



Supplementary Figure 3:

GTEX expression profiles for SETD1B (top) and SETD1A (bottom) are shown above, calculated from a model that collapses all isoforms to a single gene, with no other normalization steps applied. X axis lists tissue subtypes, with tissue category as the background colour (nervous system is yellow). Y axis is “Transcripts per Million” (TPM) as calculated across 980 subjects. Boxplots show the median, 25th and 75th centiles; outliers that fall outside 1.5 times the interquartile range are marked.

SETD1B is expressed broadly. In brain, it is preferentially expressed in cerebellum and cerebellar hemisphere. No data for bone expression are available.

Source: <https://www.gtexportal.org/home/gene/SETD1B>