

CORRECTION

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# Correction to: Metabolomic profiling identifies distinct phenotypes for ASS1 positive and negative GBM

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## Correction

In the original publication of this article [1], published on 8 February 2018, it was noticed that the acknowledgement of the source of the drug ADI-PEG20 was missing. In this Correction, the source of drug ADI-PEG20 is shown (and marked bold). This addition is made in the Methods section, under the heading ADI-PEG20 treatment.

ADI-PEG20 treatment.

SNB19 and U87 cells, cultured in DMEM + 10% FBS and normal human astrocytes, cultured in speciality media provided by lonza were seeded in replicates ( $n = 12$ ) at  $8 \times 10^4$  cells per well in 6-well dishes (Corning, NY, USA). 24 h post seeding, cells were washed with phosphate buffered saline (PBS) and cultured in the presence or absence of ADI-PEG20 (**kindly provided by Polaris Pharmaceuticals Inc, San Diego, CA**) (1  $\mu\text{g}/\text{ml}$ ) in media containing, 1 mM citrulline and 10% fetal FBS. ADI-PEG20 was added at the start of the experiment and no fresh media was added to any of the experimental plates before harvesting. ADI-PEG20 treated and untreated media ( $n = 3$ ) was included for normalization purposes. 48 h after ADI-PEG20 treatment replicate samples for each condition ( $n = 3$ ) were harvested, collecting both spent media and cells for GC-TOFMS metabolomic analysis. Additional replicates ( $n = 3$ ) of each condition were collected for total cell count determination.

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1. Mörén L, et al. Metabolomic profiling identifies distinct phenotypes for ASS1 positive and negative GBM. *BMC Cancer*. 2018;18:167. <https://doi.org/10.1186/s12885-018-4040-3>.

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