Correction to: A Risk Signature Consisting of Eight m⁶A Methylation Regulators Predicts the Prognosis of Glioma

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The original version of this article unfortunately contained a typo in caption of Fig. 4A.

The sentence "yellow represents the gene expression difference in GBM and LGG patients" should be revised as "yellow represents no significant gene expression difference in GBM and LGG patients".

The original article can be found online at <https://doi.org/10.1007/s10571-021-01135-x>.

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Fig. 4 Interactions among the top 20 differentially expressed m⁶A methylation regulators. **A** Protein–Protein Interactions (PPI) network interaction between the top 20 differentially expressed m⁶A methylation regulators. A triangle represents “Readers”, a rectangle represents “Writers”, and a circle represents “Erasers”. Red represents the gene expression difference in GBM and LGG patients, pink represents the gene expression difference in GBM or LGG patients, and

yellow represents no significant gene expression difference in GBM and LGG patients. **B** Correlation between the expression of m⁶A methylation regulators and CNV. **C** Correlation between the expression levels of m⁶A methylation regulators. The size of the sector in the square represents the correlation. The blue color represents the positive correlation, and the dark red represents the negative correlation

Hence, the correct Fig. 4A with its caption is presented here.

The original article has been corrected.

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