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Erratum to: MSRE-HTPrimer: a highthroughput and genome-wide primer design pipeline optimized for epigenetic research

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Erratum

Unfortunately, the original version of this article [1] contained an error. The name of one of the authors was included incorrectly and it read "Pulverer Walter" instead of "Walter Pulverer".

The name has been updated in the original article and is correctly included in full in this erratum.

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 Pandey RV, Pulverer W, Kallmeyer R, Beikircher G, Pabinger S, Kriegner A, et al. MSRE-HTPrimer: a high-throughput and genome-wide primer design pipeline optimized for epigenetic research. Clin Epigenetics. 2016;8:26. doi: 10.1186/s13148-016-0190-9.

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