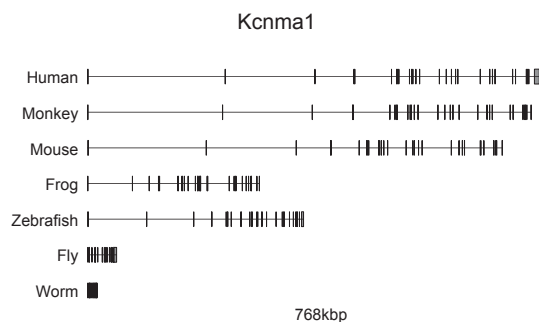
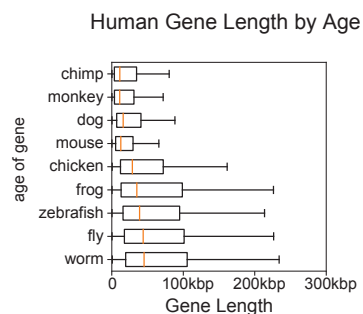


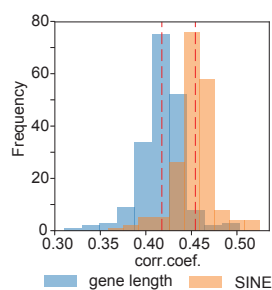
A



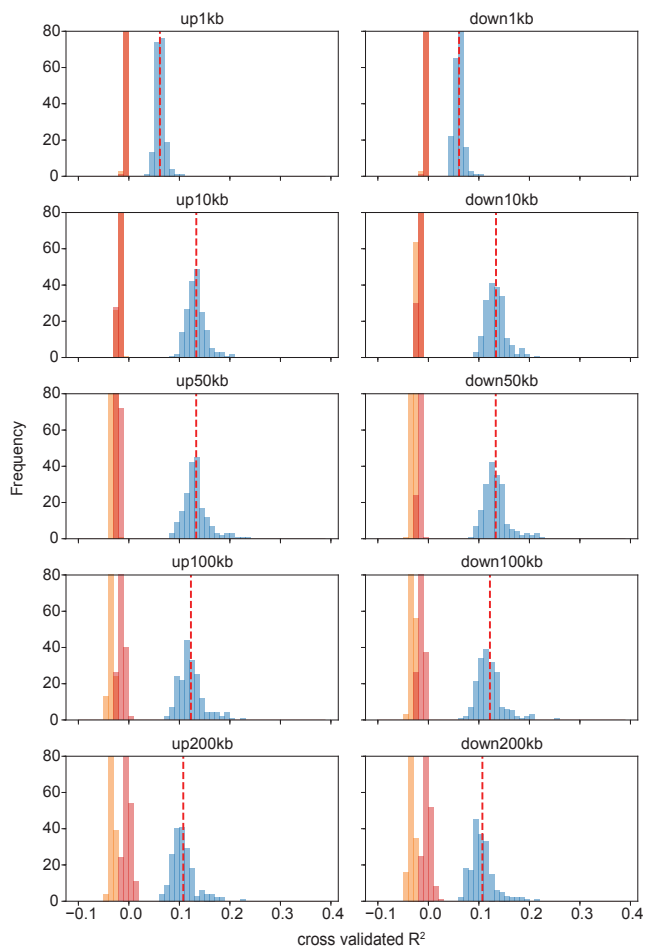
B



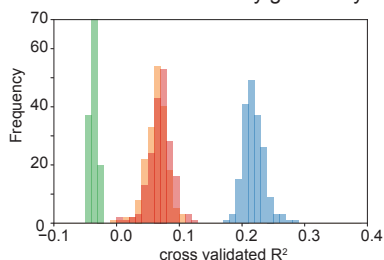
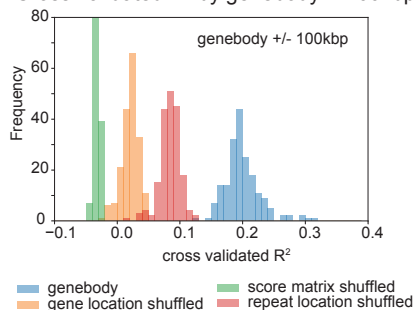
C

Pearson's r to expression rank

D

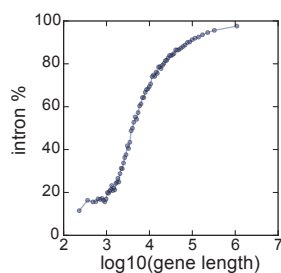
Cross-validated R^2 by Surround TE scores

E

Cross-validated R^2 by genebodyCross-validated R^2 by genebody +/- 100kbp

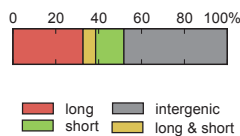
F

intron % vs. gene length



G

Percentage of base pairs by long, short and intergenic sequences



H

Percentage of TE inserted genes

