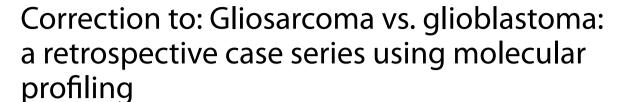
PUBLISHER CORRECTION

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Following publication of the original article [1], the authors reported an error that the wrong version of the Additional file 2 was published. The correction version of Additional file 2 is now updated.

The original article [1] has been updated.

Supplementary Information

The online version contains supplementary material available at https://doi.org/10.1186/s12883-021-02326-1.

Additional file 2. This is the statistical analysis, provided in .pdf format. This contains the R code used in the analysis and so parts of this are likely to be hard to understand for the reader not familiar with this language. The file provides the results given in the main article. It includes some additional analyses and images which, for reasons of space, do not form part of the main article. In particular, it provides details of *all* of the MTs assessed. For each gene assessed, we also provide hyperlinks (where possible), to the relevant entry at the NCBI Gene database (https://www.ncbi.nlm.nih.gov/gene).

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