

## CORRECTION

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### **Author Correction: BCAT1 restricts $\alpha$ KG levels in AML stem cells leading to IDH<sup>mut</sup>-like DNA hypermethylation**

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We have deposited the raw mass spectrometry proteome data and search results to the ProteomeXchange Consortium (<http://proteomecentral.proteomexchange.org>) via the Proteomics Identifications Database (PRIDE) partner repository with the dataset identifier PXD008307. In addition, we now supply Supplementary Data 4 to describe our sampling and labelling strategies, and to list all identified proteins and their differential expression in leukaemia stem cells (LSCs) compared to non-LSCs. In Extended Data Fig. 1a of this Letter, the flow cytometry plot depicting the surface phenotype of acute myeloid leukaemia (AML) sample DD08 was a duplicate of the plot for AML sample DD06. Extended Data Fig. 1 has been corrected online (the Supplementary Information of this Amendment contains the incorrect version of Extended Data Fig. 1, for transparency), and Supplementary Data 4 has been added to the Supplementary Information of the original Letter.

**Supplementary information** is available for this Amendment at <https://doi.org/10.1038/s41586-018-0403-9>