

# CORRECTIONS & AMENDMENTS

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## CORRECTION

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## Author Correction: Landscape of somatic mutations in 560 breast cancer whole-genome sequences

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In the Methods section of this Article, 'greater than' should have been 'less than' in the sentence 'Putative regions of clustered rearrangements were identified as having an average inter-rearrangement distance that was at least 10 times greater than the whole-genome average for the individual sample.'. The original Article has not been corrected online.