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ERRATUM

Gene expression meta-analysis identifies chromosomal regions and candidate genes involved in breast cancer metastasis

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Due to an unfortunate mistake on Springer's part, a wrong table was published in the article, [10.1007/s10549-008-9927-2](https://doi.org/10.1007/s10549-008-9927-2). Please find the correct table below.

The online version of the original article can be found under
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Table 3 Neighboring regions (constructed from gene set a and b) differently expressed between metastasizing and non-metastasizing tumors

	a		b									Mean	P-value	fdr					
	Amster-	dam	HUMAC	Huang	Rot-	terdam	Sotirou	2003	Sotirou	2006	Uppsala	Stock-							
<i>Upregulated</i>																			
CHR16Q22	CHR16Q24	4	49	4	7	44	16	3	4	4	43	5	16	9	1	14.3	<1E-6		
CHR8Q22	CHR8Q24	14	17	65	2	55	14	12	17	9	50	32	6	14	2	5	9	20.3	3.2E-05
CHR10Q22	CHR10Q2	24	26	44	6	22	22	6	10	40	8	56	1	40	10	13	2	20.8	5.5E-05
CHR17Q23	CHR17Q25	2	47	6	11	41	19	29	16	1	40	3	43	54	11	9	8	21.4	9.4E-05
CHR16P11	CHR16Q22	13	48	3	47	27	30	27	13	6	46	8	14	46	19	3	5	22.5	2.1E-04
CHR12Q12	CHR12Q13	11	21	64	24	37	9	8	22	31	6	12	25	11	6	25	61	23.4	4.5E-04
CHR20Q12	CHR20Q13	29	45	46	51	17	25	38	55	5	22	21	3	3	1	8	10	23.8	5.7E-04
CHR20Q11	CHR20Q12	3	31	16	5	2	3	1	1	32	39	49	52	21	29	44	54	24.3	8.3E-04
<i>Downregulated</i>																			
CHR14Q11	CHR14Q24	23	65	55	40	34	66	73	35	77	79	30	42	72	75	73	51	57.1	3.0E-04
CHR1P22	CHR1P13	75	54	75	60	45	30	76	73	23	72	59	43	67	59	73	59.7	2.7E-05	
CHR1P32	CHR1P31	53	61	48	60	56	59	74	21	79	47	78	39	53	74	77	78	61.3	9.0E-06
CHR1P31	CHR1P22	78	50	78	35	53	74	76	77	76	52	55	73	62	49	35	77	63.4	<1E-6

The ranking numbers indicate the ranking of each gene set out of the 79 gene sets in each dataset and the mean number indicate the ranking of the neighboring regions in the meta-analysis. Twelve significant out of a total of 79 neighboring regions are shown