Author Correction: Integrated genomic analyses of de novo pathways underlying atypical meningiomas


Nature Communications 8:14433 doi: 10.1038/ncomms14433 (2017) Published online 14 Feb 2017; Updated 20 Apr 2018

In this Article, a subset of the H3K27ac ChIP-seq data (15 benign meningiomas and 2 dura samples (Sample IDs: MN-297, MN-288, MN-292, MN-163, MN-1037, MN-105, MN-201, MN-249, MN-191, MN-1066, MN-169, MN-291, MN-24, MN-79, MN-1044, CONTROL1, CONTROL2) was reported previously in a publication by the corresponding author1. These data were created by Dr. Justin Cotney in Dr. James Noonan’s laboratory at Yale. The GEO database entry associated with this dataset has been updated to reflect this fact (GSE91372).


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