuals who took part in these studies and all researchers, clinicians, technicians and administrative staff who have enabled this work to be carried out. ABCFS thank Maggie Angelakos, Judi Maskiell, Gillian Dite. ABCS and BOSOM thanks all the collaborating hospitals and pathology departments and many individual that made this study possible, specifically, we wish to acknowledge: Annegien Broeks, Sten Cornelissen, Frans Hogervorst, Laura van ‘t Veer, Floor van Leeuwen, Emiel Rutgers. EMC thanks J.C. Blom-Leenheer, P.J. Bos,C.M.G. Crepin and M. van Vliet for data management. CGPS thanks staff and participants of the Copenhagen General Population Study. For the excellent technical assistance: Dorthe Uldall Andersen, Maria Birna Arnadottir, Anne Bank, Dorthe Kjeldgård Hansen. HEBCS thanks Taru A. Muranen, Kristiina Aittomäki, Karl von Smitten, Irja Erkkilä. KARMA thanks the Swedish Medical Research Counsel. LMBC thanks Gilian Peuteman, Thomas Van Brussel, EvyVanderheyden and Kathleen Corthouts. MARIE thanks Petra Seibold, Dieter Flesch-Janys, Judith Heinz, Nadia Obi, Alina Vrieling, Sabine Behrens, Ursula Eilber, Muhabbet Celik, Til Olchers and Stefan Nickels. ORIGO thanks E. Krol-Warmerdam, and J. Blom for patient accrual, administering questionnaires, and managing clinical information. The authors thank the registration team of the Netherlands Comprehensive Cancer Organisation (IKNL) for the collection of data for the Netherlands Cancer Registry as well as IKNL staff for scientific advice. PBCS thanks Louise Brinton, Mark Sherman, Neonila Szeszenia-Dabrowska, Beata Peplonska, Witold Zatonski, Pei Chao, Michael Stagner. The ethical approval for the POSH study is MREC /00/6/69, UKCRN ID: 1137. We thank the SEARCH team.

**Funding**

This work is supported by the Alpe d’HuZes/Dutch Cancer Society (KWF Kankerbestrijding) project 6253.

BCAC is funded by Cancer Research UK [C1287/A16563, C1287/A10118], the European Union's Horizon 2020 Research and Innovation Programme (grant numbers 634935 and 633784 for BRIDGES and B-CAST respectively), and by the European Community´s Seventh Framework Programme under grant agreement number 223175 (grant number HEALTH-F2-2009-223175) (COGS). The EU Horizon 2020 Research and Innovation Programme funding source had no role in study design, data collection, data analysis, data interpretation or writing of the report.

The Australian Breast Cancer Family Study (ABCFS) was supported by grant UM1 CA164920 from the National Cancer Institute (USA). The ABCFS was also supported by the National Health and Medical Research Council of Australia, the New South Wales Cancer Council, the Victorian Health Promotion Foundation (Australia) and the Victorian Breast Cancer Research Consortium. J.L.H. is a National Health and Medical Research Council (NHMRC) Senior Principal Research Fellow. M.C.S. is a NHMRC Senior Research Fellow. The ABCS study was supported by the Dutch Cancer Society [grants NKI 2007-3839; 2009 4363]. The work of the BBCC was partly funded by ELAN-Fond of the University Hospital of Erlangen. BOSOM was supported by the Dutch Cancer Society grant numbers DCS-NKI 2001-2423, DCS-NKI 2007-3839, and DCSNKI 2009-4363; the Cancer Genomics Initiative; and notary office Spier & Hazenberg for the coding procedure. The EMC was supported by grants from Alpe d’HuZes/Dutch Cancer Society NKI2013-6253 and from Pink Ribbon 2012.WO39.C143. The HEBCS was financially supported by the Helsinki University Hospital Research Fund, the Finnish Cancer Society, and the Sigrid Juselius Foundation.

Financial support for KARBAC was provided through the regional agreement on medical training and clinical research (ALF) between Stockholm County Council and Karolinska Institutet, the Swedish Cancer Society, The Gustav V Jubilee foundation and Bert von Kantzows foundation. The KARMA study was supported by Märit and Hans Rausings Initiative Against Breast Cancer. LMBC is supported by the 'Stichting tegen Kanker'. The MARIE study was supported by the Deutsche Krebshilfe e.V. [70-2892-BR I, 106332, 108253, 108419, 110826, 110828], the Hamburg Cancer Society, the German Cancer Research Center (DKFZ) and the Federal Ministry of Education and Research (BMBF) Germany [01KH0402]. MEC was support by NIH grants CA63464, CA54281, CA098758, CA132839 and CA164973. The ORIGO study was supported by the Dutch Cancer Society (RUL 1997-1505) and the Biobanking and Biomolecular Resources Research Infrastructure (BBMRI-NL CP16). The PBCS was funded by Intramural Research Funds of the National Cancer Institute, Department of Health and Human Services, USA. Genotyping for PLCO was supported by the Intramural Research Program of the National Institutes of Health, NCI, Division of Cancer Epidemiology and Genetics. The POSH study is funded by Cancer Research UK (grants C1275/A11699, C1275/C22524, C1275/A19187, C1275/A15956 and Breast Cancer Campaign 2010PR62, 2013PR044. PROCAS is funded from NIHR grant PGfAR 0707-10031. SEARCH is funded by Cancer Research UK [C490/A10124, C490/A16561] and supported by the UK National Institute for Health Research Biomedical Research Centre at the University of Cambridge. SKKDKFZS is supported by the DKFZ. The SZBCS (Szczecin Breast Cancer Study) was supported by Grant PBZ\_KBN\_122/P05/2004 and The National Centre for Research and Development (NCBR) within the framework of the international ERA-NET TRANSAN JTC 2012 application no. Cancer 12-054 (Contract No. ERA-NET-TRANSCAN / 07/2014).

**Compliance with ethical standards**

**Funding:** This work is supported by the Alpe d’HuZes/Dutch Cancer Society (KWF Kankerbestrijding) project 6253.

**Conflict of interest:** Author DG, MH, EW, MAA, DA, JCB, CB, SEB, MKB, JCC, KC, PD, AMD, DFE, JF, HF, MGC, LH, CAH, PH, UH, JLH, AJ, AJ2, AJ3, RK, LBK, IK, DL, LLN, AL, JL, MM, LM, HN, HSAO, SP, PDPP, MS, SS, VTHBMS, MCS, WJT, RAEMT, AJvdB, CHMvD, FEvL, CvO, LvV, QW, CW, PJW, MJH declares that he has no conflict of interest. Author DMM declares that she receives a lecture fee from Pierre Fabre and personal fees for consultancy from Astra Zeneca. Author PAF reports grants from Novartis, grants from Biontech, personal fees from Novartis, personal fees from Roche, personal fees from Pfizer, personal fees from Celgene, personal fees from Daiichi-Sankyo, personal fees from TEVA, personal fees from Astra Zeneca, personal fees from Merck Sharp & Dohme, personal fees from Myelo Therapeutics, personal fees from Macrogenics, personal fees from Eisai, personal fees from Puma, grants from Cepheid.

**Ethical approval**: all procedures performed in studies involving human participants were in accordance with the ethical standards of international, national, and institutional research committees and with the 1964 Helsinki declaration and its later amendments or comparable ethical standards.

**Informed consent:** informed consent was obtained from all individual participants included in the study.

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| **Table 1:** Predictors included in current contralateral breast cancer risk prediction tools | | | | |
| List of predictors | CBCrisk§ | Manchester formula† | PredictCBC version 1A‡ | PredictCBC version 1B‡ |
| Age at diagnosis | ✔ | ✔ | ✔ | ✔ |
| Age at first birth | ✔ |  |  |  |
| First-degree family history | ✔ | ✔ | ✔ | ✔ |
| *BRCA1/2* germline mutation |  | ✔ | ✔ |  |
| First breast cancer behavior type\* | ✔ | ✔ |  |  |
| Lymph node status |  |  | ✔ | ✔ |
| Breast density | ✔ |  |  |  |
| Tumor size |  |  | ✔ | ✔ |
| Morphology |  |  | ✔ | ✔ |
| Tumor grade |  |  | ✔ | ✔ |
| High risk pre-neoplasia | ✔ |  |  |  |
| ER status | ✔ | ✔ | ✔ | ✔ |
| HER2 status |  |  | ✔ | ✔ |
| Chemotherapy |  |  | ✔ | ✔ |
| Endocrine therapy | ✔ |  | ✔ | ✔ |
| Radiation to the breast |  |  | ✔ | ✔ |
| Trastuzumab |  |  | ✔ | ✔ |
| Oophorectomy under 40 years |  | ✔ |  |  |
| Abbreviation: ER: estrogen receptor status; HER2: human epidermal growth factor receptor 2.  \* Contralateral breast cancer risk was calculated including women diagnosed with ductal carcinoma in situ;  §Chowdhury M, Euhus D, Onega T, Biswas S, Choudhary PK (2017) A model for individualized risk prediction of contralateral breast cancer. Breast Cancer Res Treat 161 (1):153-160.  †Basu NN, Ross GL, Evans DG, Barr L (2015) The Manchester guidelines for contralateral risk-reducing mastectomy. World J Surg Oncol 13:237  ‡ Giardiello D, Steyerberg EW, Hauptmann M, Adank MA, Akdeniz D, Blomqvist C, Bojesen SE, Bolla MK, Brinkhuis M, Chang-Claude J, Czene K, Devilee P, Dunning AM, Easton DF, Eccles DM, Fasching PA, Figueroa J, Flyger H, Garcia-Closas M, Haeberle L, Haiman CA, Hall P, Hamann U, Hopper JL, Jager A, Jakubowska A, Jung A, Keeman R, Kramer I, Lambrechts D, Le Marchand L, Lindblom A, Lubinski J, Manoochehri M, Mariani L, Nevanlinna H, Oldenburg HSA, Pelders S, Pharoah PDP, Shah M, Siesling S, Smit V, Southey MC, Tapper WJ, Tollenaar R, van den Broek AJ, van Deurzen CHM, van Leeuwen FE, van Ongeval C, Van't Veer LJ, Wang Q, Wendt C, Westenend PJ, Hooning MJ, Schmidt MK (2019) Prediction and clinical utility of a contralateral breast cancer risk model. Breast Cancer Res 21 (1):144. doi:10.1186/s13058-019-1221-1 | | | | |

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| --- | --- | --- | --- | --- | --- | --- | --- |
| **Table 2:** Description of main patient and clinical factors used for evaluation of the models and formula ‡ | | | | | | | |
| Study\* / Geographic area | Europe -  other§ | Europe -  Scandinavia | Europe -  United Kingdom | Netherlands - BOSOM | Netherlands -  EMC | Netherlands -  NCR | United States and  Australia |
| N | 15,183 | 12,928 | 11,921 | 3,760 | 3,390 | 83,138 | 2,436 |
| Age at first diagnosis, *years* (%) |  |  |  |  |  |  |  |
| <30 | 152 (1.0) | 46 (0.4) | 156 (1.3) | 108 (2.9) | 46 (1.4) | 388 (0.5) | 41 (1.7) |
| 30-39 | 1,252 (8.2) | 489 (3.8) | 1,811 (15.2) | 842 (22.4) | 374 (11.0) | 4,241 (5.1) | 494 (20.3) |
| 40+ | 13,779 (90.8) | 12,393 (95.9) | 9,954 (83.5) | 2,810 (74.7) | 2,970 (87.6) | 78,509 (94.4) | 1,901 (78.0) |
| Age at first birth = unknown (%) | 15,183 (100.0) | 12,928 (100.0) | 11,921 (100.0) | 3,760 (100.0) | 3,390 (100.0) | 83138 (100.0) | 2,436 (100.0) |
| Family history (%) |  |  |  |  |  |  |  |
| Yes | 2,123 (14.0) | 818 (6.3) | 1,371 (11.5) | 737 (19.6) | 591 (17.4) | 0 (0.0) | 319 (13.1) |
| No | 8,057 (53.1) | 3,158 (24.4) | 8,210 (68.9) | 1,177 (31.3) | 2,482 (73.2) | 0 (0.0) | 1,498 (61.5) |
| Unknown | 5,003 (33.0) | 8,952 (69.2) | 2,340 (19.6) | 1,846 (49.1) | 317 (9.4) | 83,138 (100.0) | 619 (25.4) |
| First BC type = Pure invasive (%) | 15,183 (100.0) | 12,928 (100.0) | 11,921 (100.0) | 3,760 (100.0) | 3,390 (100.0) | 83,138 (100.0) | 2,436 (100.0) |
| Breast density = unknown (%) | 15,183 (100.0) | 12,928 (100.0) | 11,921 (100.0) | 3,760 (100.0) | 3,390 (100.0) | 83,138 (100.0) | 2,436 (100.0) |
| ER status (%) |  |  |  |  |  |  |  |
| Negative | 3,387 (22.3) | 1,746 (13.5) | 1,718 (14.4) | 896 (23.8) | 842 (24.8) | 14,591 (17.6) | 445 (18.3) |
| Positive | 10,071 (66.3) | 9,401 (72.7) | 7,175 (60.2) | 2,024 (53.8) | 2,427 (71.6) | 64,790 (77.9) | 1,572 (64.5) |
| Unknown | 1,725 (11.4) | 1,781 (13.8) | 3,028 (25.4) | 840 (22.3) | 121 (3.6) | 3,757 (4.5) | 419 (17.2) |
| High risk pre-neoplasia = unknown (%) | 15,183 (100.0) | 12,928 (100.0) | 11,921 (100.0) | 3,760 (100.0) | 3,390 (100.0) | 83,138 (100.0) | 2,436 (100.0) |
| Anti-estrogen therapy (%) |  |  |  |  |  |  |  |
| Yes | 7,868 (51.8) | 6,434 (49.8) | 8,712 (73.1) | 809 (21.5) | 1,559 (46.0) | 40,214 (48.4) | 363 (14.9) |
| No | 4,570 (30.1) | 1,947 (15.1) | 2,046 (17.2) | 2,739 (72.8) | 1,821 (53.7) | 42,924 (51.6) | 8 (0.3) |
| Unknown | 2,745 (18.1) | 4,547 (35.2) | 1,163 (9.8) | 212 (5.6) | 10 (0.3) | 0 (0.0) | 2,065 (84.8) |
| CBC cumulative incidence (%) |  |  |  |  |  |  |  |
| 3-year (95% CI) | 1.0 (0.8 - 1.2) | 0.7 (0.5 - 0.9) | 0.5 (0.3 - 0.7) | 1.7 (1.3 - 2.1) | 1.7 (1.2 - 2.1) | 1.3 (1.2 - 1.4) | 1.8 (0.8 - 2.8) |
| 5-year (95% CI) | 1.6 (1.4 - 1.9) | 1.0 (0.8 - 1.3) | 1.0 (0.8 - 1.3) | 3.0 (2.5 - 3.6) | 2.6 (2.1 - 3.2) | 2.4 (2.3 - 2.5) | 2.8 (1.7 - 3.8) |
| 10-year (95% CI) | 3.5 (3.1 - 3.9) | 2.1 (1.7 - 2.4) | 1.3 (1.0 - 1.5) | 5.5 (4.7 - 6.2) | 5.7 (4.9 - 6.6) | 4.6 (4.5 - 4.8) | 4.1 (3.0 - 5.3) |
| ‡More details about the main patient and clinical characteristics by study are available in the supplementary information of [5]  Abbreviations: \* The studies denoted with Europe and United States and Australia are part of the Breast Cancer Association Consortium  § Europe - other geographic area included studies from Belgium (1), Germany (2), Netherlands (2) and Poland (2). BOSOM: Breast Cancer Outcome Study of Mutation carriers; EMC: Erasmus Medical Center; NCR: Netherlands Cancer Registry BC: breast cancer; ER: estrogen receptor; CBC: contralateral breast cancer; CI: confidence interval; | | | | | | | |

**Table 3:** Calibration performance of the CBCrisk model§

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| Validation dataset | E/O ratio at 5 years (95% CI) | | E/O ratio at 10 years (95% CI) | | Calibration slope (95% CI) | |
| Europe - Other | 0.87 (076 - 0.98) | | 0.75 (0.68 - 0.81) | | 1.11 ( 0.40 - 1.83) | |
| Europe - Scandinavia | 1.59 (1.28 - 1.91) | | 1.23 (1.08 - 1.38) | | 0.86 ( 0.16 - 1.57) | |
| Europe - UK | 1.35 (1.38 - 2.17) | | 1.82 (1.53 - 2.11) | | 0.85 (-0.03 - 1.73) | |
| Netherlands - BOSOM | 0.45 (0.37 - 0.53) | | 0.50 (0.43 - 0.57) | | 1.34 ( 0.76 - 1.93) | |
| Netherlands - EMC | 0.48 (0.38 - 0.57) | | 0.43 (0.37 - 0.50) | | 1.19 ( 0.65 - 1.73) | |
| Netherlands - NCR | 0.57 (0.54 - 0.59) | | 0.54 (0.52 - 0.56) | | 1.40 ( 1.11 - 1.68) | |
| US and Australia | 0.43 (0.33 - 0.54) | | 0.56 (0.45 - 0.67) | | 1.13 ( 0.25 - 2.00) | |
|  |  |  |  |  |  |  |
| Meta-analysis | 0.86 (0.50 - 1.46) | | 0.82 (0.51 - 1.32) | | 1.26 ( 1.01 – 1.50) | |
| 95% PI | 0.20 - 3.75 | | 0.21 - 3.14 | | 1.01 - 1.50 | |
| Abbreviations: E/O: expected-observed; CI: confidence interval; UK: United Kingdom; BOSOM: Breast Cancer Outcome Study of Mutation carriers; EMC: Erasmus Medical; Center NCR: Netherlands Cancer Registry; PI: prediction interval  §Chowdhury M, Euhus D, Onega T, Biswas S, Choudhary PK (2017) A model for individualized risk prediction of contralateral breast cancer. Breast Cancer Res Treat 161 (1):153-160. | | | | | | |
|

**Table 4:** Calibration performance of the Manchester formula§

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| Validation dataset | E/O ratio at 5 years (95% CI) | | E/O ratio at 10 years (95% CI) | | Calibration slope (95% CI) | |
| Europe - Other | 1.64 (1.44 - 1.85) | | 1.46 (1.34 - 1.58) | | 0.40 (0.29 - 0.50) | |
| Europe - Scandinavia | 2.61 (2.09 - 3.12) | | 2.11 (1.85 - 2.37) | | 0.35 (0.13 - 0.57) | |
| Europe - UK | 3.34 (2.60 - 4.08) | | 3.49 (2.93 - 4.05) | | 0.42 (0.23 - 0.61) | |
| Netherlands - BOSOM | 0.81 (0.66 - 0.96) | | 0.92 (0.79 - 1.05) | | 0.45 (0.33 - 0.56) | |
| Netherlands - EMC | 0.94 (0.75 - 1.14) | | 0.87 (0.75 - 1.00) | | 0.35 (0.21 - 0.49) | |
| Netherlands - NCR | 1.00 (0.95 - 1.04) | | 1.01 (0.98 - 1.05) | | 0.37 (0.33 - 0.42) | |
| US and Australia | 0.77 (0.58 - 0.96) | | 1.02 (0.82 - 1.23) | | 0.51 (0.33 - 0.68) | |
|  |  |  |  |  |  |  |
| Meta-analysis | 1.54 (0.61 - 3.92) | | 1.53 (0.63 - 3.73) | | 0.39 (0.34 - 0.43) | |
| 95% PI | 0.11 - 20.72 | | 0.13 - 18.52 | | 0.34 - 0.43 | |
| Abbreviations: E/O: expected-observed; CI: confidence interval; UK: United Kingdom; BOSOM: Breast Cancer Outcome Study of Mutation carriers; EMC: Erasmus Medical; Center NCR: Netherlands Cancer Registry; PI: prediction interval  §Basu NN, Ross GL, Evans DG, Barr L (2015) The Manchester guidelines for contralateral risk-reducing mastectomy. World J Surg Oncol 13:237 | | | | | | |

**Table 5:** Summary of prediction performance of CBCrisk, Manchester formula, and PredictCBC version 1A and version 1B with the corresponding 95% prediction intervals (PI).

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
|  |  |  |  |  |
| Characteristics | CBCrisk§ | Manchester  formula† | PredictCBC  version 1A‡\* | PredictCBC  version 1B‡\* |
| Discrimination |  |  |  |  |
| AUC at 5 years (95% PI) | 0.59 (0.54 - 0.64) | 0.61 (0.59 - 0.63) | 0.63 (0.52 - 0.74) | 0.59 (0.46 - 0.71) |
| AUC at 10 years (95% PI) | 0.58 (0.55 - 0.61) | 0.60 (0.58 - 0.62) | 0.63 (0.53 - 0.72) | 0.59 (0.52 - 0.66) |
| Calibration |  |  |  |  |
| E/O ratio at 5 years (95% PI) | 0.86 (0.20 - 3.75) | 1.54 (0.11 - 20.72) | 1.26 (0.14 - 11.34) | 1.33 (0.14 - 12.76) |
| E/O ratio at 10 years (95% PI) | 0.82 (0.21 - 3.14) | 1.53 (0.13 - 18.52) | 1.28 (0.18 - 9.18) | 1.35 (0.19 - 10.24) |
| Slope (95% PI) | 1.26 (1.01 - 1.50) | 0.39 (0.34 - 0.43) | 0.90 (0.73 - 1.08) | 0.81 (0.50 - 1.12) |
| Abbreviations: AUC: Area under the curve; PI: prediction interval  §Chowdhury M, Euhus D, Onega T, Biswas S, Choudhary PK (2017) A model for individualized risk prediction of contralateral breast cancer. Breast Cancer Res Treat 161 (1):153-160.  †Basu NN, Ross GL, Evans DG, Barr L (2015) The Manchester guidelines for contralateral risk-reducing mastectomy. World J Surg Oncol 13:237  ‡ Giardiello D, Steyerberg E, Hauptmann M, et al. (2019) Prediction and clinical utility of a contralateral breast cancer risk model. Breast Cancer Res. doi:10.1186/s13058-019-1221-1, Figure 1 and Figure S5  \*version 1A includes *BRCA* mutation status as a variable while 1B does not. | | | | |