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Correction: Quantification of hematopoietic stem and progenitor cells by targeted DNA methylation analysis

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The wrong Additional file 4 was originally published with this article [1]. It has now been replaced with the correct file (Additional file 4).

The original article has been corrected.

Supplementary Information

The online version contains supplementary material available at https://doi.org/10.1186/s13148-023-01532-7.

Additional file 4. Table S6. Application for NNLS model with 6 CpGs.

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Reference

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