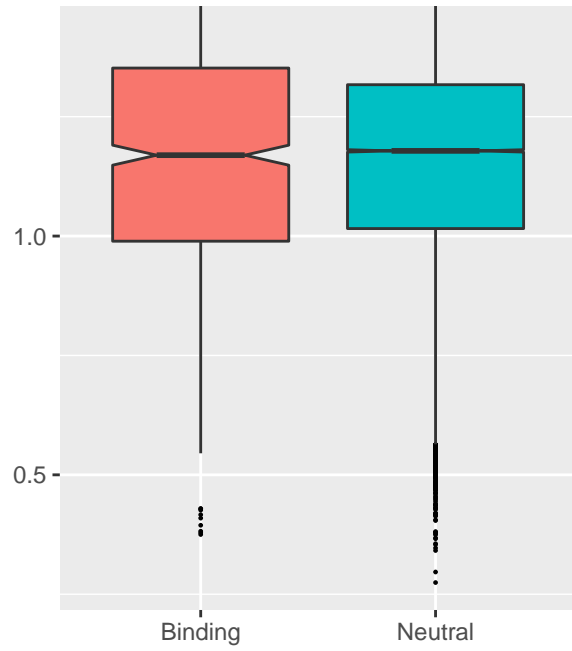
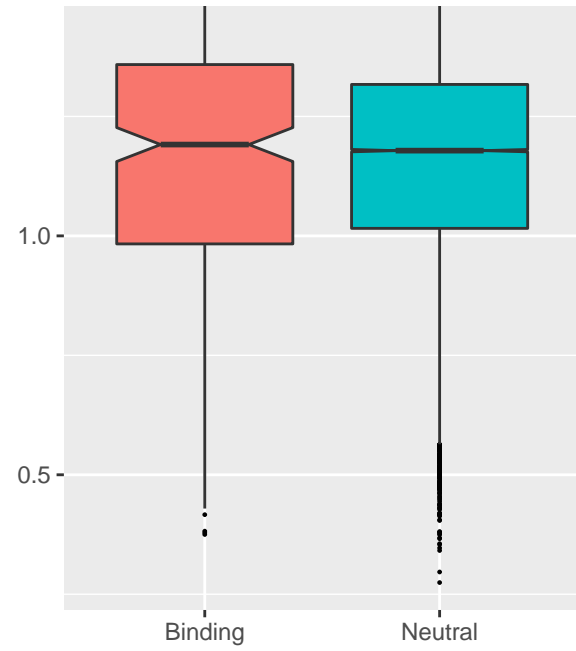


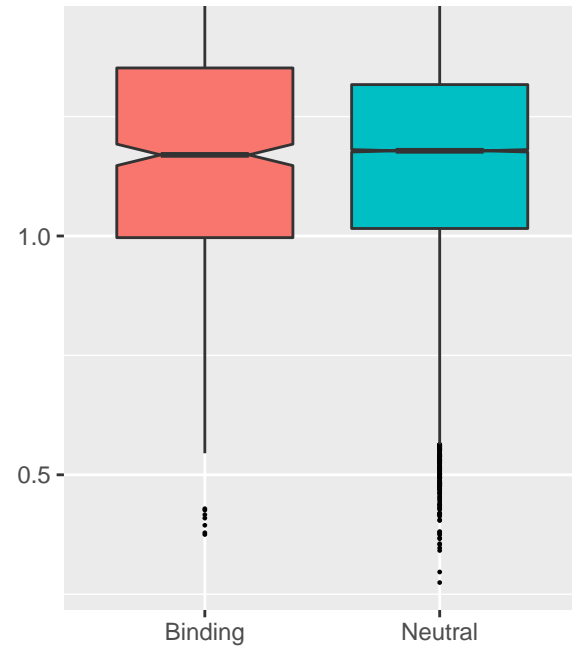
dna, jsd_SE_sum
#(binding)=749
p=0.45



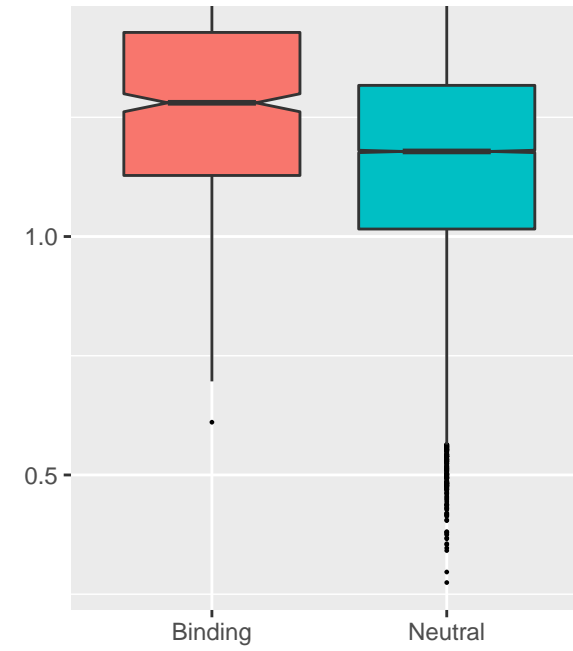
dnabase, jsd_SE_sum
#(binding)=279
p=0.15



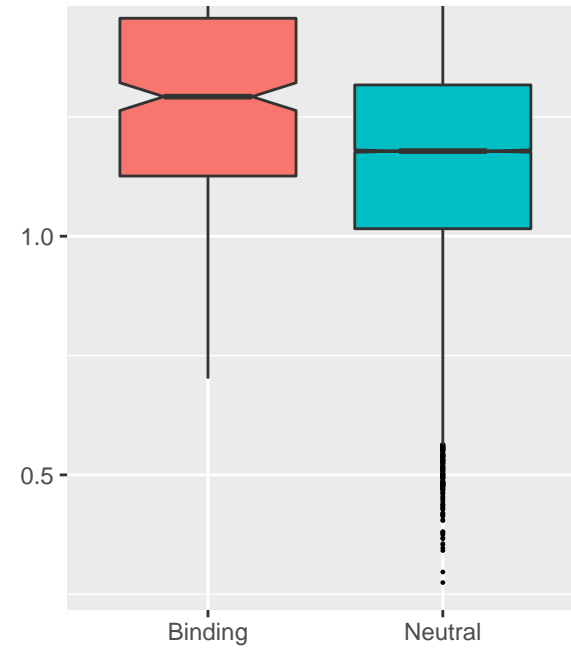
dnabackbone, jsd_SE_sum
#(binding)=625
p=0.36



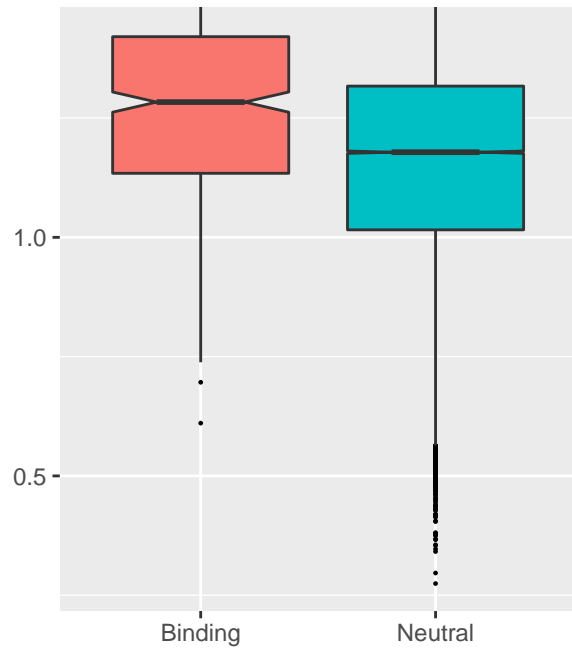
rna, jsd_SE_sum
#(binding)=629
p=4.4e-31



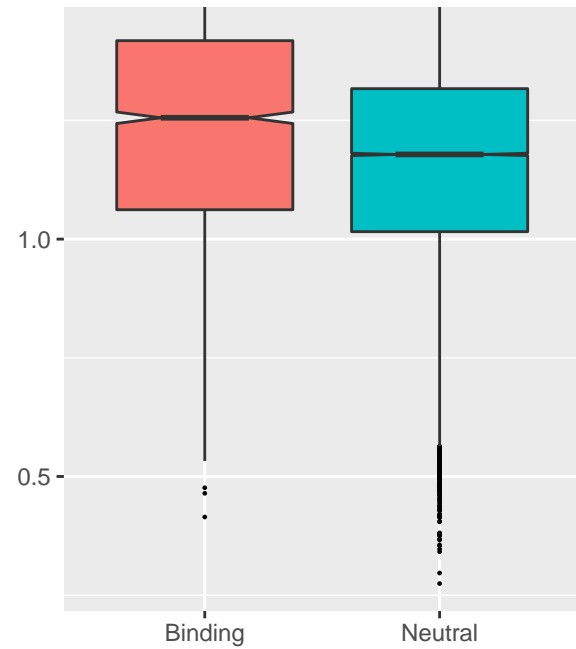
rnabase, jsd_SE_sum
#(binding)=322
p=3.9e-19



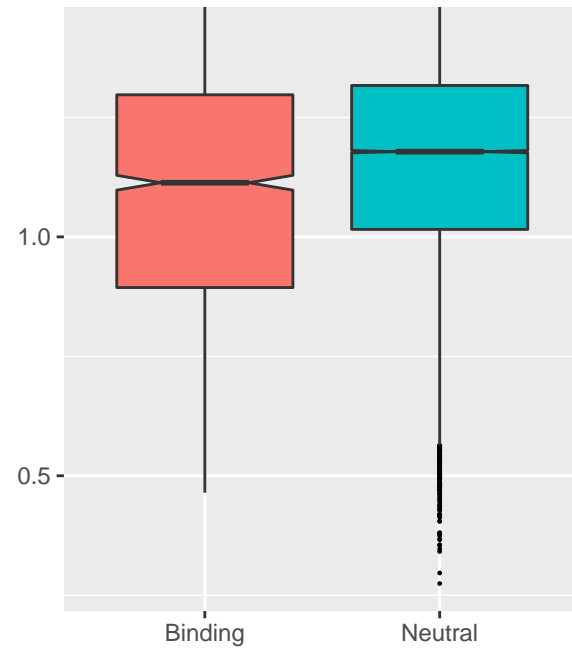
rnabackbone, jsd_SE_sum
#(binding)=458
p=1.4e-23



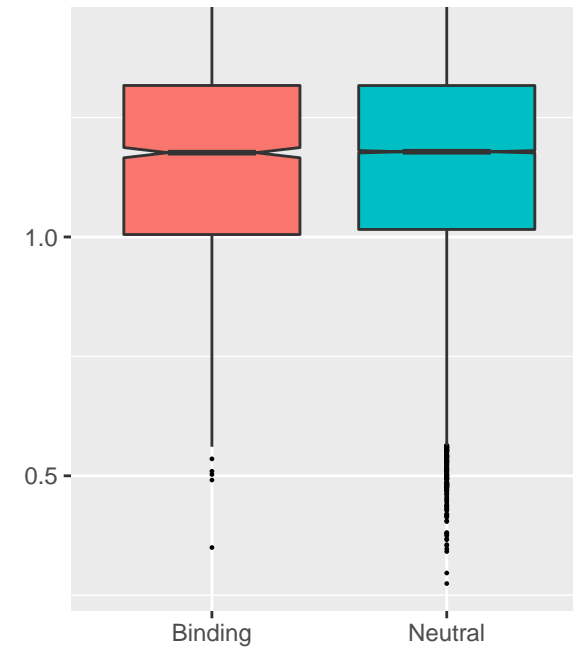
peptide, jsd_SE_sum
#(binding)=2173
p=4e-47



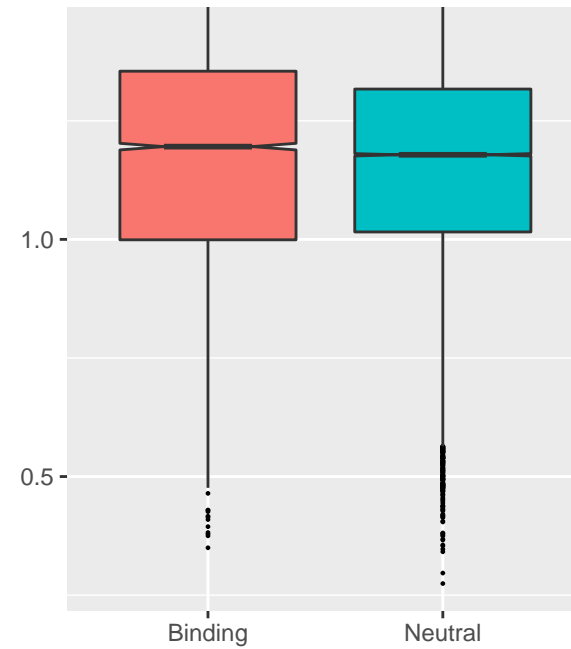
ion, jsd_SE_sum
#(binding)=1691
p=1



metabolite, jsd_SE_sum
#(binding)=2109
p=0.45



max, jsd_SE_sum
#(binding)=6486
p=1.3e-09



factor(label) Binding Neutral