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Corrigendum: Chromatin structure and context-dependent sequence features control prime editing efficiency

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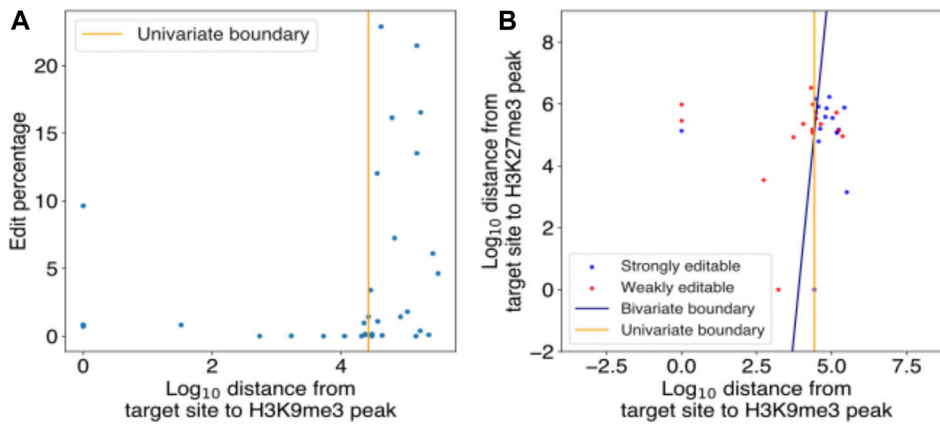
KEYWORDS

prime editing, CRISPR–Cas9, heterochromatin, nucleosome positioning, DNA-RNA hybridization, nucleotide preference, machine learning, neural network interpretation

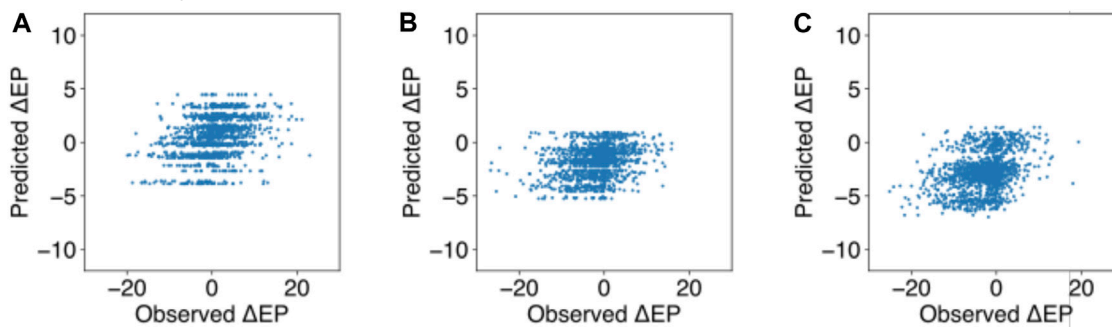
A Corrigendum on Chromatin structure and context-dependent sequence features control prime editing efficiency

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In the published article, there was an error in **Supplementary Figure S3**. Even though the figure caption for Supplementary Figure S3 was correct, Figure S3 was inadvertently duplicated from Figure S2 and appeared as:



The correct figure appears below.



The authors apologize for this error and state that this does not change the scientific conclusions of the article in any way. The original article has been updated.

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