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**CORRIGENDUM**

## Three-dimensional genome architecture: players and mechanisms

*Ana Pombo and Niall Dillon*

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In the original article, the following sentence was incorrect: “Histone marks associated with enhancers, such as histone 3 Lys4 monomethylation (H3K4me1), and with transcription repression, such as histone 3 Lys9 trimethylation (H3K9me3), were also enriched at TAD boundaries.”

The corrected sentence is as follows: “Histone marks associated with enhancers, such as histone 3 Lys4 monomethylation (H3K4me1), and with transcription repression, such as histone 3 Lys9 trimethylation (H3K9me3), were not found to be enriched at TAD boundaries.”

This has been corrected in the online version of the article.