

Correction

Correction: Genomic profiling of breast tumours in relation to *BRCA* abnormalities and phenotypes

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Following the publication of our article [1] we noticed an error in the article PDF.

Due to a production error, the final two sentences of the legend to figure 2a are incomplete. These two sentences should appear as follows:

The characters represent cluster memberships of each tumour with *BRCA1* and *BRCA2* abnormalities indicated, see bottom of the figure. It can be hypothesised here that component three reflects differences between *BRCA1*- and *BRCA2*-related tumours whereas component two reflects their similarities, see further in Additional data file 3.

Reference

1. Stefansson OA, Jonasson JG, Johannsson OT, Olafsdottir K, Steinarsdottir M, Valgeirsdottir S and Eyfjord JE: **Genomic profiling of breast tumours in relation to *BRCA* abnormalities and phenotypes.** *Breast Cancer Research* 2009, **11**:R47.