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Guidelines Breast
Version 2023.1E

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Diagnosis and Treatment of Patients with early and advanced Breast Cancer

Breast Cancer Risk, Genetics and Prevention

Breast Cancer Risk and Prevention

- **Versions 2003–2022:**

Albert / Bischoff / Blohmer / Dall / Ditsch / Fasching / Fehm / Gerber / Kiechle / Maass /
Müller-Schimpfle / Mundhenke / Park-Simon / Rhiem / Rody / Schmidt / Schmutzler / Stickeler
/ Thomssen / Witzel

- **Version 2023:**
Schütz / Thomssen



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Principles of Prevention

- **Women at increased risk for breast cancer are not considered *patients* but *healthy women* or *counselees*.**
- **A comprehensive informed consent taking into consideration all potential side effects and risks is warranted prior to offering preventive measures.**
- **Highest priority: „First, do no harm!“**

(Primum nil nocere)

Indication for Genetic Testing of *BRCA1/2* Genes and Possibly Further Risk Genes?

(Part 1 of 2 – testing according to family history)

Oxford LoE: 2b GR: B AGO: ++

Families with (each from one family branch) at least*

- three women with breast cancer independent of age
- two women with breast cancer, one diagnosed before the 51st birthday
- one woman affected by breast and one by ovarian cancer or
- one woman affected by breast and ovarian cancer or
- two women affected by ovarian cancer or
- one woman affected by bilateral breast cancer, first before 51st birthday
- one woman affected by breast cancer before the 36th birthday or
- one man affected by breast cancer

* Inclusion criteria of the German Consortium of Hereditary Breast and Ovarian Cancer (GCHBOC) based on a *BRCA1/2* mutation prevalence $\geq 10\%$ tested in 21,401 families. All mutation carriers should be registered in scientific databases, to validate the inclusion and exclusion criteria (including population-based studies).

1. Beitsch PD, Whitworth PW, Hughes K. Underdiagnosis of Hereditary Breast Cancer: Are Genetic Testing Guidelines a Tool or an Obstacle? *Journal of Clinical Oncology* 2019 37:6, 453-460
2. Couch FJ, Hart SN, Sharma P, et al. Inherited mutations in 17 breast cancer susceptibility genes among a large triple-negative breast cancer cohort unselected for family history of breast cancer. *J Clin Oncol.* 2015;33(4):304-11.
3. Meindl A, German Consortium for Hereditary B, Ovarian C. Comprehensive analysis of 989 patients with breast or ovarian cancer provides *BRCA1* and *BRCA2* mutation profiles and frequencies for the German population. *Int J Cancer.* 2002;97(4):472-80.
4. Kast K, Rhiem K, Wappenschmidt B, et al., Prevalence of *BRCA1/2* germline mutations in 21.401 families with breast and ovarian cancer. *J Med Genet* 2016;53:465-71.
5. Manchanda R, Gaba F. Population Based Testing for Primary Prevention: A Systematic Review. *Cancers (Basel).* 2018 Nov 5;10(11).
6. Rolfes M, Borde J, Möllenhoff K et al, Prevalence of Cancer Predisposition Germline Variants in Male Breast Cancer Patients: Results of the German Consortium for Hereditary Breast and Ovarian Cancer, *Cancers*, 2022, 14(13): 3292

Indication for Genetic Testing of *BRCA1/2* Genes and Possibly Further Risk Genes?

(Part 2 of 2 - testing according to disease)

Oxford LoE: 2b GR: B AGO: ++


■ Other recommended criteria:

- own disease of triple negative breast cancer diagnosed before 60th birthday
- own disease of ovarian cancer before 80th birthday
- if therapeutically relevant (e.g. PARPi; *BRCA1* and *BRCA2* only; possibly *PALB2*)

1. Couch FJ, Hart SN, Sharma P, et al. Inherited mutations in 17 breast cancer susceptibility genes among a large triple-negative breast cancer cohort unselected for family history of breast cancer. *J Clin Oncol*. 2015;33(4):304-11.
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5. Litton JK, Hurvitz SA, Mina LA, et al. Talazoparib versus chemotherapy in patients with germline *BRCA1/2*-mutated HER2-negative advanced breast cancer: final overall survival results from the EMBRACA trial. *Ann Oncol*. 2020 Nov;31(11):1526-1535. doi: 10.1016/j.annonc.2020.08.2098. Epub 2020 Aug 20. PMID: 32828825.
6. Manchanda R, Gaba F. Population Based Testing for Primary Prevention: A Systematic Review. *Cancers (Basel)*. 2018 Nov 5;10(11).
7. Meindl A, German Consortium for Hereditary B, Ovarian C. Comprehensive analysis of 989 patients with breast or ovarian cancer

provides BRCA1 and BRCA2 mutation profiles and frequencies for the German population. *Int J Cancer*. 2002;97(4):472-80.

8. Robson M, Im SA, Senkus E, et al. Olaparib for Metastatic Breast Cancer in Patients with a Germline BRCA Mutation *N Engl J Med* 2017;377:523-533



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Checklist for Recording a Possible Hereditary Burden of Breast and / or Ovarian Cancer

DKG
krebsgesellschaft
Zertifizierung

Online checklist for familial breast and
ovarian cancer:



Source: Deutsche Krebsgesellschaft e.V.

Name Patientin/Patient: _____ Geburtsdatum: _____

A. Patientin und deren Geschwister / Kinder

Auftreten bei Patientin/Patient	Anzahl	Gewichtung	Ergebnis
eines Mammakarzinoms bei der Patientin vor dem 36. Geburtstag	3	0	
eines bilateralen Mammakarzinoms bei der Patientin vor dem 50. Geburtstag	3	0	
eines unilateralen Mammakarzinoms bei der Patientin vor dem 50. Geburtstag	2	0	
eines bilateralen Mammakarzinoms bei der Patientin, das erst vor dem 50. Geburtstag	3	0	
eines un- oder bilateralen Mammakarzinoms bei der Patientin nach dem 50. Geburtstag	1	0	
eines un- oder bilateralen Mammakarzinoms bei der Patientin (Häufigkeit)	2	0	
eines Ovarialkarzinoms bei der Patientin vor dem 50. Geburtstag	3	0	
eines Ovarialkarzinoms bei der Patientin nach dem 50. Geburtstag	2	0	
Auftreten bei Kindern, Geschwister und deren Kindern			
eines Mammakarzinoms bei Schwestern/Töchtern/Männern vor dem 36. Geburtstag	3	0	
eines unilateralen Mammakarzinoms bei Schwestern/Töchtern/Männern vor dem 50. Geburtstag	2	0	
eines bilateralen Mammakarzinoms bei Schwestern/Töchtern/Männern, das erst vor dem 50. Geburtstag	3	0	
eines un- oder bilateralen Mammakarzinoms bei Schwestern/Töchtern/Männern nach dem 50. Geburtstag	1	0	
eines un- oder bilateralen Mammakarzinoms bei Brüdern/Geschwister	2	0	
eines Ovarialkarzinoms bei Schwestern/Töchtern/Männern	2	0	
B. Mütterliche Linie (incl. Mutter)			
Auftreten	Anzahl	Gewichtung	Ergebnis
eines Mammakarzinoms bei einer Angehörigen vor dem 36. Geburtstag	3	0	
eines unilateralen Mammakarzinoms bei einer Angehörigen vor dem 50. Geburtstag	2	0	
eines bilateralen Mammakarzinoms bei einer Angehörigen, das erst vor dem 50. Geburtstag	3	0	
eines un- oder bilateralen Mammakarzinoms bei einer Angehörigen nach dem 50. Geburtstag	1	0	
eines Mammakarzinoms bei einer Angehörigen (Häufigkeit)	2	0	
eines Ovarialkarzinoms bei einer Angehörigen	2	0	
Summe mütterliche Linie			B
C. Väterliche Linie (incl. Vater)			
Auftreten	Anzahl	Gewichtung	Ergebnis
eines Mammakarzinoms bei einer Angehörigen vor dem 36. Geburtstag	3	0	
eines unilateralen Mammakarzinoms bei einer Angehörigen vor dem 50. Geburtstag	2	0	
eines bilateralen Mammakarzinoms bei einer Angehörigen, das erst vor dem 50. Geburtstag	3	0	
eines un- oder bilateralen Mammakarzinoms bei einer Angehörigen nach dem 50. Geburtstag	1	0	
eines Mammakarzinoms bei einer Angehörigen (Häufigkeit)	2	0	
eines Ovarialkarzinoms bei einer Angehörigen	2	0	
Summe väterliche Linie			C
D. Der höhere Wert aus B und C			D
E. Summe aus A und D = Risiko-Score			A+D

Hier ist das Online Tool zur Checkliste „Familiärer Brust- und Eierstockkrebs“ hinterlegt:
https://www.krebsgesellschaft.de/zertdokumente.html?file=files/dkg/deutsche-krebsgesellschaft/content/pdf/Zertifizierung/Checklisten-und-Algorithmen/checkliste_erbliche_belastung_brust_gyn-220118.xlsx&cid=98969

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1. Kast K, Rhiem K, Wappenschmidt B, et al., Prevalence of BRCA1/2 germline mutations in 21.401 families with breast and ovarian cancer. J Med Genet 2016;53:465-71.
2. Rhiem K, Bücker-Nott HJ, Hellmich M, et al. Benchmarking of a checklist for the identification of familial risk for breast and ovarian cancers in a prospective cohort. Breast J. 2019;25(3):455–460. doi:10.1111/tbj.13257

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1. Kast K, Rhiem K, Wappenschmidt B, et al., Prevalence of BRCA1/2 germline mutations in 21.401 families with breast and ovarian cancer. J Med Genet 2016;53:465-71.
2. Rhiem K, Bücker-Nott HJ, Hellmich M, et al. Benchmarking of a checklist for the identification of familial risk for breast and ovarian cancers in a prospective cohort. Breast J. 2019;25(3):455–460. doi:10.1111/tbj.13257

Risk estimation for syndrome-associated breast cancer (non-BRCA)

	Oxford		
	LoE	GR	AGO
History and family history over at least three generation (including age of first disease)	2b	B	++
<ul style="list-style-type: none"> Characteristic disease <ul style="list-style-type: none"> Breast and ovarian cancer Further disease <ul style="list-style-type: none"> Pancreatic, thyroid, colorectal, stomache, hepatobiliar, urogenital, lung cancer, melanoma, osteosarcoma, leukemia, lymphoma Kidney cancer Testinal cancer Endometrial cancer Prostate cancer 			

- Antoniou AC, Casadei S, Heikkinen T, et al. Breast-cancer risk in families with mutations in PALB2. N Engl J Med. 2014;371(6):497-506.
- Benusiglio PR, Malka D, Rouleau E, et al. CDH1 germline mutations and the hereditary diffuse gastric and lobular breast cancer syndrome: a multicentre study. J Med Genet. 2013;50(7):486-9
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- di Masi A, Antoccia A. NBS1 Heterozygosity and Cancer Risk. Curr Genomics. 2008;9(4):275-81.
- Gao P, Ma N, Li M, et al. Functional variants in NBS1 and cancer risk: evidence from a meta-analysis of 60 publications with 111 individual studies. Mutagenesis. 2013;28(6):683-97.
- Goldgar DE, Healey S, Dowty JG, et al. Rare variants in the ATM gene and risk of breast cancer. Breast Cancer Res. 2011;13(4):R73.
- Engel C, Loeffler M, Steinke V, et al. Risks of less common cancers in proven mutation carriers with lynch syndrome. J Clin Oncol. 2012;30(35):4409-15.
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- Hearle N, Schumacher V, Menko FH, et al. Frequency and spectrum of cancers in the Peutz-Jeghers syndrome. Clin Cancer Res. 2006;12(10):3209-15.

10. Masciari S, Dillon DA, Rath M, et al. Breast cancer phenotype in women with TP53 germline mutations: a Li-Fraumeni syndrome consortium effort. *Breast Cancer Res Treat.* 2012;133(3):1125-30.
11. Meindl A, Hellebrand H, Wiek C, et al. Germline mutations in breast and ovarian cancer pedigrees establish RAD51C as a human cancer susceptibility gene. *Nat Genet.* 2010;42(5):410-4.
12. Song H, Dicks E, Ramus SJ, et al. Contribution of Germline Mutations in the RAD51B, RAD51C, and RAD51D Genes to Ovarian Cancer in the Population. *J Clin Oncol.* 2015;33(26):2901-7.
13. Tan MH, Mester JL, Ngeow J, et al. Lifetime cancer risks in individuals with germline PTEN mutations. *Clin Cancer Res.* 2012;18(2):400-7.
14. Weber-Lassalle N, Hauke J, Ramser J, et al. BRIP1 loss-of-function mutations confer high risk for familial ovarian cancer, but not familial breast cancer. *Breast Cancer Res.* 2018 Jan 24;20(1):7. doi: 10.1186/s13058-018-0935-9.



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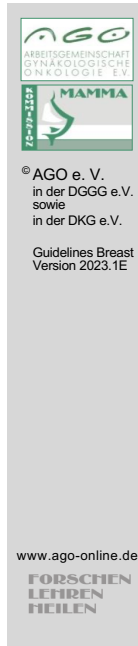
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Non BRCA-Associated Hereditary Cancer Syndromes with Increased Risk for Breast Cancer

Syndrome	Gene	Risk for malignancy
Li Fraumeni	<i>TP53</i>	Breast, endometrium, colorectal, small intestine, stomach, hepato biliary, skin, osteosarcoma, soft tissue sarcoma, urogenital, CNS, ACC, leukemia, lymphoma, lung
Cowden	<i>PTEN</i>	Breast, endometrium, thyroid, colorectal, kidney, melanoma
Hereditary diffuse gastric cancer syndrome	<i>CDH1</i>	Hereditary diffuse gastric cancer, lobular invasive breast cancer
Peutz-Jeghers Syndrome	<i>STK11/ LKB1</i>	Colorectal, small intestine, stomach, pancreas, testicle, endometrium
Lynch	<i>MLH1, MSH2, MSH6, PMS2, EPCAM</i>	Endometrium, ovary, colorectal, small intestine, stomach, hepato biliary, pancreas, kidney, urogenital, CNS
Ataxia telangiectasia (AT-Syndrome)	<i>ATM</i>	Breast cancer, leukemia, stomach, melanoma, sarcoma
Franconi Anämie	<i>BRCA2, BRIP1, RAD51C, PALB2</i>	AML, MDS, SCC, medulloblastoma, nephroblastoma, breast, pancreas, ovary

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2. Benusiglio PR, Malka D, Rouleau E, et al. CDH1 germline mutations and the hereditary diffuse gastric and lobular breast cancer syndrome: a multicentre study. *J Med Genet*. 2013;50(7):486-9
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4. di Masi A, Antoccia A. NBS1 Heterozygosity and Cancer Risk. *Curr Genomics*. 2008;9(4):275-81.
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6. Goldgar DE, Healey S, Dowty JG, et al. Rare variants in the ATM gene and risk of breast cancer. *Breast Cancer Res*. 2011;13(4):R73.
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12. Song H, Dicks E, Ramus SJ, et al. Contribution of Germline Mutations in the RAD51B, RAD51C, and RAD51D Genes to Ovarian Cancer in the Population. *J Clin Oncol.* 2015;33(26):2901-7.
13. Tan MH, Mester JL, Ngeow J, et al. Lifetime cancer risks in individuals with germline PTEN mutations. *Clin Cancer Res.* 2012;18(2):400-7.
14. Weber-Lassalle N, Hauke J, Ramser J, et al. BRIP1 loss-of-function mutations confer high risk for familial ovarian cancer, but not familial breast cancer. *Breast Cancer Res.* 2018 Jan 24;20(1):7. doi: 10.1186/s13058-018-0935-9



Non-Directive Counseling Regarding Preventive Measures

AGO ++

According to:

- The Genetic Diagnostic Law
- The Medical Devices Act (e.g. risk assessment)
- Application of software for risk calculation requires professional training and experience

Communicate:

- Absolute cancer risks within a manageable timeframe
- Risk and benefit of a multimodal intensive surveillance program
- Risk and benefit of preventive clinical methods
- Competing risks, e.g. risk of disease progression in relation to risk of a secondary primary in case women already affected by primary breast cancer

Allow appropriate time for consideration

1. Phi XA, Houssami N, Hooning MJ et al., Accuracy of screening women at familial risk of breast cancer without a known gene mutation.. Eur J of Cancer 2017;85:31-38

SOFTWARE (BOADICEA, IBIS)

1. Lee A, Mavaddat N, Wilcox AN et al. BOADICEA: a comprehensive breast cancer risk prediction model incorporating genetic and nongenetic risk factors. Genet Med. 2019 Aug;21(8):1708-1718. Erratum in: Genet Med. 2019 Feb 21;:
1. Terry MB, Liao Y, Whittemore AS et al. 10-year performance of four models of breast cancer risk: a validation study. Lancet Oncol. 2019 Apr;20(4):504-517.

Current Clinical Impact of Further Risk Genes

- Further moderate and low-risk gene variants are most likely transmitted by an oligo- or polygenic trait.
- The penetrance of such genes depends on the own and family cancer history.
- Individual low-risk variants increase the risk of disease only insignificantly. They have a multiplicative effect, so that the analysis of multiple gene regions (polygenic risk score, PRS) will be of clinical relevance.

	Oxford		
	LoE	GR	AGO
Clinical genetic testing of moderate-risk genes, e.g. gene panels	1b	B	+
Clinical genetic testing for low-risk variants (polygenic risk score, PRS)	2b	B	+*
Referral to centers of the GC-HBOC or cooperating centers	5	D	+

* Currently, moderately penetrant genes and low-risk variants should only be examined in the context of prospective cohort studies, such as that of the German consortium, in order to assess the clinical benefit.

Analyse von moderaten Risikogenen e.g. Genpanel

- Borde J, Ernst C, Wappenschmidt B et al. Performance of breast cancer polygenic risk scores in 760 female CHEK2 germline mutation carriers. J Natl Cancer Inst. 2020 Dec 29:djaa203. doi: 10.1093/jnci/djaa203. Epub ahead of print. PMID: 33372680.
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- Dorling L, Carvalho S, Allen J et al. Breast-Cancer Risk Genes — Association Analysis in More than 113,000 Women. January 20, 2021 DOI: 10.1056/NEJMoa1913948
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- Mavaddat N, Pharoah PD, Michailidou K, et al. Prediction of breast cancer risk based on profiling with common genetic variants. J Natl Cancer Inst. 2015;107(5).

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9. Michailidou K, Beesley J, Lindstrom S, et al. Genome-wide association analysis of more than 120,000 individuals identifies 15 new susceptibility loci for breast cancer. *Nat Genet.* 2015;47(4):373-80.
10. Lakeman IMM, van den Broek AJ, Vos JAM, et al. The predictive ability of the 313 variant-based polygenic risk score for contralateral breast cancer risk prediction in women of European ancestry with a heterozygous BRCA1 or BRCA2 pathogenic variant. *Genet Med.* 2021;23(9):1726-1737.
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Analyse von Niedrigisikovarianten (Polygenic risk score, PRS)

1. Jiao Y, Truong T, Eon-Marchais S, et al. Association and performance of polygenic risk scores for breast cancer among French women presenting or not a familial predisposition to the disease. *Eur J Cancer.* 2023 Jan;179:76-86.
2. Ohbe H, Hachiya T, Yamaji T et al.; Japan Public Health Center-based Prospective Study Group. Development and validation of genome-wide polygenic risk scores for predicting breast cancer incidence in Japanese females: a population-based case-cohort study. *Breast Cancer Res Treat.* 2022 Dec 20.
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Pathogenic Variants with Moderate to High Lifetime Risk for Breast Cancer

	Oxford		
	LoE	GR	AGO
Age-related risks for breast cancer			
▪ high: <i>BRCA1, BRCA2, PALB2</i>			
▪ high: <i>CDH1, PTEN, TP53; STK11[#]</i>			
▪ moderate: <i>ATM, CHEK2</i>			
▪ moderate: <i>BARD1, RAD51C, RAD51D</i>			
Clinical benefit* of a genetic test			
▪ <i>BRCA1, BRCA2</i>	1b	A	++ [°]
▪ <i>PALB2</i>	3a	B	+ [°]
▪ <i>CDH1, PTEN, TP53, STK11</i>	3b	B	+ [°]
▪ <i>ATM, BARD1, CHEK2, RAD51C, RAD51D</i>	3a	B	+/- [°]

* Take into account the effectiveness of preventive measures and competing risks when making clinical decisions.

[°] Participation in prospective registries or studies is highly recommended.

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CDH1:

1. Hearle N, Schumacher V, Menko FH, et al: Frequency and spectrum of cancers in the Peutz-Jeghers syndrome. Clin Cancer Res. 2006 May 15;12(10):3209-15
2. Kaurah P, MacMillan A, Boyd N, et al: Founder and recurrent CDH1 mutations in families with hereditary diffuse gastric cancer. JAMA. 2007 Jun 6;297(21):2360-72. doi: 10.1001/jama.297.21.2360. Epub 2007 Jun 3.
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PTEN:

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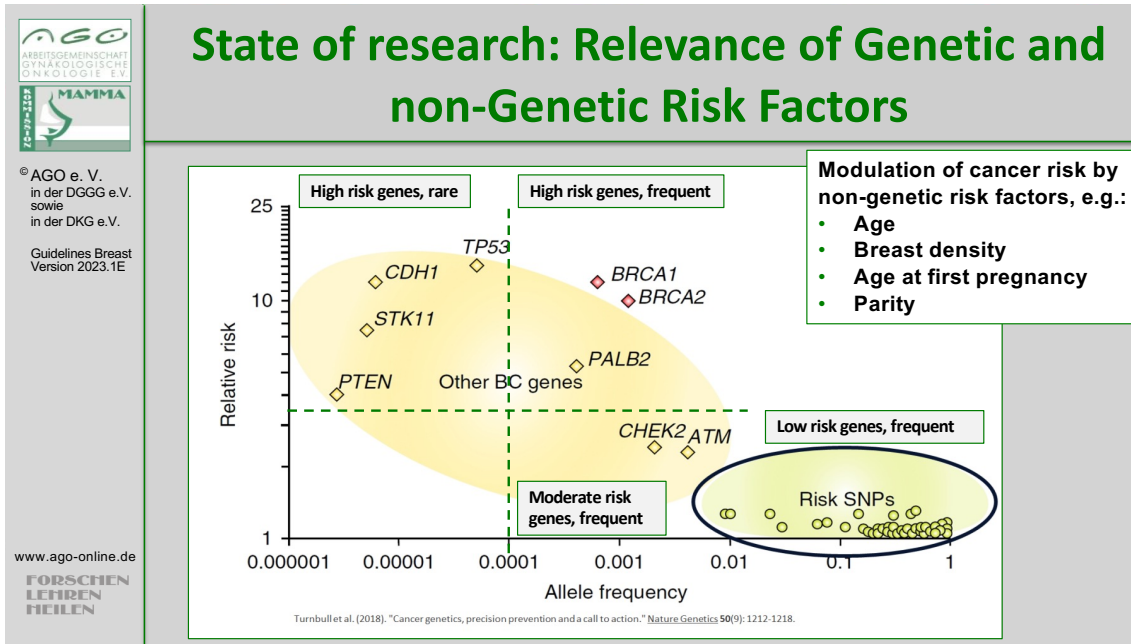
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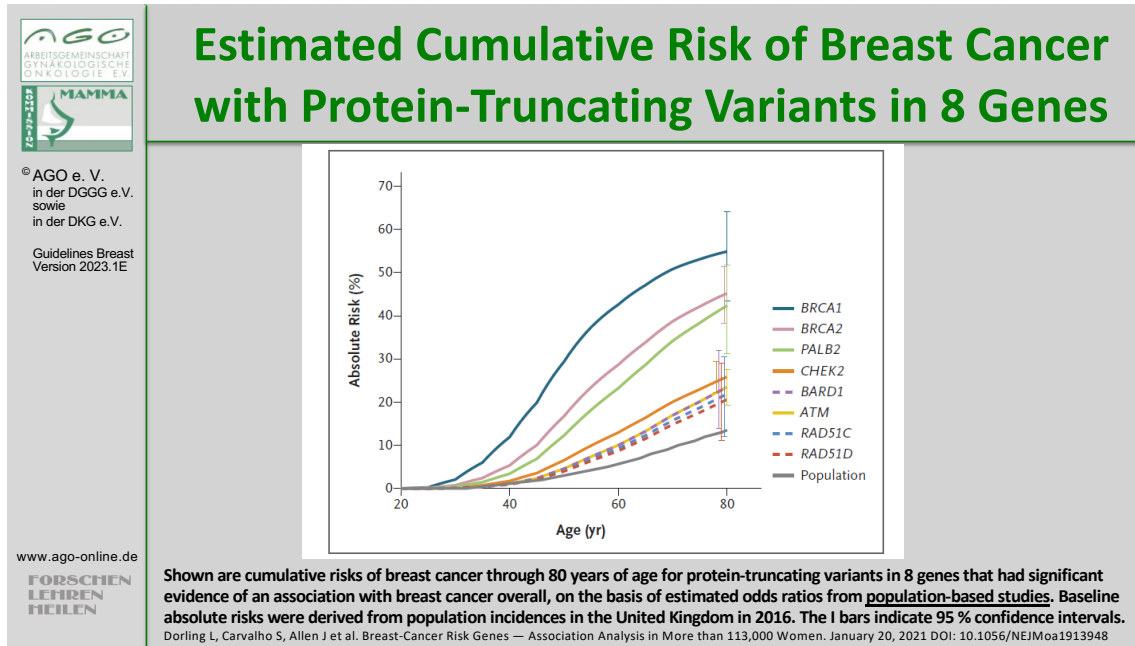
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Breast Cancer Risk for Individual Mutations (according NCCN 2023)		
Life time risk (age 20 y.)	High frequency	Rare frequency
High Risk (≥30%)	<i>BRCA1, BRCA2, PALB2</i>	<i>CDH1, PTEN, TP53, STK11</i>
Moderate Risk (17-29%)		<i>ATM, BARD1, CHEK2, RAD51C, RAD51D</i>
Low Risk (<17%)	<i>MSH2, MLH1, MSH6, PMS2, EPCAM</i>	
Unclear clinical relevance	<i>BRIP1, CDKN2A, FANCC, MRE11, MUTYH, NBN, NF1, RAD50, RECQL, RINT1, SLX4, SMARCA4, XRCC2</i>	




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Breast Cancer Risk Category

Definition of moderate/high risk for breast cancer

Breast cancer risk category

	Near population risk of breast cancer	Moderate risk of breast cancer	High risk of breast cancer
Lifetime risk from age 20	Less than 17%	Greater than 17% but less than 30%	30% or greater
Risk between ages 40 and 50	Less than 3%	3 to 8%	Greater than 8%

NICE (National Institute for Health and Care Excellence) guidance: Familial breast cancer: classification, care and managing breast cancer and related risks in people with a family history of breast cancer
Clinical guideline [CG164] Published: 25 June 2013 Last updated: 20 November 2019

1. NICE (National Institute for Health and Care Excellence) guidance: Familial breast cancer: classification, care and managing breast cancer and related risks in people with a family history of breast cancer. Clinical guideline [CG164] Published: 25 June 2013 Last updated: 20 November 2019
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IARC - classification of sequence variants (Plon et al., Human Mutation, 2008)

Proposed Classification System for Sequence Variants Identified by Genetic Testing		
Class	Description	Probability of being pathogenic
5	Definitely pathogenic	> 0,99
4	Likely pathogenic	0,95-0,99
3	Uncertain	0,05-0,949
2	Likely not pathogenic or of little clinical significance	0,001-0,049
1	Not pathogenic or no of clinical significance	< 0,001

**Only class 4 and class 5 variants are considered clinically relevant.
Class 3 are considered as Variants of Unknown Significance (VUS).**

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Variant of Unknown Significance (VUS): Problems and Questions

- „A Variant of Unknown Significance (VUS IARC class 3) is a genetic variant with unknown clinical relevance.“ (Plon et al. Hum Mutat 2008)
- Most VUS are extremely rare (≤ 3 variants in $> 80\%$ of families)
- Classification of sequence variants should be performed according to the IARC classification system
- Frequency of VUS (IARC class 3) increases with numbers of tested genes
- Clinical interpretation and decision making depending on the IARC classification system is not standardized yet
- In silico prediction tools (PolyPhen2, SIFT) are not adequate or sufficient for clinical decision making
- Additional analyses are required, e.g. in vitro splicing assay, functional assay, segregation analysis, co-occurrence analysis, large case / control studies

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Multimodal Intensive Surveillance Program*

		Oxford		
		LoE	GR	AGO
■	Program for BRCA-mutation carriers without BC			
■	For the detection of early stage cancers	2b	B	++
■	Clinical breast exam	≥ 25 years	Semi-annually	
■	Sonography	≥ 25 years	Semi-annually	
■	Mammogram	≥ 40 years	Every 1-2 years**	
■	Breast MRI	≥ 25 years	Annually	
■	For improvement of metastasis-free interval	2b	B	+
■	Radiotherapy of thoracic wall in the childhood (e.g. M. Hodgkin)	2a	B	++

*The multimodal early detection program should be carried out for women with a pathogenic mutation in risk genes and those with an increased calculated risk without a mutation within the framework of transparent quality assurance and appropriate evaluation;

** According to the recommendation of the German Consortium 2022: Depending on the assessability of the breast, the glandular parenchyma density and the previous mammographic findings every 1-2 years from the 40th-45th Age, under 40 years only after strict individual indication.

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High-Risk Breast Cancer Surveillance with MRI


	30-39 years		40-49 years		≥50 years	
	Detection rate (‰)	PPV (%)	Detection rate (‰)	PPV (%)	Detection rate (‰)	PPV (%)
BRCA1	43.2	29.4	21.8	25.5	30.5	33.3
BRCA2	22.7	23.3	24.3	27.5	16.3	23.5
BRCA1/2-non carriers with high risk	2.9	2.8	7.4	6.8	10.9	13.8

PPV: Positive predictive value

Detection performance of annual multimodality screening rounds with MRI by risk group and age.

Bick U, Engel C, Krug B, et al. High-risk breast cancer surveillance with MRI: 10-year experience from the German consortium for hereditary breast and ovarian cancer. *Breast Cancer Res Treat.* 2019;175(1):217–228. doi:10.1007/s10549-019-05152-9

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Modified Surveillance Program for BRCA-neg. Women at Moderate to High Risk or Survivors of Hodgkin Disease

Rationale:

- Increased risk of breast cancer after chest irradiation because of Hodgkin lymphoma in childhood (9–18 years)
- Increased risk of breast or ovarian cancer in women from BRCA1/2 negative families at risk that is, however, lower than in women from BRCA1/2 positive families
- Referral to centres of the GC-HBOC or cooperating centres for the evaluation of structured surveillance and follow-up

1. E-Learning DKG/FBREK, 2022
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Multimodal Surveillance Program for Female Carriers of Pathogenic BRCA Mutations after Unilateral Breast Cancer

	Oxford		
	LoE	GR	AGO
▪ Multimodal intensive surveillance program*			
▪ For detection of early stage breast cancers	2a	B	++
▪ Clinical breast exam			Semi-annually
▪ Sonography			Semi-annually
▪ Mammogram			Every 1-2 years**
▪ Breast MRI (until ACR1)			Annually
▪ For mortality reduction (10-year survival)	3a	C	+/-

* Aftercare should be carried out within the framework of transparent quality assurance and corresponding evaluation.

** According to the recommendation of the German Consortium 2022: Depending on the assessability of the breast, the glandular parenchyma density and the previous mammographic findings every 1-2 years from the 40th-45th Age, under 40 years only after strict individual indication.

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Surveillance for Male Carriers of Pathogenic BRCA Mutations*

	Oxford		
	LoE	GR	AGO
Currently, no specific surveillance is recommended →			
Early detection of cancer as part of standard care			
▪ BRCA1/2 mutation carrier: explanation of risks for cancer disease including male family members	5	D	++
▪ For breast cancer: self examination	5	D	+
▪ For prostate cancer: Compare German Guideline program	5	D	+

The lifetime risk of breast cancer in the general male population is 0.1 %. *BRCA1* mutation carriers have a risk of breast cancer of about 1 % and an up to 1.8 to 3.75 times higher risk for prostatic cancer ≤ 65y. *BRCA 2* mutation carriers have an up to 5–7 % lifetime risk for breast cancer and an up to 2.5 to 8.6 times higher risk for prostatic cancer ≤ 65y.

* Follow-up care / surveillance should be carried out as part of transparent quality assurance and appropriate evaluation.

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Surgical Prevention

	Oxford		
	LoE	GR	AGO
■ Risk-reducing unilateral or bilateral mastectomy (RRME) without the presence of clearly defined genetic risk factors	2a	B	-*
■ Axillary dissection or Sentinel lymph node excision during RRME	2a	B	--

* study participation recommended

RRME ohne gentisches Risiko

1. Kurian AW, Lichtensztajn DY, Keegan TH, et al. Use of and mortality after bilateral mastectomy compared with other surgical treatments for breast cancer in California, 1998-2011. JAMA. 2014;312(9):902-14.
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Sentinel-Lymphknoten Exzision bei RRME

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Surgical Prevention for Healthy Female *BRCA1/2* Mutation Carriers

	Oxford		
	LoE	GR	AGO
■ Risk-reducing bilateral salpingo-oophorectomy (RR-BSO)**	2a	B	
<ul style="list-style-type: none"> Reduces OvCa incidence and mortality 			++*
<ul style="list-style-type: none"> Reduces overall mortality 			++*
■ Risk-reducing bilateral mastectomy (RR-BM)			
<ul style="list-style-type: none"> Reduces BC incidence 	2b	B	+*
<ul style="list-style-type: none"> Reduces BC mortality in <i>BRCA1</i> mutation carriers*** 	2b	B	+*

* Study participation recommended
 ** The RR-BSO is recommended from about 35 years for *BRCA1* and from about 40 years for *BRCA2* mutation carriers, taking into account the age of ovarian cancer diagnosis in the family and the family planning status.
 *** No reduction in mortality could be shown for *BRCA2* mutation carriers. RRBm counselling should be individualised.

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- Kotsopoulos J, Huzarski T, Gronwald J, et al: Hereditary Breast Cancer Clinical Study Group. Bilateral Oophorectomy and Breast Cancer Risk in *BRCA1* and *BRCA2* Mutation Carriers. *J Natl Cancer Inst*. 2016 Sep 6;109(1). doi: 10.1093/jnci/djw177. Print 2017 Jan.
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Risk-reducing Interventions for BRCA1/2 Female Mutation Carriers Affected by Breast Cancer


	Oxford		
	LoE	GR	AGO
<ul style="list-style-type: none"> ▪ Risk-reducing bilateral salpingo-oophorectomy (RR-BSO) <ul style="list-style-type: none"> ▪ Reduces OvCa incidence and mortality ▪ Reduces overall mortality (contradictory results for reduction of cl BC incidence) 	2b	B	++*
<ul style="list-style-type: none"> ▪ Prophylactic contralateral mastectomy (RR-CM)* <ul style="list-style-type: none"> ▪ Reduces BC incidence and mortality 	2b	B	++*
<ul style="list-style-type: none"> ▪ Tamoxifen (reduces contralateral BC incidence) 	2b	B	+/-*
<ul style="list-style-type: none"> ▪ Indication for RR-CM should consider age at onset of first breast cancer in affected gene 	2a	B	+++*
<ul style="list-style-type: none"> ▪ RR-BM after ovarian cancer 	4	C	+/-**

* Study participation recommended

** Depends on tumor stage (FIGO I/II), recurrence free interval (≥ 5 yrs.), age

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Improved Overall Survival After Contralateral Risk-reducing Mastectomy in *BRCA1/2* Mutation Carriers with a History of Unilateral Breast Cancer: A Prospective Analysis

Analysis ^a	Group	Person years of observation	Deaths	Mortality ^b (95 % CI)	HR (95 % CI)
(a)	Surveillance	3007	65	21.6 (16.9-27.6)	Ref
	CRRM	1975	19	9.6 (6.1-15.1)	0.43 (0.26-0.72) ^c 0.49 (0.29-0.82) ^d
(b)	Surveillance	2673	56	21.0 (16.1-27.2)	Ref.
	CRRM	1837	18	9.8 (6.2-15.5)	0.46 (0.27-0.79) ^c 0.55 (0.32-0.95) ^d

^a Analysis (a) is the main analysis with start of observation being either the date of primary breast cancer (PBC) diagnosis or the date of DNA diagnosis, whichever came first. In the additional analysis (b), the observation starts either 2 years after PBC or at the date of DNA diagnosis, whichever came first, to exclude patients who presented with distant metastases or died within 2 years after PBC diagnosis (*n* = 17).

^b Per 1000 person years of observation.

^c Univariate analysis.

^d Multivariate analysis, adjusted for risk-reducing salpingo-oophorectomy. The following variables did not meet the criteria for incorporation in the multivariate Cox model as described in the Methods section, and were therefore not included in the multivariate analysis: type of mutation, year of birth, age at DNA diagnosis, age at PBC diagnosis, T-status, presence of positive lymph nodes, differentiation grade, hormone receptor status, HER2 status and treatments administered for PBC.

Abbreviations: CRRM, contralateral risk-reducing mastectomy; HR, Hazard ratio; CI, confidence interval.

We conclude that CRRM is associated with improved overall survival in *BRCA1/2* mutation carriers with a history of PBC. Further research is warranted to develop a model based on age at diagnosis and tumour and treatment characteristics that can predict survival benefit for specific subgroups of patients, aiming at further personalized counselling and improved decision making.

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Therapy of Germline Mutation-Associated Breast Cancer

	Oxford		
	LoE	GR	AGO
▪ Breast conserving surgery according common standard (adequate local tumor control in long time follow up, ~10 years observation)	2a	B	+
▪ Systemic therapy according to common standard	3a	B	+
▪ gBRCA mutation status is predictive for neoadjuvant chemotherapy in early TNBC	2b	B	
▪ gBRCA mutation status is predictive for Carboplatin (vs. Docetaxel) in metastatic breast cancer	1b	B	
PARP inhibitor (Her2-negative carcinoma):			
▪ eBC high risk:			
▪ Olaparib (in case of <i>gBRCA1/2</i> mutation)*	1b	A	++
▪ MBC:			
▪ Olaparib, Talazoparib in <i>gBRCA 1/2</i> mutation	1b	A	++
▪ Olaparib in <i>sBRCA 1/2</i> mutation (somatic mutation)	2b	B	+/-
▪ Olaparib in <i>PALB2</i> germ line mutation	2b	B	+/-

EBC: Early Breast Cancer; MBC: Metastatic Breast Cancer; * Use according to study inclusion criteria and approval

BCS bei BRCA 1/2 Mutationsträgern

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PARP-inhibitors mBC gPALB2mut

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Medical Prevention for Women at Increased Risk

	Oxford		
	LoE	GR	AGO
■ Tamoxifen for women > 35 years: Risk reduction of invasive BC, DCIS and LN	1a	A	+*
■ Raloxifen for postmenopausal women: Risk reduction of invasive BC only	1b	A	+*
■ AI for postmenopausal women	1b	A	+**

* Risk situation as defined in NSABP P1-trial (1.66% in 5 years) or according to #Tyrer-Cuzick model (IBIS-II)

** Significant risk reduction was seen for anastrozole for ovarian and endometrial cancer, as well as skin, colorectal, hematologic, thyroid and urinary tract cancers. Chemopreventive regimes should only be offered after individual and comprehensive counseling. The net benefit strongly depends on risk status, age and pre-existing risk factors for side effects.

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