

Erratum: Probes: FRET sensor design and optimization

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In the version of this article initially published, the captions and image credit information for images from the Frommer lab and the Lin lab were swapped. The image from the Frommer lab was also truncated. The errors have been corrected in the PDF and HTML versions of this article.

Corrigendum: Accurate identification of single-nucleotide variants in whole-genome-amplified single cells

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In the version of this article initially published, Lodato, M.A. *et al. Science* **350**, 94–98 (2015) (reference 2) was cited as an example of a single-cell sequencing study with high CG-to-TA transitions that applies heat lysis. However, that work used alkaline lysis on ice (Walsh, C.A. and Lodato, M.A., personal communication); therefore, we have changed the third sentence of the paper from “This pathway may explain the observed excess of such mutations in single neurons² compared with unamplified neuronal clones³” to “Amplification artifacts could, in general, explain the observed excess of such mutations in single neurons² compared with unamplified DNA from neuronal clones³.” The error has been corrected in the HTML and PDF versions of the article.