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Comparison of primary breast cancer and paired metastases: biomarkers discordance influence on outcome and therapy

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Research Article

Comparison of primary breast cancer and paired metastases: biomarkers discordance influence on outcome and therapy

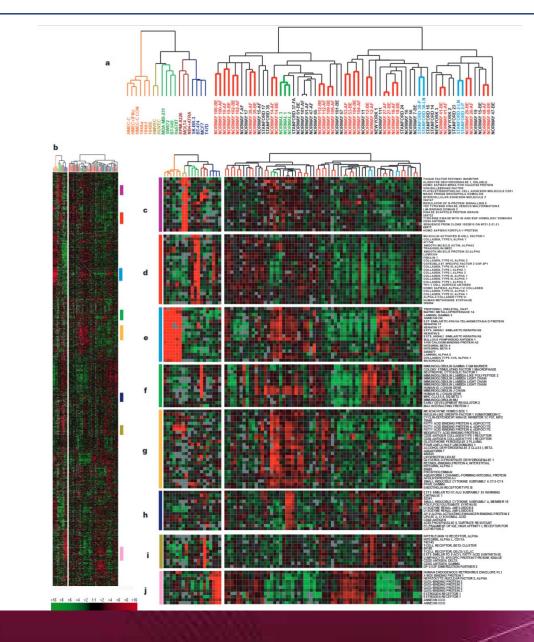
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Background

The high heterogeneity of breast cancer led to the definition of different molecular subtypes (luminal A and B, HER2-enriched and triple negative) with distinctive biological features and clinical outcomes

→ immunohistochemical-defined surrogates based on the assessment of ER, PR, HER2 and Ki67 expression



Perou et al. Nature 2000

Background

• It is commonly thought that distant metastases resemble the same profile of the primary tumor, systemic palliative therapy is usually assigned according to the primitive biological phenotype

• Discordance rates for ER, PR and HER2 expression between primary BC and paired metastases are described, suggesting both biological and therapeutic implications

Aurilio et al. EJC 2014

Metastasis biopsy is not strictly required, but NCCN guidelines recommend the re-biopsy
of metastatic sites when feasible

Aim

 To evaluate the discordance in ER, PR, HER2 and Ki67 expression in a retrospective, real-world series of metastatic breast cancer

 To investigate the possible impact of antineoplastic treatments on biomarkers expression with an exploratory hypothesis generating purpose

Methods

- From January 2004 to July 2013 we retrospectively analyzed a cohort of 544 patients with metastatic breast cancer receiving anticancer treatments at the Department of Oncology, University Hospital of Udine, Italy.
- We selected patients that underwent biopsy of local or distant recurrence with available archival ER, PR, HER2 and Ki67 data.
- Biopsies from both primary and metastatic sites were analyzed at the Pathology Department of the University Hospital of Udine.

Statistical analysis

- Descriptive analysis of clinical and pathological characteristics was performed.
 - ER and PR variations were defined according to a 1% cut off, while for Ki67 a cut off of 14, 20 and 30% was applied.
 - Modifications of both hormonal receptors and Ki67 were analyzed also on a percent basis.
- To estimate the association between received therapies and ER, PR and Ki67 changes, OR with the 95% CI were calculated through uni- and multivariate logistic regression.
- Prognostic impact of changes in ER, PR and Ki67 in terms of OS was investigated through uni- and multivariate Cox regression and described by means of Kaplan–Meier estimator plot.

Baseline features

Table 1. Population's main char	acteristics.		
Baseline features	Number	%	
Age at stage IV diagnosis:			
-<35	2	0.8	
- 35–65	137	59.1	
- 65-70	Table 1. Population's ma	ain characteristics	
->70	Baseline features	Number	%
Grading:	— pN:		
- G1	- N0	56	24.1
– G2	– N1mi	2	0.9
- G3	- N1	44	19.0
– Gx	- N2	28	12.1
pT:			16.4
- T0	- N3	38	
- T1	- Nx	64	27.5
– T2	Stage IV at diagnosis:		
– T3	– Yes	42	18.0
- T4	– No	164	70.7
-Tx	– Unknown	24	10.3
_ TA	Previous treatments:		
	– Anthracyclines	117	50.4
	– laxanes	/8	33.6
	– Antiestrogens	114	49.1
	– Aromatase inhibitors	91	39.2

Molecular characteristics

	Primary Biopsy		Secondary Biopsy		
	Number	%	Number	%	
Tumor Phenotype	166		178		
Luminal A	26	15.66	10	5.62	
Luminal B	88	53.01	100	56.18	
Luminal HER2	16	9.64	16	8.99	
HER2 Positive	17	10.24	19	10.67	
Triple Negative (TNBC)	19	11.45	33	18.54	

Molecular discordance

	Number	%	p value
ER (n=166)			
Discordant	21	12.65	
Gain	4	10.25	p = 0.0072
Loss	17	13.38	
ER continuous (n=127)			
Lower	58	45.67	p = 0.2437
Higher or Stable	69	ρ – 0.24. 59	
PgR (n=171)			
Discordant	85	49.71	
Gain	13	20.63	<i>p</i> < 0.0001
Loss	72	66.67	
PgR continuous (n=108)			
Lower	88	81.48	p < 0.0001
Higher or Stable	20	18.52	p (5.5551

HER2 (n=175)			
Discordant	6	3.43	
Gain	4	2.79	p = 0.6875
Loss	2	6.25	
Ki67 >14 (n=120)			
Discordant	39	32.5	
<14 to ≥14	31	73.80	p = 0.0003
≥14 to <14	8	10.25	
Ki67 ≥20 (n=120)			
Discordant	42	35.0	
<20 to ≥20	35	67.30	p < 0.0001
≥20 to <20	7	10.29	
Ki67 ≥30 (n=120)			
Discordant	45	37.5	
<30 to ≥30	38	52.77	p < 0.0001
≥30 to <30	7	14.58	-

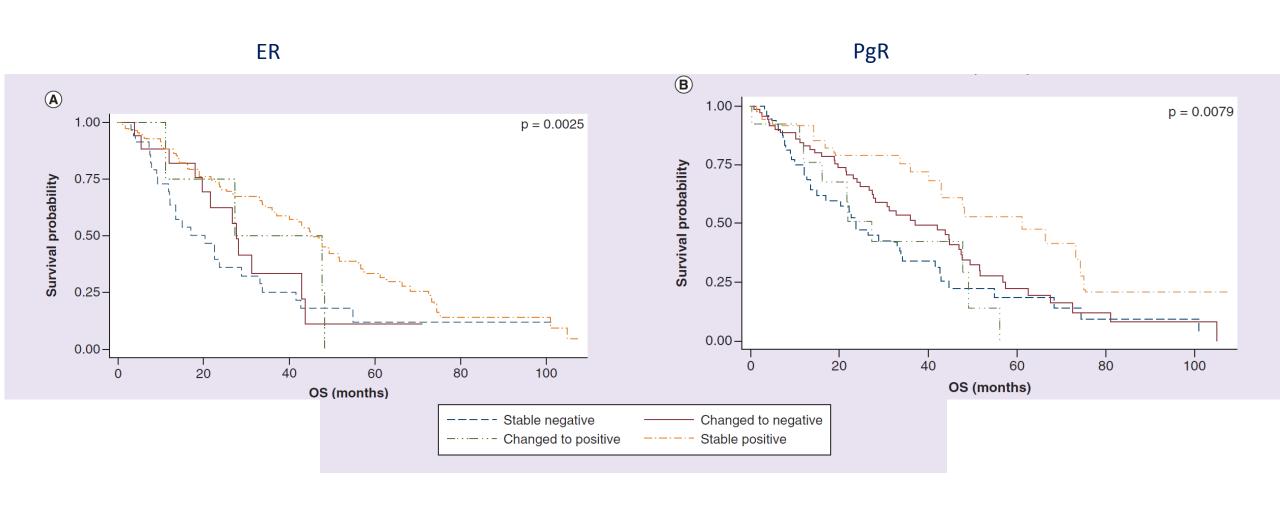
Molecular discordance

Tumor phenotype (n=126)				
Discordant	28	22.2		
Luminal A to B	7	63.6		
Luminal B to A	2	2.89		
Luminal A/B to TNBC	11	13.75		
Luminal A/B to HER2 positive	1	1.25		
Luminal A/B to Luminal HER2	2	2.50		
Luminal HER2 to Luminal A/B	1	7.69		
Luminal HER2 to HER2 positive	3	23.07		
HER2 positive to Luminal HER2	1	7.14		

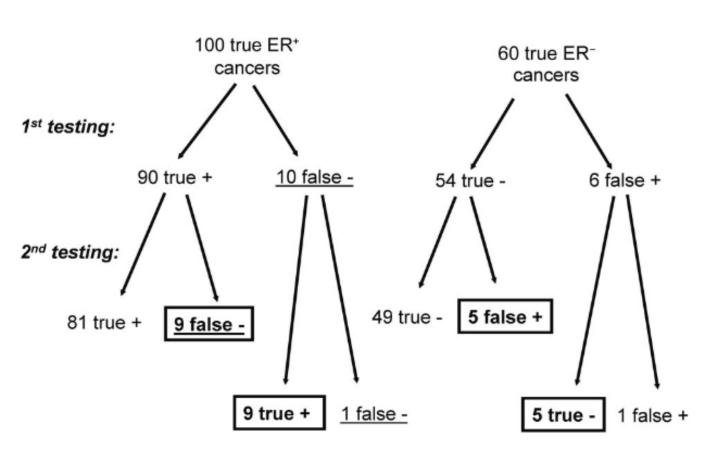
Uni- and multivariate analyses

Phenotype modification	Anthracyclines [‡]		Taxa	Taxanes [§]		Antiestrogens [¶]		Aromatase inhibitors#	
	OR (95% CI)	p-value	OR (95% CI)	p-value	OR (95% CI)	p-value	OR (95% CI)	p-value	
ER loss: – Univariate – Multivariate	1.85 (0.61–5.63) –	ns	1.43 (0.51–4.01)	ns	0.81 (0.28–2.31) –	ns	3.02 (0.92–9.84) –	ns	
ER reduction (min. 1%): – Univariate – Multivariate	1.33 (0.65–2.71) –	ns	4.03 (1.89–8.58) 3.59 (1.66–7.77)	<0.001 0.001	0.81 (0.39–1.67) –	ns	2.51 (1.21–5.19) 2.07 (0.96–4.44)	0.013 ns	
PR loss: – Univariate	6.47 (2.68–15.61)	<0.001	3.7 (1.48–9.2)	0.005	1.14 (0.47–2.74)	ns	1.85 (0.82–4.16)	ns	
 Multivariate 	5.20	0.02	1.52	ns	-		-		
PgR reduction (min. 1%): – Univariate	5.27	0.002	5.17	0.013	2.45 (0.89–6.70)	ns	2.33 (0.84–6.40)	ns	
– Multivariate	(1.82–15.18) 3.35 (0.99–11.28)	ns	(1.41–18.92) 2.61 (0.58–11.61)	ns	-		-		
K-67 <14 to ≥4: – Univariate – Multivariate	0.37 (0.16–0.87) 0.40 (0.13–1.17)	0.023 ns	0.31 (0.12–0.79) 0.55 (0.17–1.72)	0.015 ns	3.24 (1.36–7.69) 3.71 (1.45–9.47)	0.008 0.006	0.87 (0.38–1.98)	ns	

Survival analysis



Discordance: a technical issue...?



Nomura 1985 0.24 (0.12-0.39) 42 Spataro 1992 0.30 (0.26-0.35) 401 Li 1994 0.29 (0.19-0.40) 83 Shimizu 2000 0.25 (0.09-0.49) 20 Lower 2005 0.30 (0.24-0.37) 200 Broom 2009 0.18 (0.09-0.30) 62 Lindstrom 2012 0.32 (0.28-0.37) 459 Subtotal 0.27 (0.23-0.31) IHC Kamby 1989 0.28 (0.19-0.39) 92 Kuukasiärvi 1996 0.24 (0.13-0.38) 50 Sekido 2003 0.16 (0.07-0.30) 44 Gutierrez 2005 0.13 (0.04-0.27) 39 Gomez-Fernandez 2008 0.03 (0.01-0.06) 278 Wu 2008 0.20 (0.03-0.56) 10 Hoefnagel 2010 0.10 (0.07-0.15) 233 Idirisinghe 2010 0.16 (0.10-0.24) 117 Thompson 2010 0.10 (0.06-0.17) 137 Bogina 2011 0.06 (0.03-0.12) 140 Chang 2011 0.30 (0.19-0.44) 56 Curigliano 2011 0.15 (0.10-0.19) 255 Gong 2011 0.07 (0.04-0.12) 227 Nishimura 2011 97 0.10 (0.05-0.18) Sari 2011 0.36 (0.25-0.48) 75 Amir 2012 0.16 (0.09-0.25) 94 Duchnowska 2012 0.29 (0.21-0.38) 120 Jensen 2012 0.12 (0.07-0.19) 118 Montagna 2012 0.09 (0.06-0.13) 282 Aurilio 2013 0.21 (0.13-0.29) 107 235 Curtit 2013 0.17 (0.12-0.22) Dieci 2013 0.13 (0.08-0.21) 119 Ibrahim 2013 116 0.16 (0.10-0.24) 167 de Duenas 2014 0.13 (0.08-0.19) Subtotal 0.15 (0.12-0.18) Overall 0.19 (0.16-0.22) 0 .1 .2 .3 .4 .5 .6

Pusztai et al. The Oncologist 2010

Figure 2. Meta-analysis of the proportion of estrogen receptor discordance by type of assay. Abbreviations: CI, confidence interval; Prop., proportion.

Biochemical or mixed

Brennan 1979

Holdaway 1983

Raemaekers 1984

Hoehn 1979

Hull 1983

Prop. (95% CI)

0.24 (0.10-0.44)

0.19 (0.08-0.35)

0.54 (0.34-0.72)

0.21 (0.15-0.28)

0.19 (0.11-0.29)

Ν

29

37

28

167

75

...or a real fact?

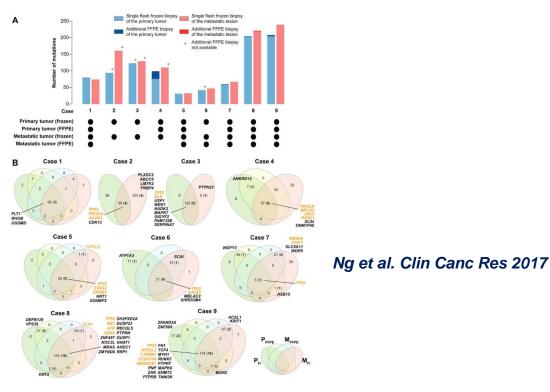


Fig. 1. Spatial and temporal heterogeneity in primary and metastatic lesions in treatment-naïve synchronous metastatic breast cancer

(A) Barplots depict the number of somatic mutations identified in single frozen biopsies of the primary tumors and metastatic lesions, and for seven patients, the additional mutations identified by analyzing an additional FFPE biopsy of the corresponding tumor. Available biopsy samples are indicated by black dots below the barplots. (B) Venn diagrams illustrate the number of somatic mutations and the number of likely pathogenic mutations (in bold) present in each of the biopsies. Genes affected by likely pathogenic mutations are listed, with those affecting cancer genes (31–33) labeled in orange. P_{Fr}: frozen biopsy of primary; P_{FFPE}: FFPE biopsy of primary; M_{Fr}: frozen biopsy of metastasis; M_{FFPE}: FFPE biopsy of metastasis. FFPE, formalin-fixed paraffin-embedded.

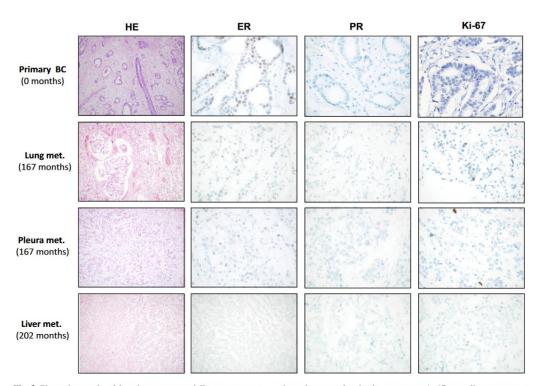


Fig. 3 Photomicrographs of the primary tumor and distant metastases of patient 10. The primary tumor was a LumB (ER+, PR- and Ki67 low, HER2 neg) IDC. After lumpectomy and sentinel lymph node biopsy (SLNB) the patient received radiation therapy, but declined any other adjuvant treatment. 14 years later metastases were discovered in the lung and pleura; the patient received multiple lines of

chemotherapy and endocrine treatment. At 17 years liver metastases occurred, treatment was switched back to chemotherapy, the patient passed away 4 months later due to progression of the disease. The lung and pleura metastases were ER weakly positive, but the liver metastasis was detected as TN

Discussion

• In our series ER discordance determined a therapeutic change. Among patients with PR discordance, the driver of therapeutic decisions was the ER status.

Our results suggest a potential positive prognostic role of stability in positive ER and PR expression.

• Possible role of previous exposure to anthracyclines, taxanes or aromatase inhibitors in hormonal receptor variations. Whether this phenomenon is due to an early clonal selection or to a real ER or PR loss is not clear and needs to be further investigated.

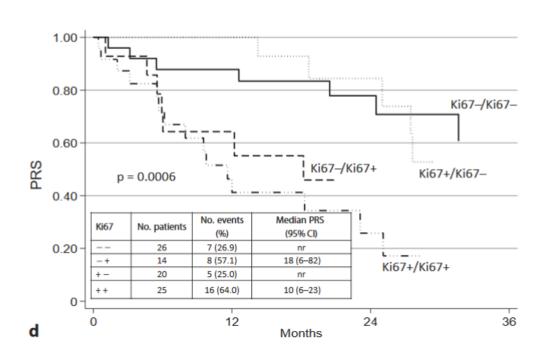
Ki-67 variation: metastatic like neoadjuvant?

Table 6 Uni- and multivariate analysis of factors for overall survival after recurrence

			Univariate	analysis	Multivariate	
Variables	Category	primary/recurrence	Relative risk	P value	P value	
Primary Tumor						
Tumor size	2.0 cm ≤2.0 cm</td <td>primary tumor</td> <td>0.98</td> <td>0.93</td> <td></td>	primary tumor	0.98	0.93		
Nodal status	0/1-3/4+	primary tumor	1.39	0.15		
Nuclear Grade	1/2/3	primary tumor	1.55	0.22		
Ki-67	50%≤/< 50%	primary tumor	5.67	0.0002	0.0006	
ER	+/-	primary tumor	0.49	0.07	0.09	
PgR	+/-	primary tumor	0.43	0.037	0.35	
p53	+/-	primary tumor	2.18	0.058	0.11	
HER2	+/-	primary tumor	0.9	0.82		
Recurrent Tumor						
DFI	5y≤/2y≤/< 2y	at recurrence	0.73	0.22		
Ki-67	50%≤/< 50%	recurrent tumor	2.88	0.009	0.32	
ER	+/-	recurrent tumor	0.36	0.013	0.51	
PgR	+/-	recurrent tumor	0.25	0.005	0.046	
p53	+/-	recurrent tumor	1.78	0.16		
HER2	+/-	recurrent tumor	0.95	0.91		
Distant metastasis	+/-	at recurrence	2.56	0.017	0.043	

ER: estrogen receptor; PgR: progesterone receptor; DFI: disease-free interval

Nishimura et al. World J Surg Oncol 2011



Ibrahim et al. Oncology 2013

^{*27} cases died of only breast cancer.

Conclusions

- Our results corroborate the importance of Ki67 as a prognostic factor and suggest a
 possible role for a proxy of acquired disease aggressiveness, consistently with data
 derived from the neoadjuvant setting
- These results may represent the basis for further prospective studies focused on the validation of Ki67 as a dynamic marker of prognosis, capable of representing the disease's evolution also in consideration of previous treatments
- If confirmed, these data could be integrated in the current follow-up strategies for a higher tailorization and precise approach.

Conclusions

 The molecular evaluation of metastatic lesions may influence the clinical decision-making process.

 Moreover, our results suggest the possible role of therapy exposure in modifying hormonal receptor status and Ki67 percentage of expression, and consequently biological behavior of metastatic BC.

