Erratum: Characterization of functional methylomes by next-generation capture sequencing identifies novel disease-associated variants

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Nature Communications 6:7211 doi: 10.1038/ncomms8211 (2015); Published 29 May 2015; Updated 29 Jul 2015

The HTML version of this Article incorrectly included all members of The Multiple Tissue Human Expression Resource Consortium in the list of authors. This error has now been corrected in the HTML; the PDF version of the paper was correct from the time of publication.