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Correction: ASPM-associated stem cell proliferation is involved in malignant progression of gliomas and constitutes an attractive therapeutic target

Sandra-Nadia Ngwabyt Bikeye¹, Carole Colin², Yannick Marie¹, Raphaël Vampouille⁴, Philippe Ravassard⁴, Audrey Rousseau⁵, Blandine Boisselier¹, Ahmed Idbaih^{1,3}, Charles Félix Calvo¹, Pascal Leuraud⁶, Myriam Lassalle⁶, Soufiane El Hallani¹, Jean-Yves Delattre^{1,3} and Marc Sanson^{1,3*}

Correction

In the original article [1], the additional file two: figure S. Two (additional file 1 here) contained two identical graphs. We present here the corrected figure S2 and the corresponding legend:

Additional material

Additional file 1: Tumor spheroid characterization. Genomic stability was examined with CGHa analysis (left = DNA profile from the initial tumor; right = DNA profile from gliomasphere at passage p28). The chromosomes are indicated on the x axis and copy number is on y axis. Yellow indicates the normal genomic copy number, while green indicates a loss and red indicates a gain in copy number. Although some differences are observed, the comparison of both profiles shows that overall genomic profile is quite well preserved (except for an amplicon on chromosome 8 present only in the gliomasphere).

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