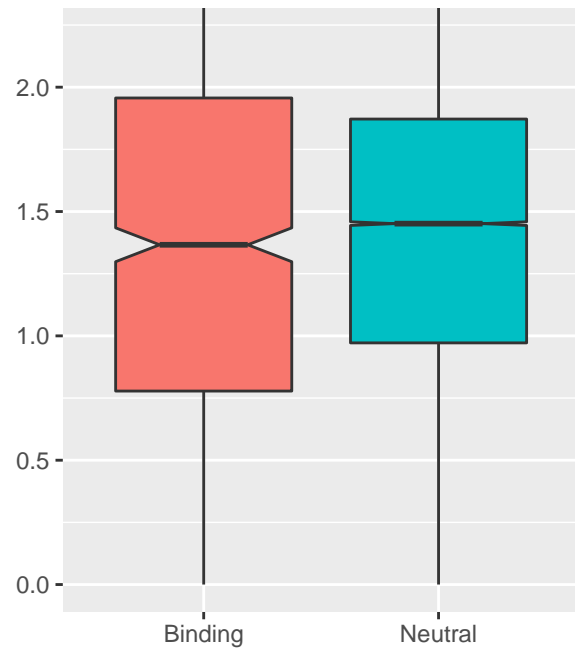
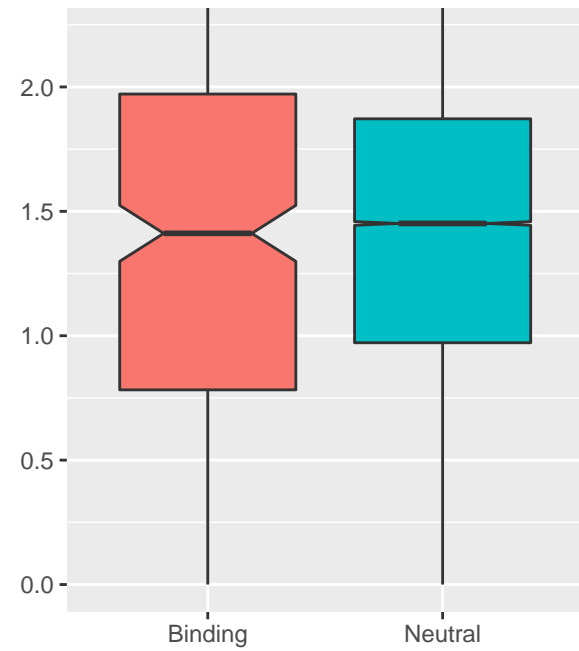


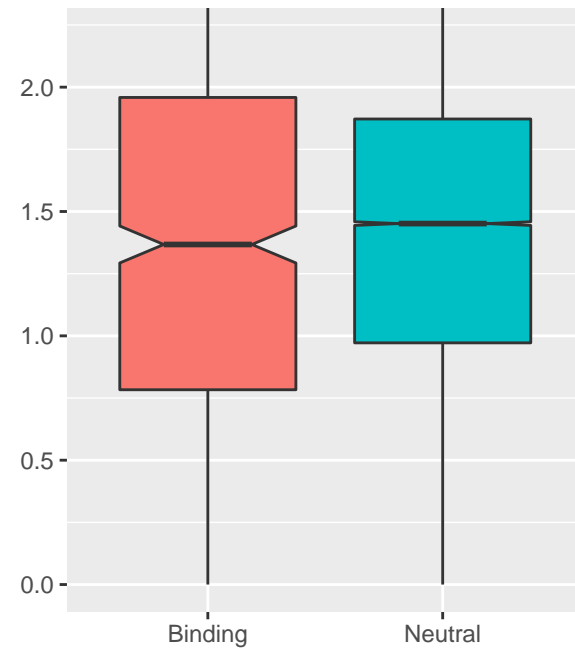
dna, jsd_mul_aa_ref_SE
#(binding)=749
p=0.99



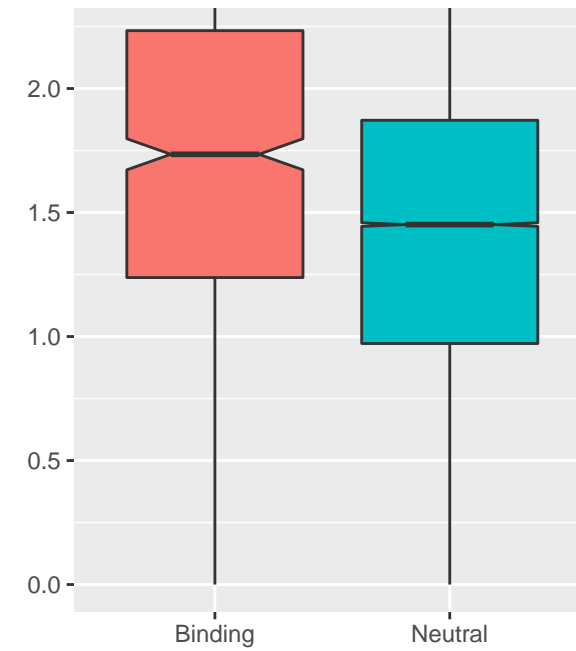
dnabase, jsd_mul_aa_ref_SE
#(binding)=279
p=0.71



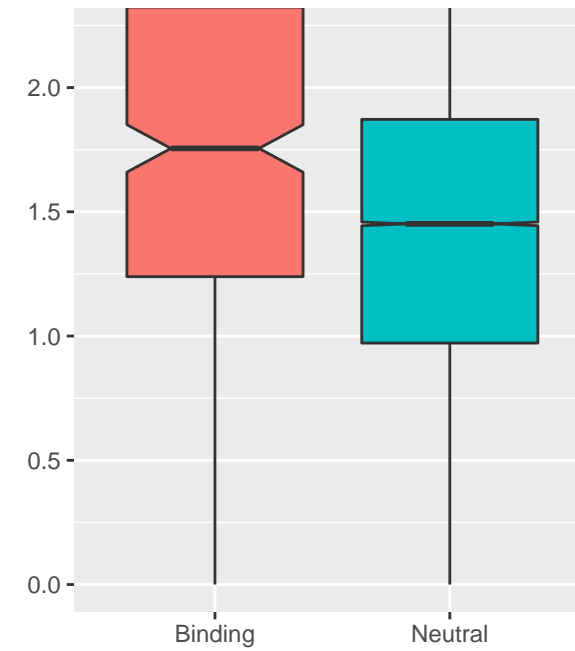
dnabackbone, jsd_mul_aa_ref_SE
#(binding)=625
p=0.97



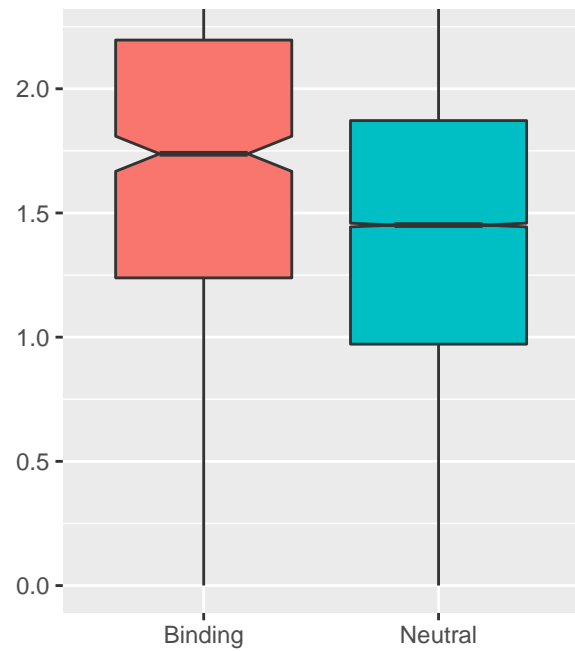
rna, jsd_mul_aa_ref_SE
#(binding)=629
p=3.2e-26



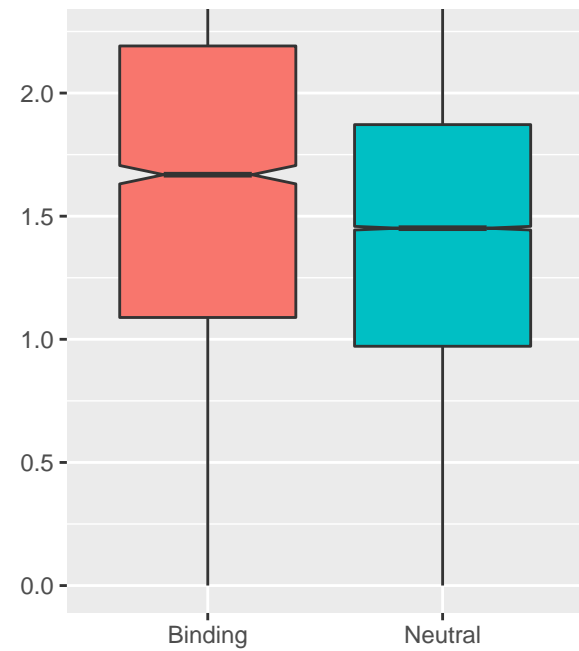
rnabase, jsd_mul_aa_ref_SE
#(binding)=322
p=5.4e-16



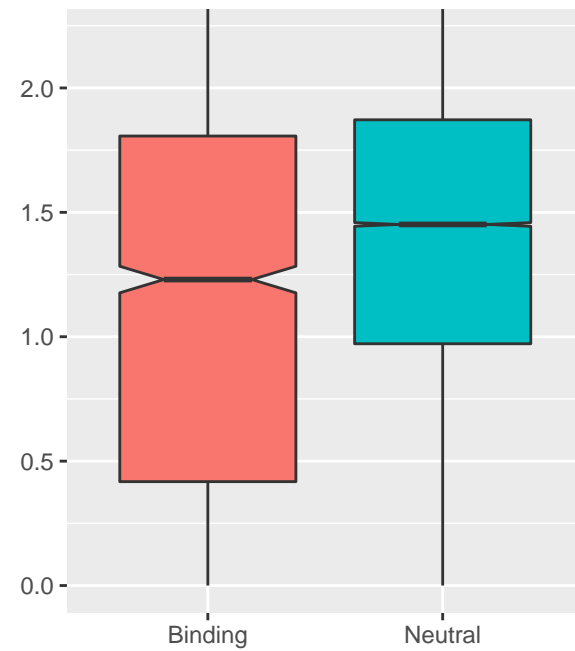
rnabackbone, jsd_mul_aa_ref_SE
#(binding)=458
p=2.2e-20



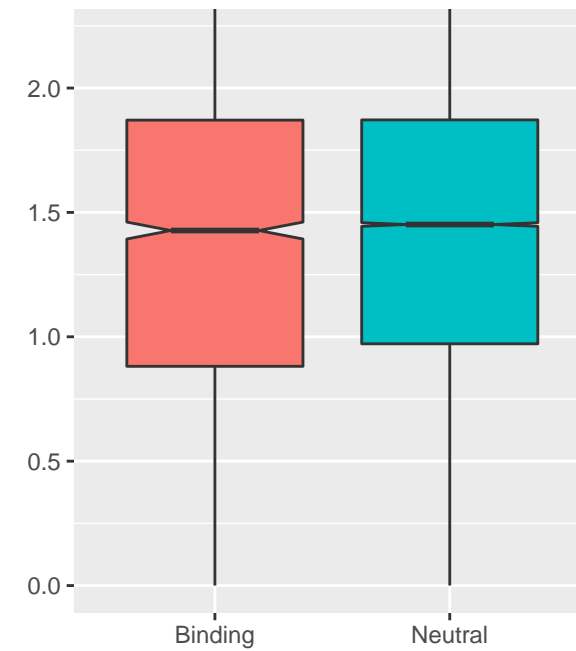
peptide, jsd_mul_aa_ref_SE
#(binding)=2173
p=6.2e-42



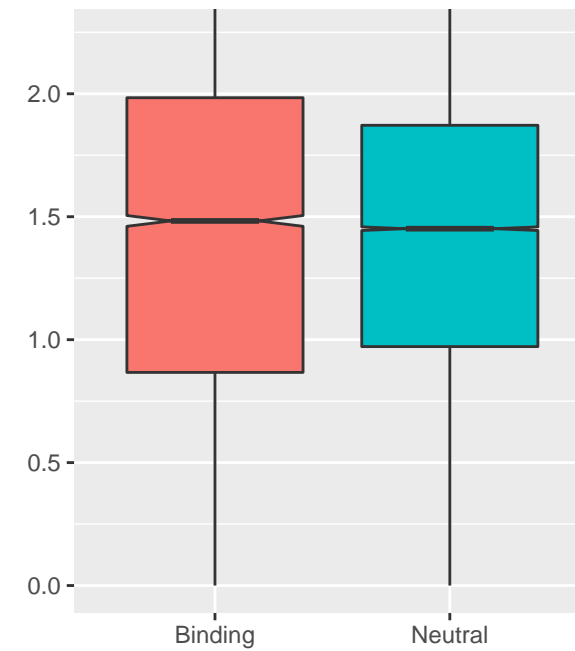
ion, jsd_mul_aa_ref_SE
#(binding)=1691
p=1



metabolite, jsd_mul_aa_ref_SE
#(binding)=2109
p=0.95



max, jsd_mul_aa_ref_SE
#(binding)=6486
p=0.0024



factor(label)  Binding  Neutral