

CORRECTION

Correction: Integrated Multiregional Analysis Proposing a New Model of Colorectal Cancer Evolution

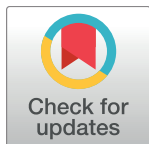
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The Data Availability Statement for this article is incorrect. The correct statement is: Sequence data are available from the DNA Data Bank of Japan (DDBJ) under the accession number JGAS00000000060 (<https://ddbj.nig.ac.jp/jga/viewer/view/study/JGAS00000000060>) for researchers who meet the criteria for access to confidential data. Array data have been deposited to the Gene Expression Omnibus database, with the accession number GSE90709 (<https://www.ncbi.nlm.nih.gov/geo/query/acc.cgi?acc=GSE90709>).

There is an error in the third sentence of the Abstract. The phrase ‘intertumor heterogeneity’ is incorrect, and should read ‘intratumor heterogeneity’. The correct sentence is: ‘Extensive intratumor heterogeneity is observed, from which we inferred the evolutionary history of the tumors.’

Reference

1. Uchi R, Takahashi Y, Niida A, Shimamura T, Hirata H, Sugimachi K, et al. (2016) Integrated Multiregional Analysis Proposing a New Model of Colorectal Cancer Evolution. *PLoS Genet* 12(2): e1005778. <https://doi.org/10.1371/journal.pgen.1005778> PMID: 26890883



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