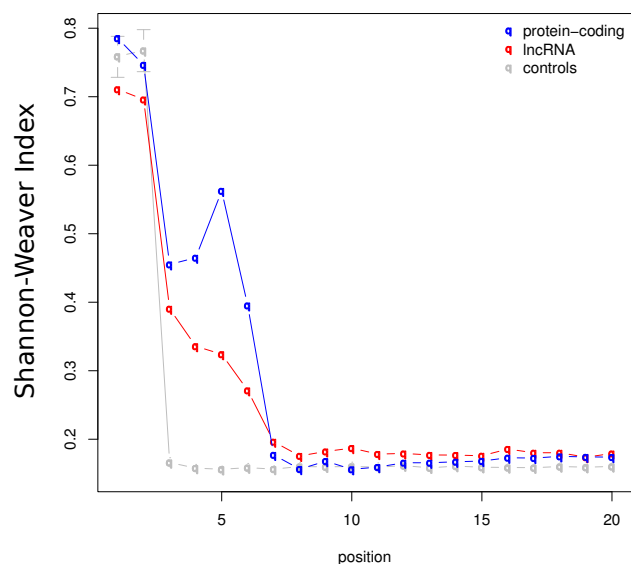


Additional File 3: Comparison of protein-coding (blue) and lncRNA (red) 5' (A) and 3' (B) splice site conservation in *D. melanogaster*. Only protein coding sequences flanking lncRNAs were used in the analysis. The control set is based on the random selection of "GT" and "AG" dinucleotides within the intergenic sequence flanking the lncRNAs in *D. melanogaster*. The Simpson-Weaver index was computed for each sites using the alignments of each splice site and its neighbouring sequences with *D. simulans*, *D. sechellia* *D. yakuba* and *D. erecta* with Muscle.

A



B

