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CORRECTION

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Correction to: HDACi mediate UNG2 depletion, dysregulated genomic uracil and altered expression of oncoproteins and tumor suppressors in B- and T-cell lines

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Correction to: J Transl Med (2020) 18:159
https://doi.org/10.1186/s12967-020-02318-8

The original publication of this article [1] contained 2 incorrect PRIDE identifiers. In this correction article the incorrect and correction information is published. The original article has been updated.

Incorrect

- The mass spectrometry proteomics data have been deposited to the ProteomeXchange Consortium [16] via the PRIDE partner repository with the dataset identifier **PXD00008293**.

Correct

- The mass spectrometry proteomics data have been deposited to the ProteomeXchange Consortium [16] via the PRIDE partner repository with the dataset identifier **PXD008293**.

Incorrect

- The SILAC mass spectrometry proteomics data have been deposited to the ProteomeXchange Consortium (16) via the PRIDE partner repository with the dataset identifier **PXD00008293**.

Correct

- The SILAC mass spectrometry proteomics data have been deposited to the ProteomeXchange Consortium (16) via the PRIDE partner repository with the dataset identifier **PXD008293**.

The original article can be found online at https://doi.org/10.1186/s12967-020-02318-8.

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1. Iveland TS, Hagen L, Sharma A, Sousa MML, Sarno A, Wollen KL, Liabakk NB, Slupphaug G. HDACi mediate UNG2 depletion, dysregulated genomic uracil and altered expression of oncoproteins and tumor suppressors in B- and T-cell lines. J Transl Med. 2020;18:159. https://doi.org/10.1186/s12967-020-02318-8.

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