

Basal markers and prognosis in luminal breast cancer

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Abstract

Purpose Basal marker expression in triple-negative breast cancers identifies basal-like tumours, and thus separates the TN group into two prognostic groups. However, the expression and prognostic significance of basal markers in luminal breast cancers are poorly described. The aim of this study was to investigate the expression and prognostic value of basal markers (CK5, CK14 and EGFR) in luminal breast cancer.

Methods A total of 1423 formalin-fixed, paraffin-embedded breast cancer tumours from a well-characterized cohort of Norwegian women, previously reclassified into molecular subtypes using IHC and ISH, were included in the study. For the present study, tumours expressing at least one of the basal markers CK5, CK14 or EGFR were defined as basal marker positive. Cumulative incidence of death from breast cancer and hazard ratio analyses were used to assess prognosis according to basal marker expression.

Results and conclusion In total, 470 cases (33.0%) were basal marker positive. A higher proportion of the basal marker-positive tumours were of histopathological grade 3 compared to basal marker negative. For hormone receptor-positive, HER2-negative cases, we found better prognosis for

basal marker-positive breast cancer compared to basal marker negative. For all subtypes combined, poorer prognosis for basal marker-negative cases was found in histopathological grade 2 tumours but not among grade 1 and 3.

Keywords Breast cancer · Luminal subtypes · Basal markers · Prognosis

Introduction

Classification of breast cancer into molecular subtypes based on gene expression patterns [23] has improved our understanding of the heterogeneity of breast cancer. Immunohistochemistry (IHC) and in situ hybridization (ISH) are often used as surrogates for gene expression analyses [5, 10, 27] and to a great extent reflect the genetic subtypes [6].

Luminal (oestrogen and/or progesterone receptor positive) tumours can be further divided into three subtypes based on their expression of Human epidermal growth factor receptor 2 (HER2) and expression of the proliferation marker Ki67. Non-luminal tumours can be classified into the HER2 subtype and the triple-negative (TN) subtype. Based on expression of basal markers, TN can be further subdivided into the basal phenotype and the non-basal or five-negative phenotype [5, 10, 15, 22]. Basal marker expression in TN breast cancers has been extensively studied. However, the prognostic value of basal markers in luminal breast cancer is still unresolved.

Luminal breast cancers are generally associated with a better prognosis compared to non-luminal. The Luminal A subtype has the best prognosis followed by the other luminal subtypes [5, 10, 11]. However, studies of long-term survival have shown that luminal breast cancers can recur

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up to decades after primary diagnosis. Luminal tumours are heterogeneous, and there is a need for further classification into prognostic groups in order to differentiate treatment and follow-up. The aim of this study was to investigate the expression and prognostic value of three basal markers [cytokeratin 5 (CK5), cytokeratin 14 (CK14) and epithelial growth factor receptor 1 (EGFR)] in luminal breast cancer tumours from a well-characterized cohort of Norwegian women with long-term follow-up.

Materials and methods

Study population

The study population has been described in detail previously [10, 14, 18, 32]. The patients are from two population-based surveys conducted in Nord-Trøndelag (NT) County in Norway, and the participants were born between 1886 and 1977. The women were followed for breast cancer occurrence, and, in total, formalin-fixed, paraffin-embedded (FFPE) tissue was available from 1423 cases. All cases were previously reclassified into molecular subtypes based on surrogate markers according to the algorithm shown in Fig. 1 [10, 31]. After linkage with the Cause of Death Registry of Norway and the Norwegian Cancer Registry, survival data were generated.

Specimen characteristics

Haematoxylin–erythrosine–safron (HES)-stained full-face sections from all tumours were already classified into histopathological type and grade according to current

guidelines [9, 19]. Methods for tissue microarray (TMA) construction, IHC and ISH are described elsewhere [10, 31]. Briefly, TMAs comprising three 1-mm-diameter tissue cores from all suitable cases were made and 4 µm sections were cut and stained. In total, 118 cases comprised only core biopsies or small tissue fragments unsuitable for TMA construction. From these, serial sections were made. IHC was done to assess ER, PR, Ki67, CK5, EGFR and HER2, and ISH was done for HER2 gene status. For the present study, IHC for CK14 was added. Two researchers assessed all IHC markers independently. Discordant results were discussed and consensus reached.

Assay methods

IHC for detection of CK14 was carried out according to the manufacturer's guidelines (Dako, Glostrup, Denmark). Sections were mounted on Superfrost + glass slides, dried at 37 °C overnight and stored at −20 °C. Slides were heated at 60 °C for 2 h and pretreated in a PT Link pre-treatment module for tissue specimens (Dako) with buffer (high pH target retrieval solution K8004) at 97 °C for 20 min. before staining. Mouse monoclonal antibody CK14 was applied (clone LL002, concentration 36 mg/l, dilution 1:20). Dako REAL™ EnVision™ detection system was used with peroxidase/diaminobenzidine (DAB)+, rabbit/mouse, code K5007, for visualization.

Scoring and reporting

Assessment and cut-off levels for all markers except CK14 have been explained previously [10, 32]. A staining index (SI) for CK14 was calculated as staining intensity

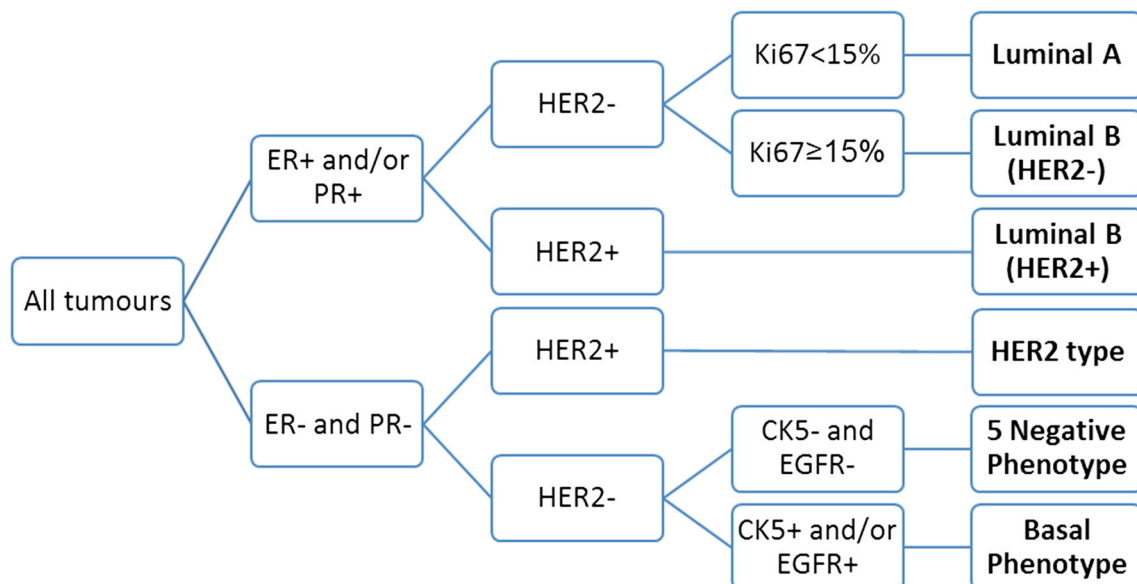


Fig. 1 Algorithm for molecular subtyping based on surrogate markers (Engstrom et al. [10])

Table 1 Clinicopathological characteristics of the 1423 cases in the study

	Basal marker positive (%)	Basal marker negative (%)	Total (%)
Number (%)	470 (33.0)	953 (67.0)	1423 (100)
Number of breast cancer deaths (%)	126 (26.8)	298 (31.3)	424 (29.8)
Number of deaths all causes (%)	250 (53.2)	635 (66.3)	885 (62.2)
Mean age at diagnosis (SD)	62.7 (12.8)	69.6 (12.2)	67.3 (12.8)
Median years of follow-up after diagnosis (IQR)	7.0 (8.9)	7.5 (8.4)	7.4 (8.5)
Histopathological type (%)			
NST (invasive carcinoma of no special type)	378 (80.3)	671 (70.4)	1049 (73.7)
ILC (invasive lobular carcinoma)	22 (4.7)	154 (16.2)	176 (12.4)
Other	70 (14.9)	128 (13.4)	198 (13.9)
Histopathological grade (%)			
Grade 1	83 (17.7)	140 (14.7)	223 (15.7)
Grade 2	179 (38.1)	564 (59.2)	743 (52.2)
Grade 3	208 (44.2)	249 (26.1)	457 (32.1)
Tumour size (%)			
≤2 cm	262 (55.7)	472 (49.5)	734 (51.6)
>2 cm, ≤5 cm	94 (20.0)	169 (17.7)	263 (18.4)
>5 cm	11 (2.3)	27 (2.8)	38 (2.7)
Uncertain, but ≥2 cm	48 (10.2)	97 (10.2)	145 (10.2)
Uncertain	55 (11.7)	188 (19.7)	243 (17.1)
Lymph node status (%)			
No metastasis	228 (48.5)	428 (44.9)	656 (46.1)
Metastasis detected	178 (37.9)	295 (30.1)	473 (33.2)
Not examined for metastasis	64 (13.6)	230 (24.1)	294 (20.7)
Stage *			
I	226 (48.1)	502 (52.7)	728 (51.2)
II	206 (43.8)	347 (36.4)	553 (38.9)
III	24 (5.1)	54 (5.7)	78 (5.5)
IV	14 (3.0)	44 (4.6)	58 (4.1)
Unknown	0	6 (0.6)	6 (0.4)

* Cancer Registry of Norway. Combined clinical and histological stage

Table 2 Basal marker status according to molecular subtypes

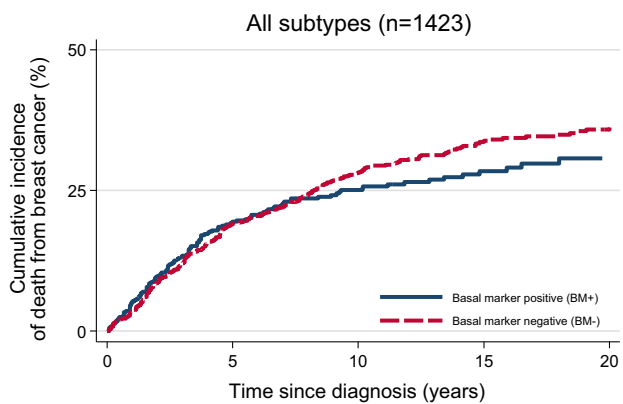
Luminal and non-luminal subtypes	Molecular subtypes	Total	Basal marker positive (%)	Basal marker negative (%)
Luminal (<i>n</i> = 1194)				
	Luminal A	705 (49.5)	181 (25.7)	524 (74.3)
	Luminal B (HER2–)	377 (26.5)	108 (28.7)	269 (71.4)
	Luminal B (HER2+)	112 (7.9)	35 (31.3)	77 (68.8)
Non-luminal (<i>n</i> = 229)				
	HER2 subtype	89 (6.3)	51 (57.3)	38 (42.7)
	Basal phenotype	92 (6.5)	92 (100.0)	0 (0)
	5 negative phenotype*	48 (3.4)	3 (6.3)	45 (93.7)
Total		1423 (100)	470 (33.0)	953 (67.0)

* Negative for ER, PR, HER2, CK5 and EGFR

Table 3 Number of basal marker-positive cases separately and in combination

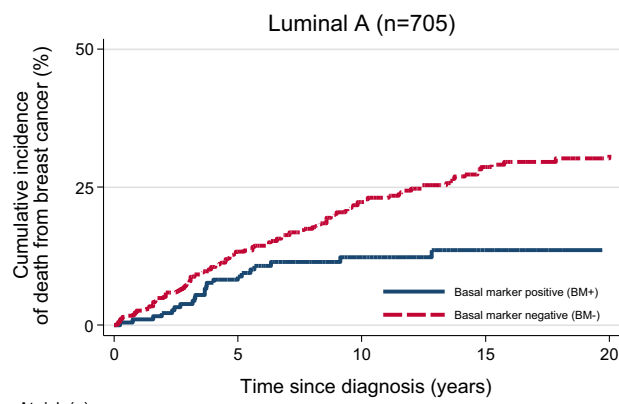
Basal markers	Number of positive (%)
Basal marker positive	470 (33.0)
CK5	350 (24.6)
CK14	332 (23.3)
EGFR	78 (5.5)
All three	37 (2.6)
One marker positive	214 (15.0)

multiplied by the proportion of cells showing positive staining. Cytoplasmic staining intensity was graded as 0 (no staining), 1 (weak), 2 (moderate) and 3 (strong). The proportion of positively stained cells was scored as 1



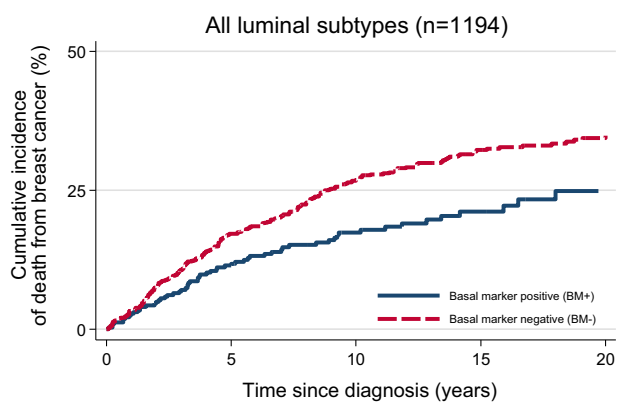
At risk (n)					
BM+	470	355	237	121	59
BM-	953	731	489	291	186

(a) Gray’s test: $p=0.26$



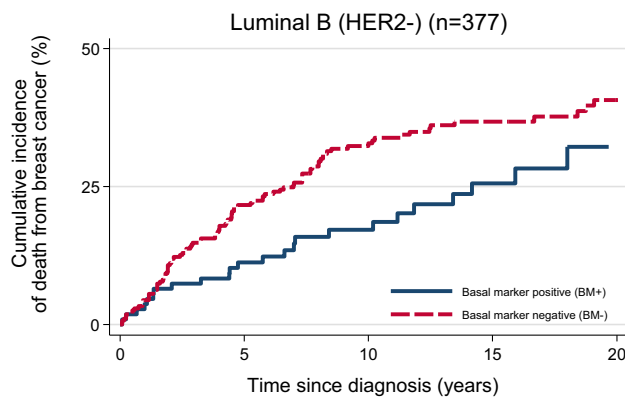
At risk (n)					
BM+	181	153	103	42	19
BM-	524	426	283	161	103

(a) Gray’s test: $p=0.003$



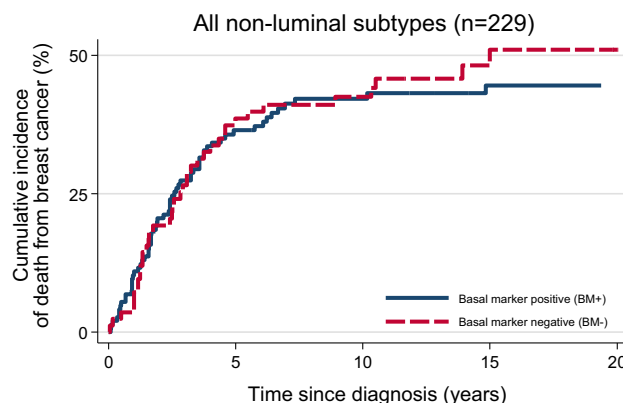
At risk (n)					
BM+	324	268	178	86	38
BM-	870	680	449	272	174

(b) Gray’s test: $p=0.003$



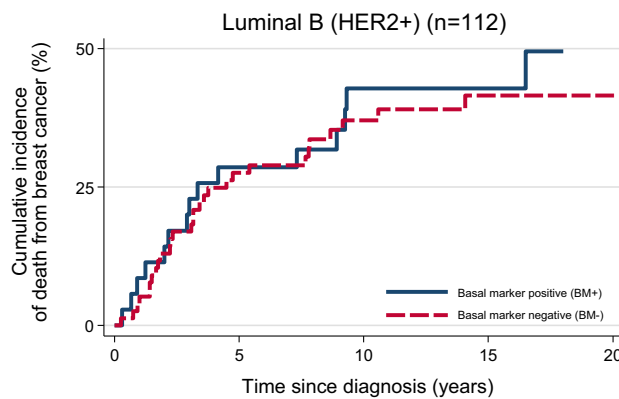
At risk (n)					
BM+	108	90	61	35	13
BM-	269	201	134	88	54

(b) Gray’s test: $p=0.04$



At risk (n)					
BM+	146	86	59	36	21
BM-	83	51	40	18	11

(c) Gray’s test: $p=0.69$



At risk (n)					
BM+	35	25	14	9	6
BM-	77	53	33	23	18

(c) Gray’s test: $p=0.79$

Fig. 2 a–c Cumulative incidence of death from breast cancer according to basal marker status. Gray’s test: **a** $p = 0.26$, **b** $p = 0.003$, **c** $p = 0.69$

(<10%), 2 (10–50%) and 3 (>50%). Cases with SI 0–1 were considered negative and cases with SI 2–9 were regarded as positive. Basal marker-positive (BM⁺) cases

Fig. 3 a–c Cumulative incidence of death from breast cancer according to basal marker status for the three luminal subtypes separately. Gray’s test: **a** $p = 0.003$, **b** $p = 0.04$, **c** $p = 0.79$

were defined as positive for at least one of the basal markers CK5, CK14 and EGFR. The REMARK recommendations for tumour marker studies were followed [20].

Table 4 Cumulative risk of death from breast cancer according to basal marker status

	5 years after diagnosis			15 years after diagnosis	
	Total (<i>n</i>)	Breast cancer deaths (<i>n</i>)	Cumulative risk of death (%) (95% CI)	Breast cancer deaths (<i>n</i>)	Cumulative risk of death (%) (95% CI)
All subtypes BM ⁺	470	90	19.2 (15.9–23.1)	120	28.4 (24.2–33.2)
All subtypes BM [−]	953	180	19.1 (16.7–21.7)	283	33.6 (30.4–37.1)
Luminal subtypes BM ⁺	324	37	11.5 (8.4–15.5)	48	21.2 (16.6–26.9)
Luminal subtypes BM [−]	870	148	17.2 (14.8–19.9)	245	32.3 (28.9–35.9)
Non-luminal subtypes BM ⁺	146	53	36.5 (29.2–44.8)	62	44.6 (36.5–53.5)
Non-luminal subtypes BM [−]	83	32	38.6 (29.1–49.9)	38	48.2 (37.4–60.2)

BM basal markers

Statistical analyses

All women were followed for breast cancer occurrence and censored at the time of death or emigration. End of follow-up was December 31st, 2010 for the oldest cohort and December 31st, 2013 for the youngest. Cumulative incidence of breast cancer death was calculated, and deaths from other causes were treated as competing events. Cumulative incidence curves were compared using Gray's test. Cox proportional hazards models were used to compare the risk of death from breast cancer according to BM status, calculating hazard ratios (HR) with 95% confidence intervals (CI). Adjustments were made for age; stage (in five categories: stage I–IV and unknown) at diagnosis according the data from the Cancer Registry [13]; and histopathological grade. Stata version 13.1 (Stata Corp., College Station, TX, USA) was used for the statistical analyses.

Ethics

The study was approved by the Regional Committee for Medical and Health Sciences Research Ethics, and dispensation from the requirement of patient consent was granted (REK, Midt-Norge, ref. nr: 836/2009).

Results

Descriptive statistics

Table 1 shows the clinicopathological characteristics of all cases. Mean age at diagnosis was 67.3 years. Women with BM⁺ tumours were younger than women with BM[−] tumours (62.7 vs. 69.6 years of age). BM⁺ tumours were more often of histopathological grade 3 (44.2 vs. 26.1%). The proportion of invasive lobular carcinoma was lower for BM⁺ cases (4.7 vs. 16.2%). Of the 1423 cases included

in the study, 1194 were luminal and 229 were non-luminal (Table 2). Of the luminal cases, 324 (27.1%) were BM⁺. Table 3 shows the number of basal marker-positive cases for each marker and in combination.

Survival analyses

Figure 2a–c and Table 4 show cumulative incidence of death from breast cancer according to BM status for all breast cancer cases and for luminal and non-luminal cases separately. For all cases, the curves overlap the first eight years (Fig. 2a), and the cumulative risk of death after 5 years of follow-up was 19.2% (95% CI 15.9–23.1) for BM⁺ and 19.1% (95% CI 16.7–21.7) for BM[−]. However, the corresponding risks of death after 15 years of follow-up were 28.4% (95% CI 24.2–33.2) for BM⁺ and 33.6% (95% CI 30.4–37.1) for BM[−].

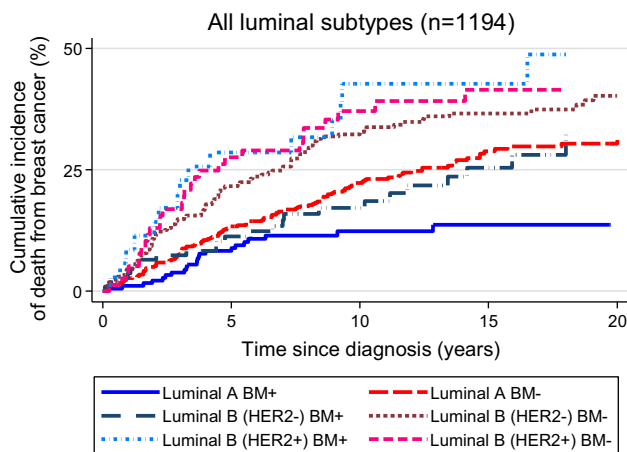
BM[−] luminal tumours had a poorer prognosis than BM⁺ luminal tumours (Gray's test: $p = 0.003$) (Fig. 2b). The cumulative risk of death from BM⁺ luminal breast cancer was 11.5% (95% CI 8.4–15.5) after 5 years of follow-up and 21.2% (95% CI 16.6–26.9) after 15 years. The corresponding risks for BM[−] luminal cases were 17.2% (95% CI 14.8–19.9) and 32.3% (95% CI 28.9–35.9), respectively. For all non-luminal cases combined, no difference in prognosis was seen according to BM status (Gray's test: $p = 0.69$) (Fig. 2c).

Figure 3a–c and Table 5 show cumulative incidence of death from breast cancer according to BM status for all three luminal subtypes separately. For Luminal A and Luminal B (HER2[−]), BM⁺ cases had better prognosis compared to BM[−] (Gray's test: $p = 0.003$ and $p = 0.04$, respectively) (Fig. 3a–b). For Luminal B (HER2⁺), no difference was seen according to BM status (Gray's test: $p = 0.79$) (Fig. 3c).

Figure 4 is a summary and relative comparison of Figs. 3a–c showing cumulative risks of death from breast cancer for all luminal subtypes according to BM status. There was no clear difference in prognosis for Luminal A

Table 5 Cumulative risk of death from breast cancer for luminal subtypes according to basal marker status

	Total (<i>n</i>)	5 years after diagnosis		15 years after diagnosis	
		Breast cancer deaths (<i>n</i>)	Cumulative risk of death (%) (95% CI)	Breast cancer deaths (<i>n</i>)	Cumulative risk of death (%) (95% CI)
Luminal A BM+	181	15	8.3 (5.1–13.4)	22	13.6 (9.0–20.2)
Luminal A BM–	524	69	13.4 (10.7–16.6)	125	28.6 (24.4–33.4)
Luminal B (HER2–) BM+	108	12	11.3 (6.6–19.1)	22	25.4 (17.2–36.7)
Luminal B (HER2–) BM–	269	58	21.7 (17.2–27.1)	91	36.6 (30.9–43.1)
Luminal B (HER2+) BM+	35	10	28.6 (16.5–46.6)	14	42.7 (27.6–61.6)
Luminal B (HER2+) BM–	77	21	27.6 (18.9–39.1)	29	41.5 (30.6–54.5)

**Fig. 4** Cumulative incidence of death from breast cancer according to basal marker status for all three luminal subtypes. Gray's test: $p < 0.0001$

BM[–] compared to Luminal B (HER2[–]) BM⁺. The cumulative risk of death from Luminal A BM[–] breast cancer was 13.4% (95% CI 10.7–16.6) and 28.6% (95% CI 24.4–33.4) after 5 and 15 years of follow-up, respectively. The corresponding cumulative risk of death for Luminal B (HER2[–]) BM⁺ was 11.3% (95% CI 6.6–19.1) and 25.4% (95% CI 17.2–36.7).

Figures 5 a–c and Table 6 show results from survival analyses for BM⁺ and BM[–] for histopathological grades 1, 2 and 3 separately. For grade 2 tumours, the prognosis was significantly poorer for BM[–] compared to BM⁺ (Gray's test: $p = 0.0004$) (Fig. 5b). For histopathological grades 1 and 3 separately, we found no clear differences in prognosis according to BM status.

Cox regression analysis was done to compare rates of breast cancer death according to BM status for all cases combined, for all luminal cases combined and for each luminal subtype separately (Table 7). Comparing BM⁺ and BM[–] for all breast cancer cases irrespective of subtypes, there was a tendency toward better prognosis for BM⁺. However, there was no difference after adjusting for age. The first ten years after diagnosis, there were no clear

differences in prognosis, but from ten years and onwards, BM[–] had poorer prognosis compared to BM⁺ [hazard ratio (HR) 1.98 (95% CI 1.09–3.63)] (Table 7). For all luminal cases combined, BM⁺ had better prognosis compared to BM[–] [HR 1.71 (95% CI 1.30–2.25)]. Luminal A BM[–] had higher risk of death from breast cancer compared to Luminal A BM⁺ [HR 2.16 (95% CI 1.39–3.33)]. Corresponding results were found for Luminal B (HER2[–]). However, HRs were attenuated after adjustment for age, stage and histopathological grade, where age had the greatest attenuating effect.

Discussion

We have studied the expression and prognostic impact of basal markers CK5, CK14 and EGFR in a well-characterized cohort of women with breast cancer. The cohort comprises 1423 breast cancer patients with long-term follow-up. We have studied risk of death from breast cancer according to BM status for all breast cancer cases combined, for all luminal cases combined and for each luminal subtype separately. We found better prognosis for BM-positive luminal breast cancer compared to BM-negative. This result was restricted to the Luminal A and Luminal B (HER2[–]) subtypes. For Luminal B (HER2⁺) tumours, we found no clear differences according to BM status.

HRs were weakened after adjustment for known prognostic markers (Table 7). Adjusting for age had the greatest influence on the results. Nevertheless, the tendency toward better prognosis for BM⁺ cases compared to BM[–] remained. When adjusting for stage and histopathological grade, only minor changes were seen. The mean age of the women with BM⁺ tumours was lower compared to the women with BM[–] tumours (62.7 vs. 69.6 years of age). The time of diagnosis in this cohort spans over several decades and treatment would have been largely determined by age at diagnosis and time of diagnosis. A high proportion of women in our cohort received no therapy after surgery, but new treatment modalities were gradually

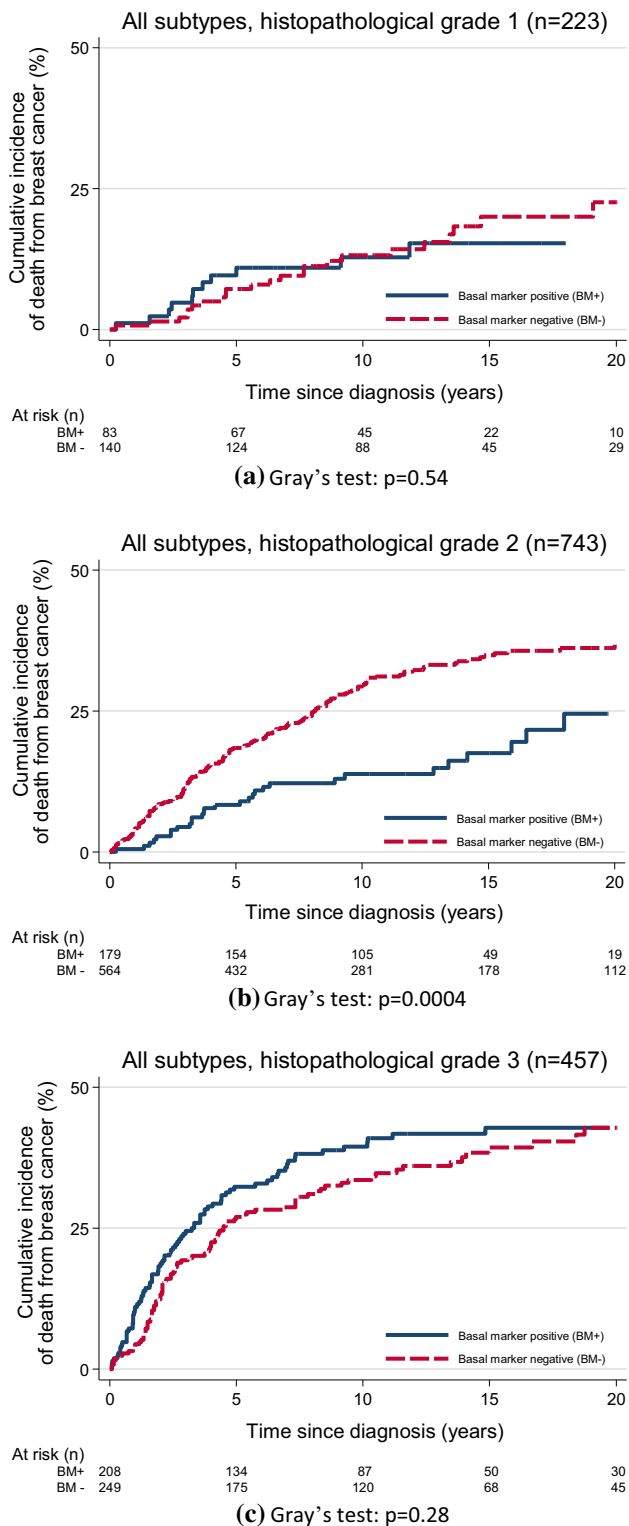


Fig. 5 a–c Cumulative incidence of death from breast cancer for histopathological grade 1, 2 and 3 according to basal marker status. Gray's test: **a** $p = 0.54$, **b** $p = 0.0004$, **c** $p = 0.28$

introduced in the latter part of the study period. Previously, our group has shown improved prognoses for Luminal A, Luminal B (HER2⁻) and Basal phenotype diagnosed after

1995 compared to cases diagnosed before 1995 [31], probably partly due to the implementation of modern treatment protocols. These changes over time may have influenced the results.

In the present study, basal marker expression had prognostic value for histopathological grade 2 tumours but not for grade 1 and 3. Grade 2 tumours represent a heterogeneous group with diverging risk of recurrence and breast cancer death [8, 10, 25]. Therefore, further subclassification of grade 2 tumours is of interest. However, even in a large study like this, some groups were too small for further subclassification, and in some of the results presented here confidence intervals were broad, probably due to low numbers of cases.

The expression and prognostic impact of basal markers in luminal breast cancer have not been widely studied. Previous studies by Blows et al. [5] and Sung et al. [27] had diverging results regarding prognosis compared to ours. In order to compare with Blows et al., we analysed prognosis based on overall survival (data not shown), but the results did not change. Sung et al. found that basal-positive [(CK5+ or CK5/6+) and/or EGFR+] Luminal A tumours were smaller, had lower histopathological grade and more often tended to be lymph node-negative compared to basal-negative tumours. For all cancer subtypes combined, we found a higher proportion of histopathological grade 3 tumours among BM+ compared to BM- (Table 1). However, within the Luminal A subtype, we found similar results to Sung et al. with regard to tumour size, histopathological grade and lymph node status (data not shown).

In addition to the high number of cases and long follow-up, a further strength of the present study is that all laboratory tests were carried out in the same laboratory, with the same technical and scientific personnel, using the same IHC markers and algorithm in all cases. Histopathological diagnoses were revised, and assessment of IHC markers was performed by two researchers independently. However, in a cohort such as this, preanalytical conditions may have varied, possibly influencing the results. Furthermore, as discussed above, the patients in the study may have received differing adjuvant treatment, if any, due to time period and age at diagnosis.

Basal markers are often focally expressed in breast cancer [17, 30]. TMA is a high throughput method enabling analyses of tumour tissue from a high number of cases under the same conditions [6, 16, 24]. However, the method may have limitations regarding representativity and false-negative cases cannot be excluded. In addition, consensus-based cut-off levels for basal markers are not established. The results must be interpreted in light of these factors.

In previous studies of prognosis for molecular subtypes based on surrogate markers, we found a poorer prognosis

Table 6 Cumulative risk of death from breast cancer for histopathological grade 1, 2 and 3 according to basal marker status

	Total (<i>n</i>)	5 years after diagnosis		15 years after diagnosis	
		Breast cancer deaths (<i>n</i>)	Cumulative risk of death (%) (95% CI)	Breast cancer deaths (<i>n</i>)	Cumulative risk of death (%) (95% CI)
Histopathological grade 1 BM+	83	8	9.6 (4.9–18.4)	11	15.4 (8.6–26.7)
Histopathological grade 1 BM–	140	10	7.2 (4.0–13.0)	22	20.1 (13.4–29.5)
Histopathological grade 2 BM+	179	15	8.4 (5.1–13.5)	26	17.5 (12.0–25.3)
Histopathological grade 2 BM–	564	103	18.5 (15.5–22.0)	173	34.9 (30.7–39.5)
Histopathological grade 3 BM+	208	67	32.4 (26.4–39.2)	83	42.8 (35.9–50.4)
Histopathological grade 3 BM–	249	67	27.0 (21.9–33.0)	88	38.4 (32.2–45.4)

Table 7 Risk of death from breast cancer according to basal marker status

	Number of cases	Deaths from breast cancer	Unadjusted		Adjusted for age		Adjusted for stage		Adjusted for histopathological grade		Adjusted for age, stage and grade	
			HR	95% CI	HR	95% CI	HR	95% CI	HR	95% CI	HR	95% CI
All subtypes												
BM ⁺	470	126	1.00	1.00–1.51	1.00	0.80–1.24	1.00		1.00		1.00	
BM [–]	953	298	1.23		1.00		1.22	0.99–1.51	1.30	1.06–1.61	1.10	0.88–1.38
	1423	424										
All subtypes, follow-up first ten years after diagnosis												
BM ⁺	470	112	1.00	0.91–1.42	1.00	0.74–1.17	1.00	0.88–1.39	1.00	0.97–1.52	1.00	
BM [–]	953	253	1.14		0.93		1.11		1.21		1.02	0.80–1.29
	1423	365										
All subtypes, follow-up from ten years after diagnosis ^a												
BM ⁺	186	14	1.00	1.09–3.63	1.00	0.83–2.93	1.00	1.26–4.39	1.00	1.08–3.67	1.00	0.89–3.31
BM [–]	318	45	1.98		1.56		2.35		1.99		1.71	
	504	59										
All luminal subtypes												
BM ⁺	324	64	1.00	1.30–2.25	1.00	1.00–1.78	1.00	1.27–2.20	1.00	1.29–2.23	1.00	0.95–1.71
BM [–]	870	259	1.71		1.34		1.67		1.69		1.28	
	1194	323										
Luminal A												
BM ⁺	181	24	1.00	1.39–3.33	1.00	0.95–2.37	1.00	1.32–3.16	1.00	1.33–3.19	1.00	0.88–2.20
BM [–]	524	132	2.16		1.50		2.04		2.06		1.39	
	705	156										
Luminal B (HER2 [–])												
BM ⁺	108	25	1.00	1.16–2.80	1.00	0.97–2.45	1.00	1.19–2.94	1.00	1.18–2.87	1.00	0.93–2.42
BM [–]	269	96	1.80		1.54		1.87		1.84		1.50	
	377	121										
Luminal B (HER2 ⁺)												
BM ⁺	35	15	1.00	0.53–1.83	1.00	0.39–1.59	1.00	0.43–1.65	1.00	0.50–1.73	1.00	0.32–1.35
BM [–]	77	31	0.98		0.79		0.84		0.93		0.66	
	112	46										

HR Hazard ratio, CI Confidence interval

^a Conditional on surviving the first 10 years

for HER2⁺ and TN subtypes the first five years after diagnosis. For those who survived five years, the prognosis did not differ between the subtypes [10, 32]. We also found a steadily declining survival for the luminal HER2⁻ subtypes throughout the follow-up period. Also in the present study, the period of time from diagnosis has impact on prognosis. As shown in Fig. 2a, the prognostic effect of basal markers may change with time from breast cancer diagnosis. Tischkowitz et al. showed that the prognostic value of different immunohistochemical markers varied with time after diagnosis and that after 10 years, basal markers might be of higher prognostic value than hormone receptors and HER2 [29]. This, along with the results in the present study, confirms that long-term follow-up is necessary to discover prognostic differences in breast cancer.

A number of studies have shown poorer prognosis for BM + TN breast cancer [5, 7, 29]. There are few published studies on the prognostic impact of BM in luminal breast cancer, but our results indicate a better prognosis for BM⁺ HER2⁻ luminal tumours. Different prognostic markers and combinations of markers may have differing impact in the various subtypes of breast cancer. The expression of different sets of markers may reflect different routes of cell differentiation [21]. Interaction between, and mutual impact of the various prognostic markers remain to be elucidated. The prognostic impact of basal markers may differ depending on the presence or absence of hormone receptors or HER2 status.

Basal breast cancer is not clearly defined and a number of basal markers have been used to define this subtype [1, 3, 30, 33]. CKs are cytoplasmic proteins important for the cytoskeleton of eukaryotic cells and are, to a certain extent, differentially expressed in basal and luminal epithelial cells [26]. EGFR is a tyrosine kinase receptor, and activation plays a central role in regulation of proliferation, growth and cell survival. Overexpression of EGFR in epithelial tumours may lead to aggressive growth and invasion, and ultimately, a poorer prognosis [12, 28]. CK5 or CK5/6 are among the most widely used basal markers [3, 5]. EGFR is upregulated in the basal-like subtype of breast cancer [23] and is considered useful in combination with basal markers [22]. We chose CK5 instead of CK5/6 to identify basal-like phenotypes due to higher sensitivity [4]. In our material, there were few EGFR-positive cases compared to other studies [28] and therefore its value was limited. We found that three cases in the non-luminal subtype, 5NP, expressed CK14. Among luminal tumours, CK14 was the only positive basal marker in 88 cases. Other basal markers such as CK17, vimentin, nestin and p63 [1, 2] could have increased the number of tumours with basal traits in this material.

In conclusion, we found better prognosis for basal marker-positive breast cancer (defined as positive for CK5

and/or CK14 and/or EGFR) compared to basal marker-negative cases. This finding was restricted to hormone receptor-positive, HER2-negative cases. For all subtypes combined, we found better prognosis for basal marker-positive cases in histopathological grade 2 tumours but not in grades 1 and 3. Time from diagnosis may be crucial for the prognostic value of basal markers and in the present study, the differences were most apparent from ten years after diagnosis. There are few studies of the prognostic value of basal markers in luminal subtypes. The results are diverging and further studies are necessary.

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Authors' contribution MJE contributed to conception and design of the study, interpretation of the IHC markers, carried out statistical analyses, interpretation of the results and drafted the manuscript. MV contributed to discussion and interpretation of the results, interpretation of the IHC markers, review and revision of the manuscript. AMB contributed to discussion and interpretation of the results, interpretation of the IHC markers, review and revision of the manuscript. All authors read and approved the final manuscript.

Compliance with ethical standards

Conflict of interest The authors declare that they have no competing interests.

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