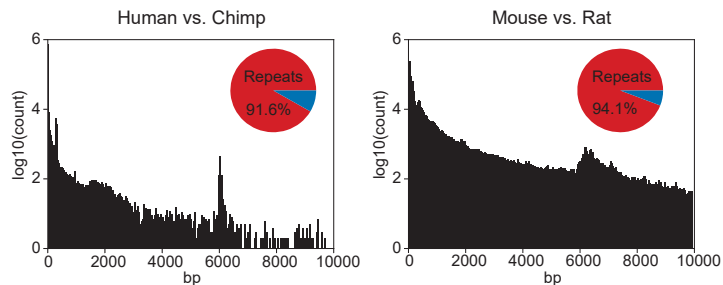


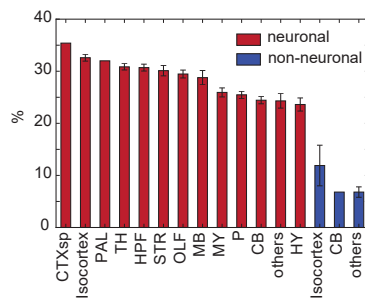
A

Lengths of inserted segments



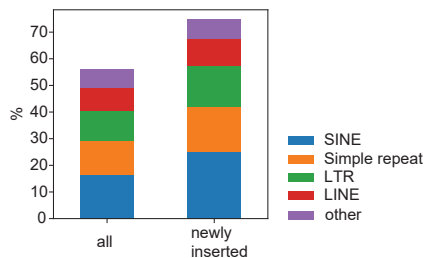
B

% of longest 500 genes with FPKM > 10



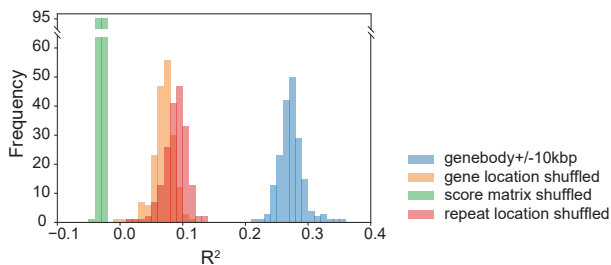
C

Overlap of ATAC peaks and repeats



D

Cross-validated regression



E

Gene Length Difference Between Species

