## CORRIGENDUM

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## Corrigendum: DNA-binding factors shape the mouse methylome at distal regulatory regions

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In the original version of this Article Erik van Nimwegen (Biozentrum of the University of Basel and Swiss Institute of Bioinformatics, Klingelbergstrasse 50-70, CH 4056 Basel, Switzerland) was inadvertently omitted from the author list. In the 'Author contribution' section the sentence beginning "Bioinformatic and statistical analyses..." should read "Bioinformatic and statistical analyses were conceived and conducted by M.B.S., L.B., R.I. and E.v.N." In addition, in the Supplementary Information section 'Unsupervised Segmentation of CpG Methylome', the HMM emission means were given as percentages instead of fractions, contained a typo in the third variance parameter, and should read as follows: "Emission probabilities of HMM states were modelled as Gaussian distributions with means 0.8, 0.28, 0.03 and variances 0.01, 0.005, 0.0026...". The PDF and HTML versions of this paper have been corrected, and the Supplementary Information has been replaced.