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Erratum to: Generation of an algorithm based on minimal gene sets to clinically subtype triple negative breast cancer patients

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Erratum

Since publication of our article [1], we have noticed an error in labelling the p-value of one of the results. The original document read:

"The direction of association of BL1 with pCR or minimal residual cancer burden (RCB) in the 2188-gene model was similar (OR = 1.91) but did not reach significance (OR = 0.14)."

This has been corrected to:

"The direction of association of BL1 with pCR or minimal residual cancer burden (RCB) in the 2188-gene model was similar (OR = 1.91) but did not reach significance (p = 0.14)."

We regret the error.

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References
1. Ring BZ, Hout DR, Morris SW, Lawrence K, Schweitzer BL, Bailey DB, et al. Generation of an algorithm based on minimal gene sets to clinically subtype triple negative breast cancer patients. BMC Cancer. 2016;16:143. doi:10.1186/s12885-016-2198-0.

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