

CORRECTION

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Correction to: Medulloblastoma cerebrospinal fluid reveals metabolites and lipids indicative of hypoxia and cancer-specific RNAs

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Correction to: *Acta Neuropathologica Communications* (2022) 10:25
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Following publication of the original article [1], the authors identified an error in the author name of Iqbal Mahmud.

The incorrect author name is: Iqbal Mohamad

The correct author name is: Iqbal Mahmud

In addition, the authors identified an error in Fig. 1, panel B. The correct figure is given below.

The original article [1] has been corrected.

The original article can be found online at <https://doi.org/10.1186/s40478-022-01326-7>.

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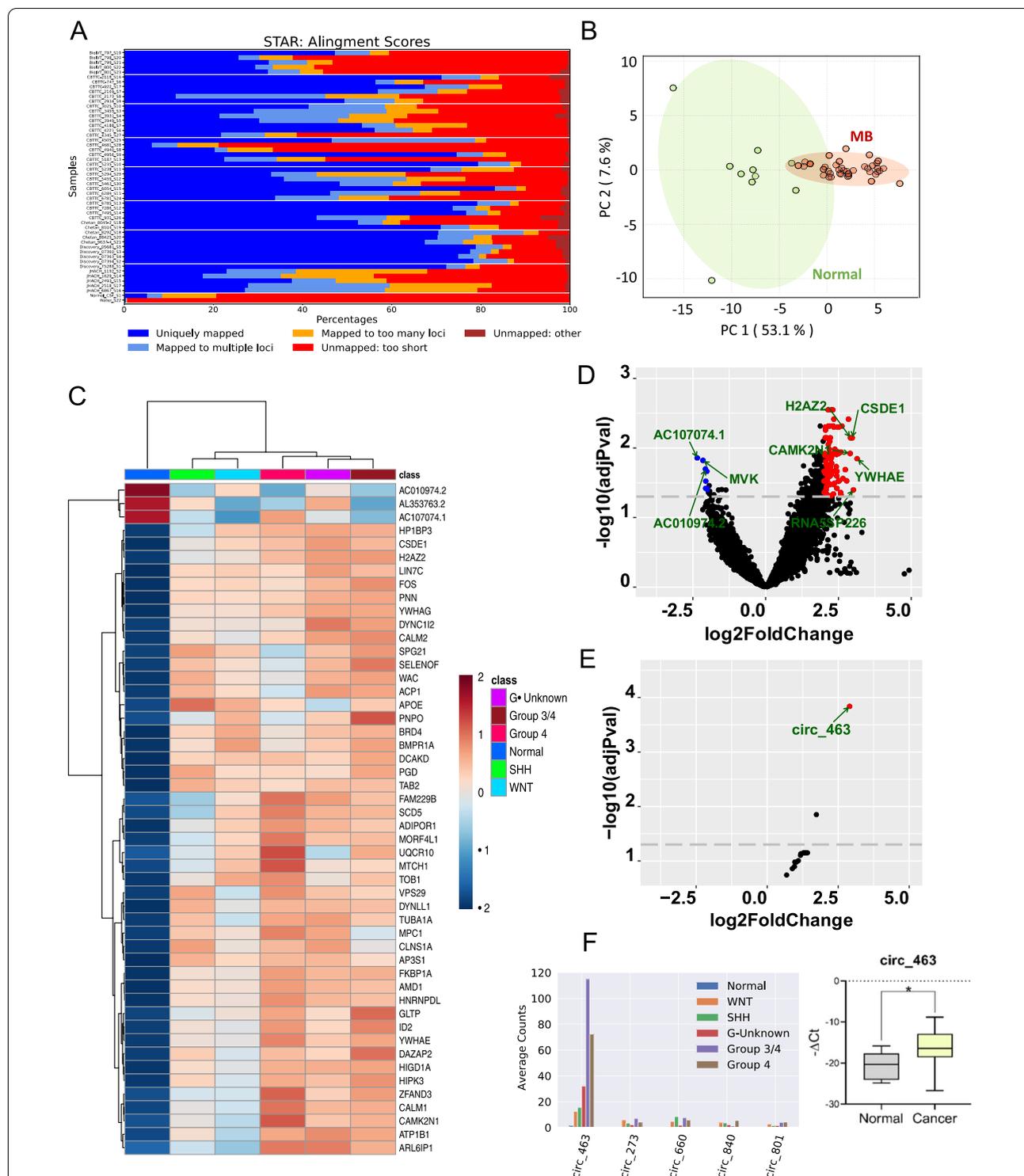


Fig. 1 Global transcriptomic differences in the CSF of patients with (n = 40) and without (n = 11) MB. **A** Mapping rate of each CSF sample. **B** Principal component analysis of CSF samples using the 48 most differentially expressed genes showing clear separation of normal CSF samples from MB CSF samples. **C** Unsupervised clustering of samples using the 48 most differentially expressed genes showing clear separation of normal CSF samples from MB CSF samples. **D** Volcano plot showing significantly up- or downregulated genes in CSF. **E** Volcano plot for differentially expressed circRNAs between normal vs MB CSF samples. **F** Top 5 circRNAs expression in different subgroups and qRT-PCR validation of circ-463

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