

ERRATUM

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Erratum to: A coherent approach for analysis of the Illumina HumanMethylation450 BeadChip improves data quality and performance in epigenome-wide association studies

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After the publication of this work [1] it was noticed that an author was omitted from the author list and author contributions. Please see below updated author contributions.

Authors' contributions

Data analysis and implementation of analysis pipeline: BL, AD, ML, RS. Manuscript writing: BL, AD, ML, JCC. Intellectual guidance in data-analysis and interpretation: JCC, ML, WZ, WRS, STT, UA, JS, MRJ, PE, MIM, JSK. Project management: JSK, JCC.

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