scientific reports



OPEN Author Correction: PR-LncRNA signature regulates glioma cell activity through expression of SOX factors

Published online: 20 June 2022

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Correction to: Scientific Reports https://doi.org/10.1038/s41598-018-30836-5, published online 24 August 2018

The original version of this Article contained errors in Figure 3.

In Figure 3C, the representative image for ASO10A was inadvertently duplicated as ASO1B. Data from the fourth experimental replicate was also omitted from Figure 3D. The original Figure 3 and accompanying legend appear below.

The original Article has been corrected.

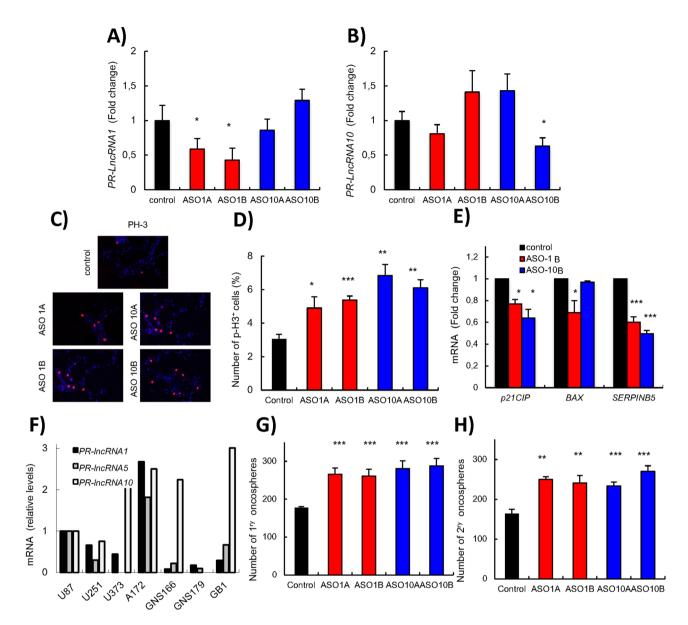


Figure 3. *PR-LncRNA1* and *10* silencing leads to increased proliferation and stemness. U87-MG cells were transfected with specific ASOs for the *PR-LncRNAs* indicated. (**A,B**) Transfected cells were examined for *PR-LncRNA1* and *PR-LncRNA10* expression by quantitative reverse transcription polymerase chain reaction (n = 4). (**C**) Representative immunofluorescence of P-H3 in U87MG cells under the conditions indicated. (**D**) Quantification of the number of P-H3 positive cells under the conditions indicated (n = 4). (**E**) Quantification of mRNA levels of $p21^{cip}$, *Bax and SerpinB5* in cells transfected with ASOs for *PR-LncRNA1* and 10 and compared to cells with a control ASO (**F**) Expression of *PR-LncRNA 1,5* and 10 in indicated conventional cell lines (U87-MG, U251, U373 and A172) and glioma stem cells (GNS166, GNS179 and GB1) (**G**) Quantification of primary oncospheres formed in ASO-transfected cells after 10 days in culture (n = 3). (**H**) Quantification of number of secondary oncospheres generated from disaggregating primary oncospheres in ASO-transfected and control cells. Numbers were assessed after 10 days in culture (n = 3). Asterisks (*,**and ***) indicate statistical significance (p < 0.05, p < 0.01, and p < 0.001, respectively).

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