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## Prediction of "BRCAness" in breast cancer by array comparative genomic hybridization

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### Citation

Joosse, S. A. (2012, March 27). *Prediction of "BRCAness" in breast cancer by array comparative genomic hybridization*. Retrieved from <https://hdl.handle.net/1887/18632>

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**Issue Date:** 2012-03-27

## List of Abbreviations

ALDH1	aldehyde dehydrogenase 1 family, member A1	F-CSGE	fluorescent conformational sensitive gel electrophoresis
APC	adenomatous polyposis coli	FE test	Fisher's exact test
ATM	ataxia telangiectasia mutated	FFPE	Formalin-fixed, paraffin embedded
BAC	bacterial artificial chromosome	GAPDH	glyceraldehyde-3-phosphate dehydrogenase
BARD1	BRCA1 associated RING domain 1	GE	gene expression
BCS	breast conserving surgery	GEP	gene expression profiling
BER	base excision repair	GEO	gene expression omnibus
BIC	breast cancer information core	HBOC	hereditary breast and ovarian cancer
bp	base pair	HD	helix-rich domain
BRCA	breast cancer, early onset	HDPB	high-dose platinum-based
BRCT	BRCA1 C terminus (domain)	H&E	hematoxylin and eosin
CHEK2	checkpoint kinase 2	HR	homologous recombination
CHK1	checkpoint kinase 1	HRD	homologous recombination deficient/deficiency
CGH	comparative genomic hybridization	IDC	invasive ductal carcinoma
CI	confidence interval	IGF	insulin-like growth factor
CISH	chromogenic in situ hybridization	IHC	immunohistochemistry
CNA	copy number alteration	ILC	invasive lobular carcinoma
CSC	cancer stem cell	IVS	intervening sequence
CTRM	c-terminal RAD51 binding motive	kbp	kilo base pairs
DCIS	ductal carcinoma in situ	KRT	keratin
DGGE	denaturing gradient gel electrophoresis	LC	lobular carcinoma
DHPLC	denaturing high-performance liquid chromatography	LCIS	lobular carcinoma in situ
DNA	deoxyribonucleic acid	LOD	logarithm of odds
DOL	degree of labeling	LOH	loss of heterozygosity
DSB	double-strand break	LOOCV	leave-one-out cross-validation
DSS1	deleted in split hand/foot protein 1	LUMC	Leiden University Medical Center
EDTA	ethylenediaminetetraacetic acid	Mbp	mega base pairs
EGFR	epidermal growth factor receptor	MLPA	multiplex ligation dependent probe amplification
ER	estrogen receptor	MRI	magnetic resonance imaging
ERBB2	human epidermal growth factor receptor 2	MYC	v-myc myelocytomatosis viral oncogene homolog
ESR1	estrogen receptor 1		
FAMA	fluorescent-assisted mismatch analysis		

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NCBI	National Center for Biotechnology Information
NGS	next generation sequencing
NHEJ	non-homologous end-joining
NKI	Netherlands Cancer Institute
NLS	nuclear localization signals
NPP	negative predictive power
OB	oligonucleotide binding
PALB2	partner and localizer of BRCA2
PAM	prediction analysis of microarrays
PARP	poly ADP ribose polymerase
PBM	prophylactic bilateral mastectomy
PBSO	prophylactic bilateral salpingo-oophorectomy
PCR	polymerase chain reaction
PPP	positive predictive power
PR	progesteron receptor
PTEN	phosphatase and tensin homolog
PTT	protein truncation test
RAD51	RAD51 homolog
RAP80	receptor-associated protein 80
RING	really interesting new gene
RNA	ribonucleic acid
RT-PCR	realtime PCR
SC	shrunken centroids
SCD	SQ-cluster domain
SCP	single-stranded conformational polymorphism
SDS	sodium dodecyl sulfate
SNP	single nucleotide polymorphism
SSB	single-strand break
SSC	saline-sodium citrate
STK11	serine/threonine kinase 11
STR	short tandem repeat
TBE	tris/borate/EDTA
TDGS	two-dimensional gene scanning
TP53	tumor protein 53
TNM	tumor/node/metastasis
ULS	universal linkage system
UV	unclassified variant
VEGF	vascular endothelial growth factor

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## Curriculum Vitae

Simon Joosse was born on August 6, 1983 in IJmuiden (Velsen), the Netherlands. He started studying medical biology at the faculty of Applied Sciences at Inholland University in Alkmaar in 2000. His graduation report on the subject "Gene inactivation for cell immortality" was carried out under the supervision of Dr. R.L. Beijersbergen at the department of Molecular Carcinogenesis at the Netherlands Cancer Institute, Amsterdam, in 2004. After graduation, he worked at the Netherlands Cancer Institute as a technician for two years and performed the Master Course on Experimental Oncology in 2006, organized by the Oncology school of Amsterdam. In 2007, Simon started working as a PhD student at the department of Experimental Therapy in the Netherlands Cancer Institute, under the supervision of Dr. P.M. Nederlof in the group of Dr. L.J. van 't Veer. In 2009, he moved to Hamburg, Germany, where he started working as scientist for the Department of Tumor Biology, University Medical Center Hamburg-Eppendorf, while simultaneously finishing his PhD studies performed at the Netherlands Cancer Institute. The results of this research are described in this thesis.





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## List of Publications

1. **Joosse SA**, Hannemann J, Spötter J, Bauche A, Andreas A, Müller V, Pantel K. *Changes in keratin expression during metastatic progression of breast cancer: impact on the detection of circulating tumor cells*. Clin Cancer Res. 2012; [Online publication ahead of print]
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4. Didraga MA, van Beers EH, **Joosse SA**, Brandwijk KI, Oldenburg RA, Wessels LF, Hogervorst FB, Ligtenberg MJ, Hoogerbrugge N, Verhoef S, Devilee P, Nederlof PM. *A Non-BRCA1/2 hereditary breast cancer sub-group defined by aCGH profiling of genetically related patients*. Breast Cancer Res Treat. 2011 Nov; 130(2):425-36.
5. **Joosse SA**, Brandwijk KI, Mulder M, Wesseling J, Hannemann J, Nederlof PM. *The genomic signature of BRCA1 deficiency in sporadic basal-like breast tumors*. Genes Chromosomes Cancer. 2011 Feb; 50:71-81.
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  12. Koski TA, Lehtonen HJ, Jee KJ, Ninomiya S, **Joosse SA**, Vahteristo P, Kiuru M, Karhu A, Salmakorpi H, Vanharanta S, Lehtonen R, Edgren H, Nederlof PM, Hietala M, Aittomäki K, Herva R, Knuutila S, Aaltonen LA, Launonen V. *Array comparative genomic hybridization identifies a distinct DNA copy number profile in renal cell cancer associated with hereditary leiomyomatosis and renal cell cancer*. Genes Chromosomes Cancer. 2009 Jul; 48(7):544-51.
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  16. Tischkowitz M, Hamel N, Carvalho MA, Birrane G, Soni A, van Beers EH, **Joosse SA**, Wong N, Novak D, Quenneville LA, Grist SA; kConFab, Nederlof PM, Goldgar DE, Tavtigian SV, Monteiro AN, Ladas JA, Foulkes WD. *Pathogenicity of the BRCA1 missense variant M1775K is determined by the disruption of the BRCT phosphopeptide-binding pocket: a multi-modal approach*. Eur J Hum Genet. 2008 Jul; 16(7):820-32.
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# Acknowledgements

This work has been made possible with the help of many people, whom I would like to thank here.

First of all, Dr. Petra Nederlof, thank you for giving me the opportunity to perform a PhD project under your supervision. Although my time as a PhD student in your lab was shorter as originally planned, we were able to produce some fine papers! I have learned an enormous amount and the years in your group were most productive.

Many thanks to my promoter Prof. Dr. Peter Devilee. You were always very kind and helpful to me and made it possible that I could finish my promotion even when I moved to Germany. Your music rocks!

I am indebted to Prof. Dr. Laura van 't Veer. It has been a real pleasure working in your group and having been witness of your groundbreaking research. Thank you for supporting me, especially in my last months at the NKI.

Prof. Dr. med. Klaus Pantel is gratefully acknowledged for allowing me to finish my PhD work at the UKE in Hamburg.

Erik van Beers, Renske Fles, and Kim Brandwijk - it has been a great experience working with you in our group! Erik, you have really taught me everything about breast cancer, array CGH, how to write a manuscript (properly and verbose), what it is like being a vegetarian, saxophone, and any other aspect of life on which we had time to talk about.

Much of the work in this thesis has been made possible with the help of the histopathology department, thank you very much for the endless amount of slides you cut and stained!! Frans Hogervorst, thank you for helping me on those occasions when I didn't understand about

those different mutation annotations again. Carla van Tiggelen and Sonja Springer for the administration of hundreds of samples. Hans Peterse, you have taught me much about pathology, thank you for always making time for me, spending hours behind the microscope, and being real critical about our work.

I would also like to acknowledge the people from our department C2 (formerly H6), especially: Adrian Begg, thank you for your time and for teaching me all about DNA repair; Els Wagenaar, it has been fun throwing away culture plates without names; Hans te Poele, it is a real joy knowing you, incredible how you could do your experiments, help every one with his/her (little) problems, run around though the building, and still have time for a chat; Thea Eggenhuizen, you knew the whole organization around the building, you have been a great help; and all other members of H6/C2, it has been a wonderful time.

My gratitude goes out to Ron Kerkhoven, Mike Heimerikx, Wim Brugman, and Marja Nieuwland from the microarray laboratory; Lodewyk Wessels for helping me with the bioinformatics and introducing me to MatLab, in which most calculation in this thesis have been performed; and Roderick Beijersbergen, for showing me the ins and outs of basic research during my internship at H2.

Last, but not least I would like to give special thanks to Juliane and Jannik. Thank you for proofreading everything I wrote (Juliane) and criticizing my plots (Jannik). Maar in het bijzonder bedank ik jullie dat jullie er elke dag voor mij zijn en de vreugde in mijn leven brengen.





