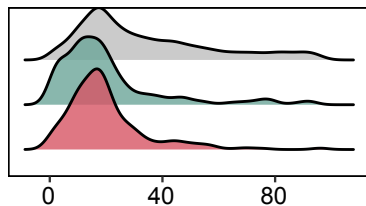
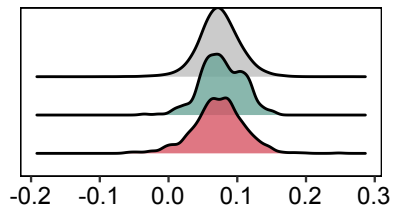
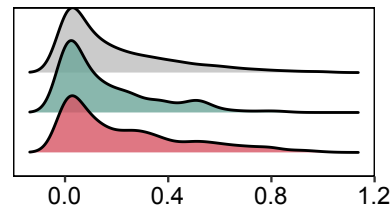
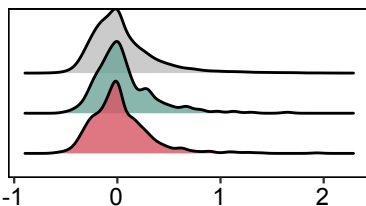
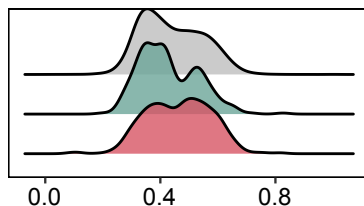
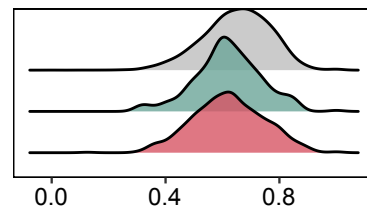
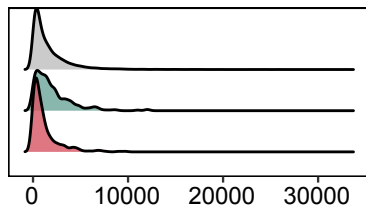
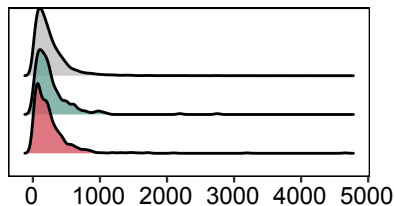
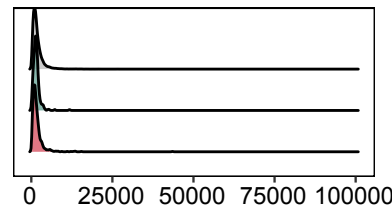
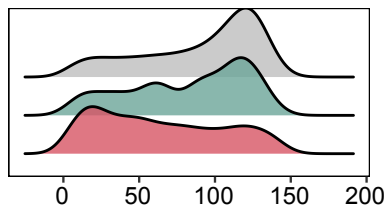


Broad conservation**Codon Pair Bias****Disorder****Distance-to-Median****GC content 3'UTR****GC content 5'UTR****Length 3'UTR****Length 5'UTR****Length CDS****Number of transcription factors****PPI degree**