

## Correction

**Correction: Increased copy number at 3p14 in breast cancer**Ingrid Ljuslinder<sup>1</sup>, Beatrice Malmer<sup>1</sup>, Irina Golovleva<sup>2</sup>, Marcus Thomasson<sup>1</sup>, Kjell Grankvist<sup>3</sup>, Thomas Höckenström<sup>4</sup>, Stefan Emdin<sup>5</sup>, Yvonne Jonsson<sup>1</sup>, Håkan Hedman<sup>1</sup> and Roger Henriksson<sup>1</sup><sup>1</sup>Department of Radiation Sciences, Oncology, Umeå University, Umeå, Sweden<sup>2</sup>Department of Medical Biosciences, Medical and Clinical Genetics, Umeå, Sweden<sup>3</sup>Department of Medical Biosciences, Clinical Chemistry, Umeå University, Umeå, Sweden<sup>4</sup>Department of Pathology, Umeå University, Umeå, Sweden<sup>5</sup>Department of Surgical and Perioperative Sciences, Division of Surgery, Umeå University, SwedenCorresponding author: Ingrid Ljuslinder, [ingrid.ljuslinder@onkologi.umu.se](mailto:ingrid.ljuslinder@onkologi.umu.se)

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It was recently noticed that an error had been made in the research article by Ljuslinder and colleagues [1] published by *Breast Cancer Research* in July 2005. The legend of Table 4 was incorrect and should read as featured opposite.

**Reference**

1. Ljuslinder I, Malmer B, Golovleva I, Thomasson M, Grankvist K, Höckenström T, Emdin S, Jonsson Y, Hedman H, Henriksson R: **Increased copy number at 3p14 in breast cancer.** *Breast Cancer Research* 2005, **7**:R719-R727.

**Table 4****EGFR and ERBB2 quantitative RT-PCR mRNA results in eight patients and matched controls (group B)**

Patient number	T	NN	T/NN <sup>a</sup>
1 <sup>b</sup>			
EGFR	4,677	12,493	0.37
ERBB2	24,066	21,817	1.1
2 <sup>b</sup>			
EGFR	3,321	14,519	0.22
ERBB2	61,389	6,666	9.2
4			
EGFR	20,778	14,842	1.39
ERBB2	9,143	6,742	1.35
5 <sup>b</sup>			
EGFR	9,730	15,867	0.61
ERBB2	13,506	3,056	4.41
6			
EGFR	1,305	15,871	0.08
ERBB2	10,677	10,080	1.06
7			
EGFR	1,024	12,162	0.08
ERBB2	7,639	5,871	1.30
8 <sup>b</sup>			
EGFR	4,936	28,111	0.17
ERBB2	17,106	9,263	1.85
9			
EGFR	5,540	1,437	3.85
ERBB2	7,330	5,540	1.32

<sup>a</sup>T/NN: mRNA expression levels of EGFR/ERBB2 in tumour tissue (T) samples divided by values in matched non-neoplastic (NN) tissue samples. As described in the results, at p.9, ratios > 1.2 are regarded as significant overexpression and ratios < 0.8 are regarded as significant underexpression in neoplastic tissue compared to non-neoplastic tissue. <sup>b</sup>Patients with increased *LRIG1* copy number. No RNA from patient 3 was available for EGFR/ERBB2 analysis.