

Integrating clinicopathological and molecular data in the breast cancer patient: towards precision medicine
Engels, C.C.

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Author: Engels, Charla Chábeli

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Chapter 8

The clinical value of HER-2 overexpression and PIK3CA mutations in the older breast cancer population: a FOCUS study analysis

Charla C. Engels*, Mandy Kiderlen*, Esther Bastiaannet, Ronald van Eijk, Antien Mooyaart, Vincent T.H.B.M. Smit, Anton J.M. de Craen, Peter J.K. Kuppen, Judith R. Kroep, Cornelis J.H. van de Velde, Gerrit Jan Liefers

* Both authors contributed equally

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ABSTRACT

Background

Studies to confirm the effect of acknowledged prognostic markers in older breast cancer patients are scarce. The aim of this study was to evaluate the prognostic value of HER-2 overexpression and PIK3CA mutations in older breast cancer patients.

Design

Female breast cancer patients aged 65 years or older, diagnosed between 1997-2004 in a geographical region in The Netherlands, with an invasive, non-metastatic tumour and tumour material available, were included in the study. The primary endpoint was relapse free period and secondary endpoint was relative survival. Determinants were immunochemical HER-2 scores (0/1+, 2+ or 3+) and PIK3CA as a binary measure.

Results

Overall, 1,698 patients were included, and 103 had a HER-2 score of 3+. HER-2 over-expression was associated with a higher recurrence risk (5 years recurrence risk 34% vs. 12%, adjusted p=0.005), and a worse relative survival (10 years relative survival 48% vs. 84% for HER-2 negative; p=0.004). PIK3CA mutations had no significant prognostic effect.

Conclusion

We showed, in older breast cancer patients, that HER-2 overexpression was significantly associated with a worse outcome, but PIK3CA mutations had no prognostic effect. These results imply that older patients with HER-2 overexpressing breast cancer might benefit from additional targeted anti-HER-2 therapy.

INTRODUCTION

Over the last decades an increased aged population in Western countries paralleled a marked increase in the incidence of age-related tumours, such as breast cancer ¹. In the United States of America, more than 40% of the women diagnosed with breast cancer were 65 years of age or older in 2013 ². Due to the continuously increasing life expectancy, it is assumed that this number will further increase the coming years.

Human epidermal growth factor receptor-2 (HER-2) overexpression occurs in approximately 15-20 percent of invasive breast carcinomas ^{3;4}. Amplification of the HER-2 gene is associated with a more aggressive tumour phenotype ⁵, and if left untreated, is associated with worse clinical outcomes ⁶⁻⁹. Detection of the amplification of this oncogene is widely performed and often routinely used in clinical settings ¹⁰, mainly for allocation of anti-HER-2 therapy, consisting of trastuzumab or pertuzumab in the (neo)adjuvant setting ¹¹⁻¹⁶ and in the metastatic setting of lapatinib and trastuzumabemtansine ^{17;18}. These anti-HER-2 therapies, in addition to cytotoxic chemotherapy, can result in a substantial reduction of recurrences, both in node negative and node positive breast cancer patients ¹⁹⁻²¹.

A well-known accomplice of HER-2 overexpression is the PIK3CA mutation, leading to aberrant activation of the Phosphatidylinositol 3-kinase (PI3K)/AKT pathway, co-occurring in approximately 40% of HER-2 amplified tumors where it supports tumor growth ^{22;23}. The aberrant activation of the PI3K pathway correlates with reduced response to HER-2-directed therapies and accelerates HER-2 mediated breast epithelial transformation and metastatic progression ^{5;24;25}. In the younger population PI3K inhibitors are considered promising novel therapeutic modalities for the treatment of breast cancer.

An important shortcoming in current clinical breast cancer research is that the majority of studies are performed in a relatively young or fit elderly breast cancer population. This hampers the extrapolation of results of prognostic as well as therapeutic studies for older patients ^{26;27}.

The high incidence of cancer in the growing elderly population encourages to investigate the effect of potential prognostic and/or predictive markers in this population specifically. Some studies suggest that HER-2 positive breast cancer occurs less frequently in older patients ^{28;29}, but other studies fail to confirm this ³⁰. The clinical value of HER-2 overexpression and its treatment in the older population, well-known for competing clinical events, is not known. Because the accrual of older patients in clinical trials for anti-HER-2 therapy was poor, and an increased incidence of cardiac adverse events of trastuzumab was reported, its use is less widespread in older women than younger women ³¹. However, evidence for the omission of anti-HER-2 therapy is lacking.

In summary, there are no conclusive data on the prognostic effect of HER-2 overexpression and PIK3CA mutations in the elderly breast cancer patient. Therefore, in order to restore the clinical interest of this neglected, but potentially very valuable treatment target for the elderly breast cancer patients, this study aims to elucidate the prognostic value of HER-2 overexpression, PIK3CA mutations and the interplay between these two markers in a population-based cohort of older breast cancer patients, that was not exposed to any anti-HER-2-therapy.

MATERIAL AND METHODS

Patients and tumors

For this study, patients with invasive, non-metastatic breast cancer from the FOCUS cohort (Female breast cancer in the elderly, Optimizing Clinical guidelines USing clinic-pathological and molecular data) who received surgery and had formalin fixed paraffin embedded (FFPE) intra-operative tumor samples available with successful measurements of the HER-2 status and/or PIK3CA mutations were included. The FOCUS cohort has been described extensively in previous publications 32 . Briefly, the cohort consists of all women aged \geq 65 years at time of diagnosis, with invasive and *in situ* breast cancer, diagnosed between 1997 and 2004 in the South Western region of The Netherlands. Follow-up on survival status was available until the 1st of January 2013. All tumor samples were handled in a coded fashion, according to national ethical guidelines ("Code for Proper Secondary Use of Human Tissue", Dutch Federation of Medical Scientific Societies).

Immunohistochemistry for HER-2

The immunohistochemical staining against HER-2 was performed on tissue sections of 4µm from intra-operatively derived FFPE tumor material of the FOCUS cohort processed into a Tissue Micro Array (TMA) according to the previously described protocol ³³. Sections were incubated at room temperature with Polyclonal Rabbit Anti-Human c-erbB-2 Oncoprotein (DAKO, Denmark (A0485); 1:100, diluted in 1% PBSA) for 20 minutes, followed by Envision anti-rabbit (DAKO, Denmark, Cytomation K4003) for 20 minutes. DAB was used for visualization of positively stained breast tumor tissue on the TMA and counterstained with haematoxylin. All slides were stained simultaneously to avoid inter-assay variation. Strongly c-erbB-2 Oncoprotein positive brain tumor tissue served as positive- and a negative-control, the latter was obtained by omitting the primary antibody.

Evaluation of immunostaining

Microscopic quantification of the c-erbB-2 Oncoprotein was performed by two independent observers (C.E and A.M.). C-erbB-2 Oncoprotein was scored as follows: 0 for no

staining at all or incomplete or faint/barely perceptible membrane staining in <10% of the invasive tumor cells; 1+ for a faint/barely perceptible partial membrane staining in >10% of the tumor cells; 2+ for weak to moderate complete membrane staining in >10% of the tumor cells; and 3+ for strong to complete membrane staining in >30% of the tumor cells (Figure 1). For all patients, the highest score out of the three punches of the

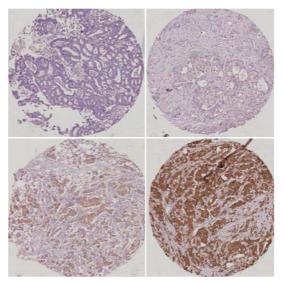


Figure 1: Immunohistochemical staining for HER-2 overexpression (*ERBB2*). Top left: HER2 score 0. Top right: HER2 score 1+. Bottom left: HER2 score 2+; and bottom right: HER2 score 3+.

same tumor was used for statistical analysis. If one or more punches were missing, the highest score of the remaining punch(es) was included for analyses.

PIK3CA mutation analysis

DNA was extracted from 2.0 mm diameter FFPE breast cancer tissue cores of 896 patients, using a fully automated system (Tissue Preparation System with VERSANT Tissue Preparation Reagents, Siemens Healthcare Diagnostics, Tarrytown, NY, USA) as described previously ³⁴. Hydrolysis probes assays were performed for the major known mutations (hotspots) in exon 9, c.1624G>A; p.E542K, c.1633G>A; p.E545K and in exon 20 the c.3140A>G; p.H1047R as described before ³⁵. Hydrolysis probe assays were analyzed using qPCR analysis software (CFX manager version 3/0, Bio-Rad). Mutation detection was performed by two observers (C.E. and R.E.) using DNA variant analysis software (Mutation Surveyor version 4.0.9, Softgenetics, State College, PA, USA). All primers and probes used for the assays are listed in Supplementary Table 1.

Statistical analysis

Statistical analyses were performed using the statistical packages SPSS (version 20.0 for Windows, IBM SPSS statistics) and Stata SE 12.0.

The primary endpoint was Relapse-Free Period (RFP), defined as the time from date of diagnosis until any recurrence (any registered loco-regional recurrence, distant recurrence or contralateral breast cancer). The Cumulative Incidence Competing Risks method was used for calculating the cumulative incidence of recurrence, taking into account the competing risk of death ³⁶. Fine & Gray competing risks regression analyses were used for univariable and multivariable analysis for RFP, taking into account the competing risk of death without recurrence ³⁷. Multivariable analyses were adjusted for age, TNM stage, grade, estrogen receptor (ER) and progesterone receptor (PR)) and type of breast surgery, axillary surgery, radiotherapy, endocrine therapy and chemotherapy (no anti-HER-2-therapy). Because the total follow-up time was longer than the followup time for RFP, the secondary endpoint was relative survival, calculated as the ratio between the observed survival in the cohort and the expected survival as calculated from the age-, sex- and year-matched background population ³⁸. Patients with missing data on the determinant of interest due to material handling were excluded from the statistical analyses regarding that determinant. First, it was checked if the patients with missing data had no significant survival difference from the patients without missing values, to confirm that there is no association with general prognosis.

Immunohistochemical HER-2 scores 0 and 1+ were considered HER-2 negative and HER-2 scores 2+ and 3+ were analyzed as separate categories. In clinical practice, tumors with an immunohistochemical HER-2 score of 2+ are considered borderline positive, therefore, it is recommended to validate the immunohistochemical HER-2 2+ results using in situ hybridization ³⁹⁻⁴¹. Unfortunately, it was not feasible to perform this additional test in our population. PIK3CA mutations were analyzed as dichotomous variable (negative or positive) in all patients, and stratified for HER-2 status.

RESULTS

Patient and tumour characteristics

A total of 1,932 tumour blocks were available for immunohistochemical staining. After material handling and staining, 1,698 patients were available for HER-2 analyses, and 912 patients for PIK3CA analyses. For the combined HER-2/PIK3CA analyses, 896 postmenopausal breast cancer patients were included in the analyses. The weighted Kappa for the HER-2 immunohistochemistry was 0.78 (SE 0.03), indicative of good inter-observer agreement'. There was no significant difference in overall survival between all patients

Table 1: Patient characteristics, tumor characteristics and treatment

Part Part		HER-2	negative	HER-2	2+	HER-2	23+	P
Name of sears (mean, SD) 76.1 ₹2.1 76.0 ₹2.1 75.0 ₹2.1 0.272 Number of comorbidities 76.0 ₹2.1 75.0 ₹2.1 0.272 Number of comorbidities 0.481		(N=1,3	26)	(N=26	9)	(N=10	3)	
Number of comorbidities		N	%	N	%	N	%	difference**
0-1 627 47,3 586 44,2 113 8.5 10.4 2-4 586 44,2 114 42,4 28 10.4 5 or more 113 8.5 39 37,9 13 12.6 TNM stage I 478 36.0 87 32.3 21 20.4 III 145 10.9 36.5 13.4 14 13.6 Missing 32 24 8 30.0 2 19 Grade ***********************************	Age in years (mean, SD)	76.1 (7	.2)	76.0 (7	7.2)	75.0 (6	5.9)	0.272
2-4 586 44.2 114 42.4 28 10.4 5 or more 113 8.5 39 37.9 13 12.6 TNM stage I 478 36.0 87 32.3 21 20.4 III 145 10.9 36 13.4 14 13.6 Missing 2 24 8 3.0 2 1.9 Grade	Number of comorbidities							0.481
Sor more 113 8.5 39 37.9 13 12.6 TNM stage Jana 18.8 36.0 87 32.3 21 20.4 II 478 36.0 87 32.3 21 20.4 III 145 10.9 36 13.4 14 13.6 Missing 32 2.4 8 3.0 2 1.9 Grade Footh Footh Footh 1 184 13.9 27 10.0 4 3.9 2 422 31.8 87 32.3 16 15.5 3 316 23.8 90 33.5 62 60.2 Missing 404 30.5 65 24.2 21 20.4 Morphology Till 8 81.0 91 88.3 Secure of Colspan="6">Missing 10ctal 157 11.8 18 6.7 4<	0-1	627	47.3	586	44.2	113	8.5	
Name	2-4	586	44.2	114	42.4	28	10.4	
Note	5 or more	113	8.5	39	37.9	13	12.6	
III	TNM stage							0.050
III	1	478	36.0	87	32.3	21	20.4	
Missing 32 2.4 8 3.0 2 1.9 Grade < 0.001 1 184 13.9 27 10.0 4 3.9 2 422 31.8 87 32.3 16 15.5 3 316 23.8 90 33.5 62 60.2 Missing 404 30.5 65 24.2 21 20.4 Morphology • 0.001 Ductal 969 73.1 218 81.0 91 88.3 Lobular 157 11.8 18 6.7 4 3.9 Other/missing 200 15.1 33 12.3 8 7.8 ER/PR • 180 13.6 33 12.3 54 52.4 Positive 980 73.9 209 77.7 35 34.0 Missing 166 12.5 27 10.0 14<	II	671	50.6	138	51.3	66	64.1	
Carace	III	145	10.9	36	13.4	14	13.6	
1 184 13.9 27 10.0 4 3.9 2 422 31.8 87 32.3 16 15.5 3 316 23.8 90 33.5 62 60.2 Missing 404 30.5 65 24.2 21 20.4 Morphology Ductal 969 73.1 218 81.0 91 88.3 Lobular 157 11.8 18 6.7 4 3.9 Other/missing 200 15.1 33 12.3 8 7.8 Cherymissing 180 13.6 33 12.3 54 52.4 Positive 980 73.9 209 77.7 35 34.0 Missing 166 12.5 27 10.0 14 13.6 Kife7 ***C********************************	Missing	32	2.4	8	3.0	2	1.9	
2 422 31.8 87 32.3 16 15.5 3 316 23.8 90 33.5 62 60.2 Missing 404 30.5 65 24.2 21 20.4 Morphology • • • • • • • • • • • • • • • • • • •	Grade							<0.001
3 316 23.8 90 33.5 62 60.2 Missing 404 30.5 65 24.2 21 20.4 Morphology 0.001 Ductal 969 73.1 218 81.0 91 88.3 Lobular 157 11.8 18 6.7 4 3.9 Other/missing 200 15.1 33 12.3 8 7.8 ER/PR	1	184	13.9	27	10.0	4	3.9	
Morphology 73.1 218 81.0 91 88.3 Lobular 969 73.1 218 81.0 91 88.3 Lobular 157 11.8 18 6.7 4 3.9 Other/missing 200 15.1 33 12.3 8 7.8 ER/PR TS TS 209 77.7 35 34.0 Positive 980 73.9 209 77.7 35 34.0 Missing 166 12.5 27 10.0 14 13.6 Ki67 TS 28 84.8 75 72.8 70.001 Negative 1094 82.5 228 84.8 75 72.8 72.8 Positive 84 6.3 20 7.4 21 20.4 20.04 Missing 148 11.2 21 7.8 7 6.8 7 6.8 BCS 510 38.5 95	2	422	31.8	87	32.3	16	15.5	
Morphology 73.1 218 81.0 91 88.3 Lobular 157 11.8 18 6.7 4 3.9 Other/missing 200 15.1 33 12.3 8 7.8 ER/PR	3	316	23.8	90	33.5	62	60.2	
Ductal 969 73.1 218 81.0 91 88.3 Lobular 157 11.8 18 6.7 4 3.9 Other/missing 200 15.1 33 12.3 8 7.8 ER/PR < 0.001 Negative 180 13.6 33 12.3 54 52.4 Positive 980 73.9 209 77.7 35 34.0 Missing 166 12.5 27 10.0 14 13.6 Ki67 < 1001	Missing	404	30.5	65	24.2	21	20.4	
Lobular 157 11.8 18 6.7 4 3.9 Other/missing 200 15.1 33 12.3 8 7.8 ER/PR < 0.001 Negative 180 13.6 33 12.3 54 52.4 Positive 980 73.9 209 77.7 35 34.0 Missing 166 12.5 27 10.0 14 13.6 Ki67 < 0.001 Negative 1094 82.5 228 84.8 75 72.8 Positive 84 6.3 20 7.4 21 20.4 Missing 148 11.2 21 7.8 7 6.8 Breast surgery* 510 38.5 95 35.3 24 23.3 Mastectomy 816 61.5 174 64.7 79 76.7 Axillary surgery* 181 13.7 35 </td <td>Morphology</td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td>0.001</td>	Morphology							0.001
Other/missing 200 15.1 33 12.3 8 7.8 ER/PR < 0.001 Negative 180 13.6 33 12.3 54 52.4 Positive 980 73.9 209 77.7 35 34.0 Missing 166 12.5 27 10.0 14 13.6 Ki67 < 1094 82.5 228 84.8 75 72.8 Positive 84 6.3 20 7.4 21 20.4 Missing 148 11.2 21 7.8 7 6.8 Breast surgery* 510 38.5 95 35.3 24 23.3 Mastectomy 816 61.5 174 64.7 79 76.7 Axillary surgery* 181 13.7 35 13.0 16 15.5 Sentinel node 311 23.5 71 26.4 16 15.5	Ductal	969	73.1	218	81.0	91	88.3	
ER/PR <0.001 Negative 180 13.6 33 12.3 54 52.4 Positive 980 73.9 209 77.7 35 34.0 Missing 166 12.5 27 10.0 14 13.6 Ki67 <0.001	Lobular	157	11.8	18	6.7	4	3.9	
Negative 180 13.6 33 12.3 54 52.4 Positive 980 73.9 209 77.7 35 34.0 Missing 166 12.5 27 10.0 14 13.6 Ki67	Other/missing	200	15.1	33	12.3	8	7.8	
Positive 980 73.9 209 77.7 35 34.0 Missing 166 12.5 27 10.0 14 13.6 Ki67 < 0.001 Negative 1094 82.5 228 84.8 75 72.8 Positive 84 6.3 20 7.4 21 20.4 Missing 148 11.2 21 7.8 7 6.8 Breast surgery* D.007 BCS 510 38.5 95 35.3 24 23.3 Mastectomy 816 61.5 174 64.7 79 76.7 Axillary surgery* 181 13.7 35 13.0 16 15.5 Sentinel node 311 23.5 71 26.4 16 15.5 ALND 834 62.9 163 60.6 71 68.9 Adjuvant radiotherapy No.947	ER/PR							<0.001
Missing 166 12.5 27 10.0 14 13.6 Ki67 < 0.001 Negative 1094 82.5 228 84.8 75 72.8 Positive 84 6.3 20 7.4 21 20.4 Missing 148 11.2 21 7.8 7 6.8 Breast surgery* BCS 510 38.5 95 35.3 24 23.3 Mastectomy 816 61.5 174 64.7 79 76.7 Axillary surgery* 181 13.7 35 13.0 16 15.5 Sentinel node 311 23.5 71 26.4 16 15.5 ALND 834 62.9 163 60.6 71 68.9 Adjuvant radiotherapy 868 51.7 142 52.8 54 52.4	Negative	180	13.6	33	12.3	54	52.4	
Ki67 <0.001 Negative 1094 82.5 228 84.8 75 72.8 Positive 84 6.3 20 7.4 21 20.4 Missing 148 11.2 21 7.8 7 6.8 Breast surgery* • • • • • • • • • • • • • • • • • • •	Positive	980	73.9	209	77.7	35	34.0	
Negative 1094 82.5 228 84.8 75 72.8 Positive 84 6.3 20 7.4 21 20.4 Missing 148 11.2 21 7.8 7 6.8 Breast surgery* BCS 510 38.5 95 35.3 24 23.3 Mastectomy 816 61.5 174 64.7 79 76.7 Axillary surgery* 181 13.7 35 13.0 16 15.5 Sentinel node 311 23.5 71 26.4 16 15.5 ALND 834 62.9 163 60.6 71 68.9 Adjuvant radiotherapy No 686 51.7 142 52.8 54 52.4	Missing	166	12.5	27	10.0	14	13.6	
Positive 84 6.3 20 7.4 21 20.4 Missing 148 11.2 21 7.8 7 6.8 Breast surgery* 0.007 BCS 510 38.5 95 35.3 24 23.3 Mastectomy 816 61.5 174 64.7 79 76.7 Axillary surgery* 181 13.7 35 13.0 16 15.5 Sentinel node 311 23.5 71 26.4 16 15.5 ALND 834 62.9 163 60.6 71 68.9 Adjuvant radiotherapy No 686 51.7 142 52.8 54 52.4	Ki67							<0.001
Missing 148 11.2 21 7.8 7 6.8 Breast surgery* 0.007 BCS 510 38.5 95 35.3 24 23.3 Mastectomy 816 61.5 174 64.7 79 76.7 Axillary surgery* 181 13.7 35 13.0 16 15.5 Sentinel node 311 23.5 71 26.4 16 15.5 ALND 834 62.9 163 60.6 71 68.9 Adjuvant radiotherapy 0.947 No 686 51.7 142 52.8 54 52.4	Negative	1094	82.5	228	84.8	75	72.8	
Breast surgery* 0.007 BCS 510 38.5 95 35.3 24 23.3 Mastectomy 816 61.5 174 64.7 79 76.7 Axillary surgery* 181 13.7 35 13.0 16 15.5 Sentinel node 311 23.5 71 26.4 16 15.5 ALND 834 62.9 163 60.6 71 68.9 Adjuvant radiotherapy 0.947 No 686 51.7 142 52.8 54 52.4	Positive	84	6.3	20	7.4	21	20.4	
BCS 510 38.5 95 35.3 24 23.3 Mastectomy 816 61.5 174 64.7 79 76.7 Axillary surgery* 181 13.7 35 13.0 16 15.5 Sentinel node 311 23.5 71 26.4 16 15.5 ALND 834 62.9 163 60.6 71 68.9 Adjuvant radiotherapy 0.947 No 686 51.7 142 52.8 54 52.4	Missing	148	11.2	21	7.8	7	6.8	
Mastectomy 816 61.5 174 64.7 79 76.7 Axillary surgery* 181 13.7 35 13.0 16 15.5 Sentinel node 311 23.5 71 26.4 16 15.5 ALND 834 62.9 163 60.6 71 68.9 Adjuvant radiotherapy 0.947 No 686 51.7 142 52.8 54 52.4	Breast surgery*							0.007
Axillary surgery* 0.296 No axillary surgery 181 13.7 35 13.0 16 15.5 Sentinel node 311 23.5 71 26.4 16 15.5 ALND 834 62.9 163 60.6 71 68.9 Adjuvant radiotherapy 0.947 No 686 51.7 142 52.8 54 52.4	BCS	510	38.5	95	35.3	24	23.3	
No axillary surgery 181 13.7 35 13.0 16 15.5 Sentinel node 311 23.5 71 26.4 16 15.5 ALND 834 62.9 163 60.6 71 68.9 Adjuvant radiotherapy No 686 51.7 142 52.8 54 52.4	Mastectomy	816	61.5	174	64.7	79	76.7	
Sentinel node 311 23.5 71 26.4 16 15.5 ALND 834 62.9 163 60.6 71 68.9 Adjuvant radiotherapy 0.947 No 686 51.7 142 52.8 54 52.4	Axillary surgery*							0.296
ALND 834 62.9 163 60.6 71 68.9 Adjuvant radiotherapy 0.947 No 686 51.7 142 52.8 54 52.4	No axillary surgery	181	13.7	35	13.0	16	15.5	
Adjuvant radiotherapy 0.947 No 686 51.7 142 52.8 54 52.4	Sentinel node	311	23.5	71	26.4	16	15.5	
No 686 51.7 142 52.8 54 52.4	ALND	834	62.9	163	60.6	71	68.9	
	Adjuvant radiotherapy							0.947
Yes 640 48.3 127 47.2 49 47.6	No	686	51.7	142	52.8	54	52.4	
	Yes	640	48.3	127	47.2	49	47.6	

	HER-2	negative	HER-2	2+	HER-	2 3+	Р	
	(N=1,3	26)	(N=26	9)	(N=10	03)	for	
	N	%	N	%	N	%	difference**	
Adjuvant endocrine therapy							0.008	
No	728	54.9	127	47.2	66	64.1		
Yes	598	45.1	142	52.8	37	35.9		
Adjuvant chemotherapy							0.003	
No	1250	94.3	257	95.5	89	86.4		
Yes	76	5.7	12	4.5	14	13.6		

Table 1: Patient characteristics, tumor characteristics and treatment (continued)

Abbreviations: ER: estrogen receptor PR: progesterone receptor BCS: breast conserving surgery ALND=axillary lymph node dissection

from the original cohort (with or without tumour blocks available) and the patients with available data on respectively HER-2 or PIK3CA (Log Rank p=0.537 and p=0.298).

Patient, tumour and treatment characteristics are shown in Table 1 by HER-2 score. Overall, mean age at diagnosis was 76 years (standard deviation 7.2 years). The majority of patients presented with early stage breast cancer (stage I 34.5%, stage II 51.5%, stage III 11.5%) of ductal morphology (75.3%). In the majority of the patients a mastectomy (63.0%) was performed. Merely six percent of patients were treated with adjuvant chemotherapy (not including anti-HER-2-therapy). In this cohort, 78% of the breast cancers were classified as HER-2 negative (N=1,326). Two hundred sixty-nine patients had a HER-2 score of 2+ (16%), and 103 patients had a HER-2 score of 3+ (6%). HER-2 overexpression was significantly associated with higher tumor grade (p<0.001), ductal tumor morphology (p=0.001), negative hormone receptor status (p<0.001), and a higher Ki-67 proliferation rate (p<0.001). In addition, HER-2 overexpression was associated with undergoing mastectomy rather than breast conserving surgery (p=0.007), less endocrine therapy (p=0.008) and more chemotherapy (without anti-HER2 therapy) (p=0.003).

Among all patients from whom PIK3CA status was available, 30% had a PIK3CA mutation. In our data, PIK3CA mutations were not associated with HER-2 overexpression (p=0.7).

Relapse free period

For RFP, median follow-up was 5.3 years (range: 0-13.5 years). HER-2 negative patients had a cumulative recurrence risk of 12% at 5 years, as compared to 12% for patients with a HER-2 score of 2+ and 34% for patients with a HER-2 score of 3+. In adjusted analysis, patients with a score of 2+ had an equal recurrence free period at 5 years as compared to HER-2 negative patients (adjusted sub-distribution Hazard Ratio (SHR) 0.95, 95% Con-

^{*}The most extended therapy was taken into account **p-values are calculated by the Pearson Chi-Square for categorical variables, and with an oneway ANOVA test for continuous variables. P-values in bold font indicate a statistically significant difference between the groups at the p-level of <0.05.

		Recurrence		959	% CI			95	% CI	_
	N of events	risk at 5 years (%)	SHR	lower	upper	P	SHR*	lower	upper	P
HER2										
negative:										
0/1+	155	12%	1 (Ref	ference)			1 (Refe	rence)		
2+	32	12%	1.06	0.76	1.47	0.747	0.95	0.67	1.35	0.790
3+	34	34%	2.68	1.86	3.86	<0.001	1.81	1.19	2.74	0.005
РІКЗСА										
negative	103	17%	1 (Refe	erence)			1 (Referen	ce)		
positive	42	16%	0.98	0.71	1.35	0.892	1.00	0.72	1.41	0.987

Table 2: Relapse free period (Fine & Gray regression)

^{*}adjusted for age, morphology, grade, hormone receptor status, tumor stage, type of breast surgery, type of axillary surgery, radiotherapy, endocrine therapy, chemotherapy

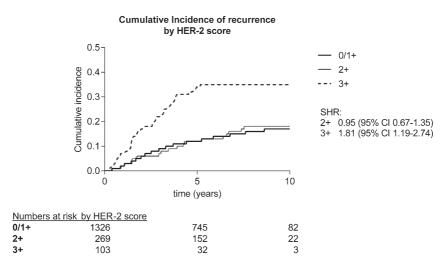


Figure 2: Cumulative incidence of recurrence by HER-2 score.

fidence Interval (CI) 0.67-1.35; p=0.8), patients with a score of 3+ had significantly worse relapse free period (adjusted SHR 1.81, 95% CI 1.19-2.74, p=0.005) (Table 2). Cumulative incidence curves are depicted in Figure 2. Among patients without PIK3CA mutations, the cumulative incidence of recurrences at 5 years was 17%, as compared to 16% of patients with a PIK3CA mutation. There was no statistically significant difference in RFP between patients with and without a PIK3CA mutation (p=0.2), also after stratifying for HER-2 status (Supplementary file 2).

Table 3: Relative survival

	N of observed deaths / N	Relative survival		95	% CI	_		95	% CI	
	of Expected deaths	at 10 years (%)	RER	lower	upper	P	RER*	lower	upper	P
HER2										
negative: 0/1+	712 / 536	84%	1 (Ref	erence)			1 (Ref	erence)		
2+	135 / 105	88%	0.91	0.46	1.82	0.796	0.78	0.45	1.35	0.373
3+	69 / 22	48%	4.75	3.10	7.28	<0.001	2.07	1.26	3.39	0.004
PIK3CA										
negative	344 / 250.6	82%	1 (Ref	erence)			1 (Ref	erence)		
positive	149 / 105	81%	0.94	0.50	1.76	0.855	1.24	0.78	2.01	0.194

^{*}adjusted for age, morphology, grade, hormone receptor status, tumor stage, type of breast surgery, type of axillary surgery, radiotherapy, endocrine therapy, chemotherapy

Relative Survival

Median follow-up was 8.9 years (range 0-17.0 years) for vital follow-up status. The results of the relative survival analyses are shown in Table 3. Patients with a HER-2 score of 2+, again showed an equal outcome as compared to the HER-2 negative patients (88% vs. 84% at 10 years; multivariable Relative Excess Risk (RER) 0.78, 95% CI 0.45-1.35; p=0.4). Patients with a HER-2 score of 3+ had a significantly worse relative survival at 10 years (48%; RER 2.07, 95% CI 1.26-3.39; p=0.004). Patients without PIK3CA mutations had a 10 years relative survival of 82%, as compared to 81% of patients with a PIK3CA mutation. There was no statistically significant difference in relative survival between patients with and without a PIK3CA mutation, also after stratifying for HER-2 status.

DISCUSSION

The main finding of our study is that HER-2 overexpression is of significant prognostic value in the older breast cancer population, even when taking the competing risk of mortality into account. Patients with HER-2 3+ tumours showed a significantly higher risk of recurrence, as compared to patients with a HER-2 negative tumour. Interestingly, patients with HER-2 2+ tumours had a similar recurrence risk as patients with HER-2 scores of 0 and 1+, who are considered to have HER-2 negative tumours. In this specific breast cancer population, PIK3CA mutations had no prognostic value.

With the results of this study, we identified a subgroup of older breast cancer patients with a significantly worse clinical outcome. Therefore, our results raise the hypothesis

that women aged 65 and older with HER2 3+ tumours might benefit from additional therapy, which includes anti-HER-2 therapy. In contrast to the study performed by Syed *et al.*, in which they claim a more indolent tumour character in the older breast cancer patient, we show that HER2+ positive tumours in the elderly are significantly associated with higher proliferative Ki67 presence in the tumour ⁴². Our results could justify more aggressive anti-cancer treatment in the older breast cancer patients harbouring tumours with disadvantageous characteristics, as they, similar to their presence in the younger breast cancer patients, are associated with worse survival rates.

It is essential to realize that in the FOCUS cohort, it is very unlikely that patients received anti-HER-2-therapy, as the FDA approved trastuzumab for the treatment of breast cancer in 2006, and patients included in the FOCUS cohort were diagnosed and treated for their breast cancer between 1997 and 2004 ⁴³. Moreover, the receipt of adjuvant chemotherapy was very low among the patients in our cohort, but is similar to previous observational studies in older breast cancer patients ^{44,45}.

An important question that still needs to be addressed is whether older breast cancer patients will benefit from anti-HER-2 therapy. Anti-HER-2 therapy is notorious for serious, mainly cardiac related, adverse events. However, recent studies have shown that cardiac adverse events from anti-HER-2 therapy are less severe and present with a lower incidence than initially assumed ^{21;46;47}. One of these studies was a side study of the HERA trial, in which the primary aim was to compare 1 versus 2 years of trastuzumab treatment. In this study patients with a median follow-up of 8-years had a relatively low incidence of cardiac events (cumulative incidence of confirmed LVEF decrease was 7% in the 2-year arm, 4% in the 1-year arm and, 1% in the observation arm). Moreover, the majority (87%) of HER-2 therapy induced cardiac events appeared to be reversible after stopping trastuzumab treatment ⁴⁷. Chavez-MacGregor *et al.* showed that trastuzumab-related cardiotoxicity did increase according to age. However, also in this older breast cancer population most of the cases were reversible, stressing the need for adequate monitoring ⁴⁸. It should also be noted that most (>80%) of the older patients who initiate trastuzumab complete this therapy ⁴⁹

In current medical practice, anti-HER-2 therapy is frequently omitted from treatment options in older breast cancer patients. This is probably due to the current standards that advise the combination with chemotherapy, but also due to the fear of cardiac toxicity. One of the major characteristics of the older cancer population is the clinical heterogeneity among patients of the same chronological age. Therefore, merely taking into account chronological age may result in unfair survival chances due to under-treatment of fit elderly. Given the results of our current study, it could be suggested that the fit elderly breast cancer patient, especially when the Left Ventricular Ejection Fraction (LVEF) is acceptable (more than 50%), should be treated with the same adjuvant-regimen as

the younger HER-2 positive breast cancer patients; consisting of a taxane based chemotherapy supplemented with one year of trastuzumab 41. Moreover, older patients with less desirable clinical conditions, or when there is a strong preference to omit chemotherapy, dual HER-2 blockage (trastuzumab combined with pertuzumab) could be considered. In the neoadjuvant setting, such chemotherapy-free regimens have shown to be associated with very few side effects, but unfortunately also with a lower tumour response 46. Currently, underrepresentation of elderly cancer patients in clinical trials hampers the implementation of widely agreed clinical practice recommendations, such as for HER-2 overexpressing breast cancer, which might result in under-treatment (suboptimal clinical outcome) or over-treatment (risk of significant toxicities) in the older breast cancer population. Currently, a randomized clinical trial is recruiting older (70+) patients with metastatic breast cancer disease, in which a chemotherapy-free regimen of trastuzumab and pertuzumab is compared with this combination and the addition of chemotherapy (EORTC-75111-10114) 50. We believe that clinical trials focusing on such specific therapies can change clinical practice for this specific breast cancer population in the coming years, as finally more insight is provided about an often neglected but increasingly important subgroup of the breast cancer population. It is for this reason that the results of this trial are eagerly awaited.

In our study, no clinically prognostic value was retrieved from PIK3CA mutation analyses in the elderly, HER-2 positive or HER-2 negative breast cancer patients. Currently, it is believed that PIK3CA mutations in the co-occurrence of HER-2 positive breast disease result in poor prognosis ⁵¹. In contrast, other studies have shown good prognosis with mutated PIK3CA in hormone receptor–positive tumors. The potential explanation of this finding is that continued activation of PI3K may have an inhibitory effect on HER-2 signalling ⁵¹. Based on the results from our study, PIK3CA mutations might not have any prognostic value in the elderly breast cancer patient.

The major strength of our study is that we used the largest consecutive series of older breast cancer patients from a population-based cohort, from which tumour material was available. Therefore, our study is not affected by selection bias. Our study is unique because most previous prognostic studies on HER-2 expression are performed in a younger breast cancer population or as part of a clinical trial. It has been shown that older patients who are registered in a clinical trial have a significantly better overall health than patients of the same age in the general population ⁵². Therefore, these studies cannot be simply extrapolated to the older breast cancer patients. This observation is in agreement with the American Cancer Society, whom states that older patients (≥65years) represent 45% of all breast cancer cases and are a particularly vulnerable, and underrepresented patient group ^{53,54}. Furthermore, recent articles by Chavez-MacGregor

et al., and Vaz-Luis et al. ^{49;53} point out that a relatively small portion of older breast cancer patients receive adjuvant chemotherapy (with/without trastuzumab therapy), which is attributed to undertreatment, a well described phenomenon among the elderly in the USA, and also in Europe. Herewith urging the need for better identification of older patients who could benefit from anti-HER2 therapy through real-world information instead of selective clinical trials.

Our present study demonstrates that HER-2 overexpression is a strong and independent predictor of worse prognosis, tested in a representative population of older breast cancer patients and even after taking into account competing risk of mortality due to other reasons. Therefore, from an under-treatment point of view, the results of this study should convince oncologists that anti-HER2 therapy should be at least considered for every HER-2 positive patient, regardless of age.

A limitation of the study is that there was no tumour material available for all patients in the cohort, mostly due to logistical reasons. We minimized the chance that this selection affected our study aims, by ruling out the association of having missing values on the determinants of interest (HER-2 amplification and PIK3CA mutation) with overall mortality. Second, in contrast to clinical practice, no confirmatory in situ hybridization was performed for the HER-2 2+ patients. However, based on the immunohistochemical data, the HER-2 2+ patients did not show a different clinical outcome compared to HER-2 negative patients. Based on these data, one could question the additional value of the costly in situ hybridization in this specific sub-group of the older breast cancer population, as this group does not have a worse prognosis than the HER-2 negative patients.

In conclusion, this population-based study among elderly breast cancer patients showed a strong prognostic effect of HER-2 overexpression, defined as an immunohistochemical score of 3+, on higher recurrence risk as well as worse relative survival. Herewith, we defined a subgroup of older breast cancer patients who are at high risk for worse clinical outcome, with a strong likelihood of under-treatment. Future research should point out whether it is possible to establish an effective anti-HER-2 regimen with minimal toxicity for the elderly breast cancer population.

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SUPPLEMENTARY TABLES

Supplementary table 1: primers used for PIK3CA mutation analysis

			,					
Assay Name	Primer Name	Primer Sequence	Reporter 1 Name	Dye	Dye Reporter 1 Sequence	Reporter 2 Name	Dye	Dye Reporter 2 Sequence
p.E542K*	PIK3CA_p.E542K_F PIK3CA_p.E542K_R	AGCTCAAAGCAATTTCTA- CACGAGAT GCACTTACCTGTGACTC- CATAGAAA	PIK3CA_p.E542K_V	VIC	PIK3CA_p.E542K_V VIC CCTCTCTGTGAAATCA	PIK3CA_p.E542K_M FAM CCTCTCTAAAATCA	FAM 6	CCTCTCTAAAATCA
p.E545K*	PIK3CA_p.E545K_F PIK3CA_p.E545K_R	TCAAAGCAATTTCTACAC-GAGATCCT GAGATCCT GCACTTACCTGTGACTC-CATAGAAA	PIK3CA_p.E545K_V VIC		CTCTCTGAAATCACT- GAGCAG	PIK3CA_p.E545K_M	FAM 6	FAM CTCTGAAATCACTAAGCAG
p.H1047R*	p.H1047R* PIK3CA_p.H1047R_F GCAAGAGG PIK3CA_p.H1047R_R TATTTCATG GCTGTTTAA GATCCAA	GCAAGAGCTTTGGAG- TATTTCATG GCTGTTTAATTGTGTGGAA- GATCCAA	PIK3CA_p. H1047R_V	VIC	CCACCATGATGTGCATC	PIK3CA_p. H1047R_M	FAM (FAM CACCATGACGTGCATC
Exon 9**	PIK3CA_x9_M13F PIK3CA_x9_M13R	TGTAAAACGACGGCCAGT- GGGAAAA TGACAAAGAACAGC CAGGAAACAGCTAT- GACCTCCATTTT AGCACTTACCTGTGAC						
Exon 20**	PIK3CA_x20_M13F PIK3CA_x20_M13R	TGTAAAACGACGGC- CAGTCTGAGCA AGAGGCTTTGGAG CAGGAAACAGCTAT- GACCCCTATGC						
* * *	PR_M13F	TGTAAAACGACGGCCAGT						
* * *	PR_M13R	CAGGAAACAGCTATGACC						

* Hydrolysis probes assays, ** Genomic PCR, *** Sanger sequencing, F=Forward primer, R=Reverse primer

Supplementary table 2: Relapse free period (Fine & Gray regression) and Relative survival, PIK3CA, stratified by HER-2

	10 N	Recurrence risk at 5 years	g T	95	12 % CI	q	***************************************	6	12% CI	q
	N OI recurrences	(%)	Z E	lower	upper	L	י אבוס	lower	upper	L
PIK3CA in HER-2	PIK3CA in HER-2 negative patients									
negative	72	15%	1 (Reference)	ence)			1 (Reference)			
positive	30	16%	1.00	0.68	1.48	0.987	1.10	0.73	1.65	0.653
PIK3CA in HER-2 2+	22+									
negative	15	16%	1 (Reference)	ence)			1 (Reference)			
positive	9	14%	0.86	0.38	1.97	0.737	0.92	0.35	2.44	0.863
PIK3CA in HER-2 3+	3+									
negative	16	35%	1 (Reference)	ence)			1 (Reference)			
positive	9	35%	1.05	0.40	2.77	0.921	n/a**			
	N of observed deaths / N of	Relative survival at 10 years		95	12 % CI			66	95% CI	
	expected deaths	(%)	RER	lower	upper	P	RER*	lower	upper	Ь
PIK3CA in HER-2	PIK3CA in HER-2 negative patients									
negative	261 / 198	84%	1 (Reference)	ence)			1 (Reference)			
positive	113/83	82%	0.89	0.38	2.09	0.795	1.14	0.56	2.32	0.710
PIK3CA in HER-2 2+	22+									
negative	46/37	%88	1 (Reference)	ence)			1 (Reference)			
positive	24/17	81%	0.82	0.16	4.16	0.813	n/a**			
PIK3CA in HER-2 3+	13+									
negative	31 / 13	%29	1 (Reference)	ence)			1 (Reference)			
positive	11/3	54%	1.46	0.55	3.85	0.443	n/a**			

*adjusted for age, morphology, grade, hormone receptor status, tumor stage, type of breast surgery, type of axillary surgery, radiotherapy, endocrine therapy, chemotherapy. **not applicable due to too few events

