

Supplementary material 3. Homopolymer size variation across five *Drosophila* species in relationship with exon splicing pattern in *D. melanogaster*. The homopolymer size variation is assessed using the total length of a neighbor joining tree built using the absolute size variation between species. CSE: constitutively spliced exon, ONE: exon found in gene with a single annotated protein isoform. The box shows the first and third quartiles, the dotted lines extend to the 5th and 95th percentiles, the size of the notch indicates level of uncertainty associated with the median.

