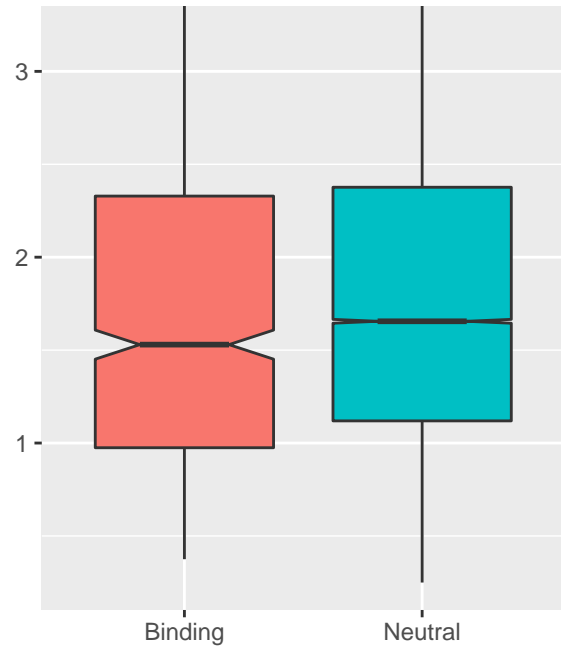
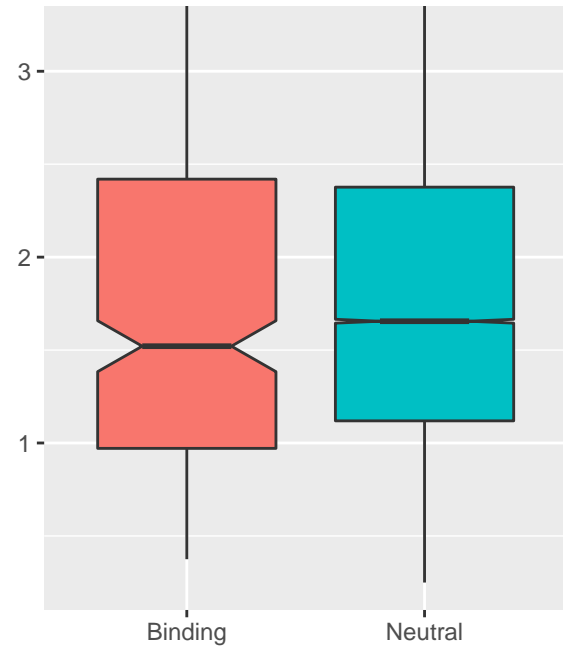


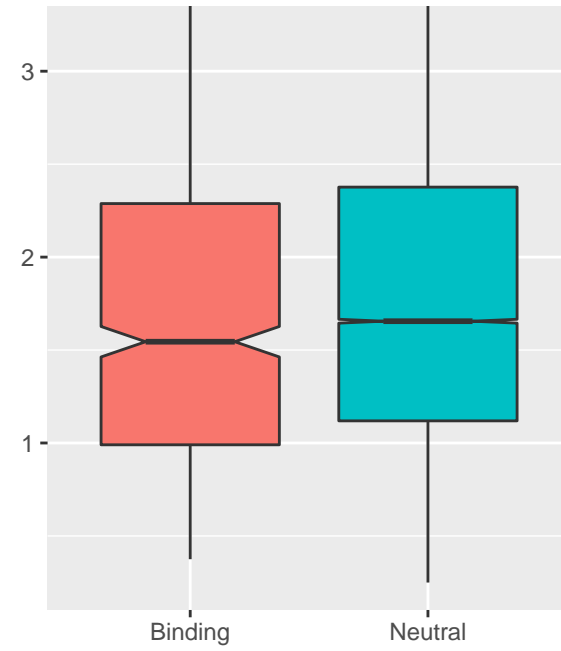
dna, jsd_100way_instances_major_#(binding)=749
p=1



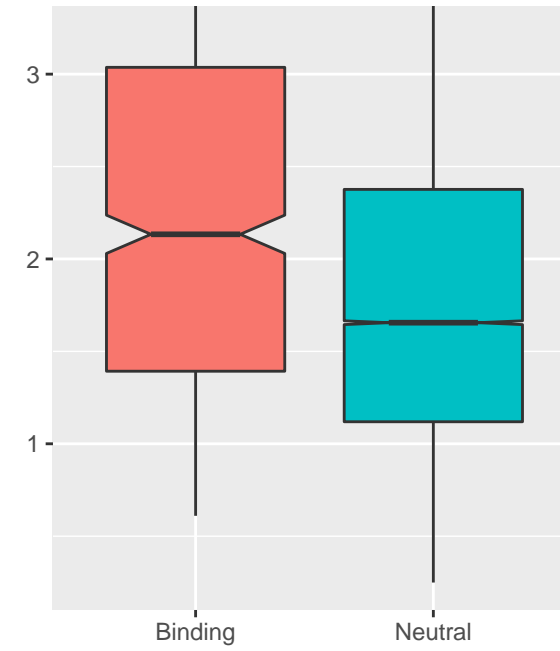
dnabase, jsd_100way_instances_r#(binding)=279
p=0.98



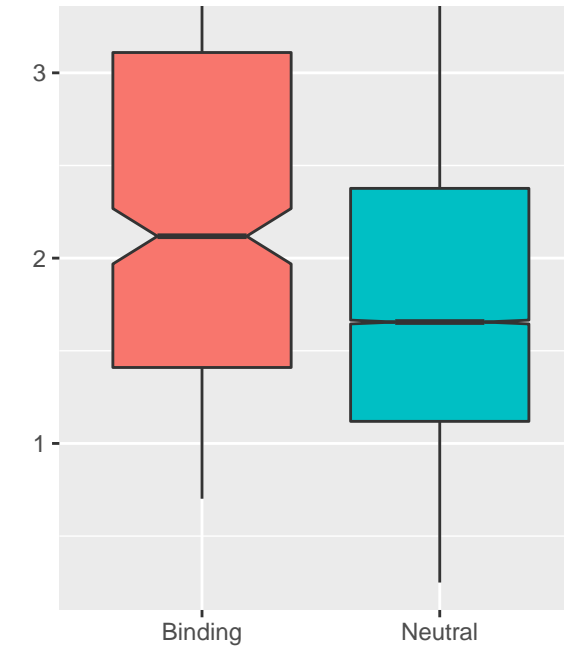
dnabackbone, jsd_100way_instanc#(binding)=625
p=1



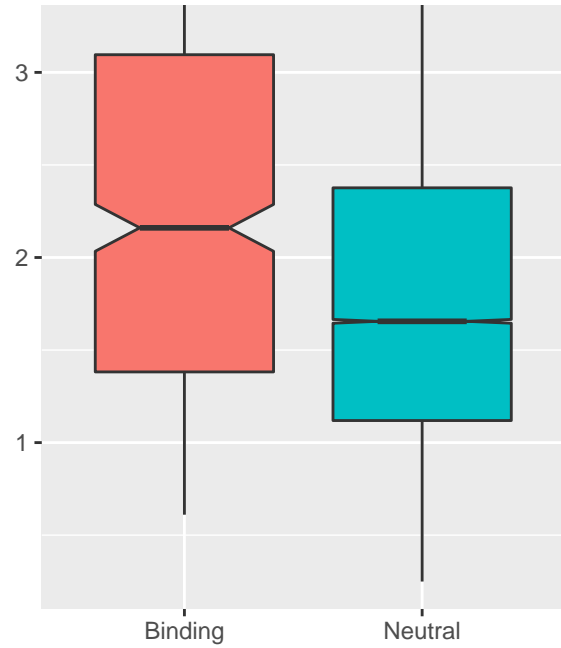
rna, jsd_100way_instances_major_#(binding)=629
p=1.9e-28



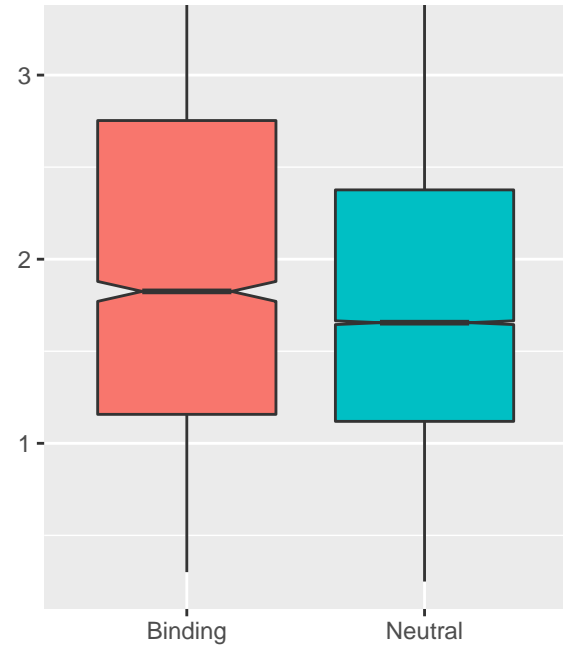
rnabase, jsd_100way_instances_m#(binding)=322
p=4.4e-17



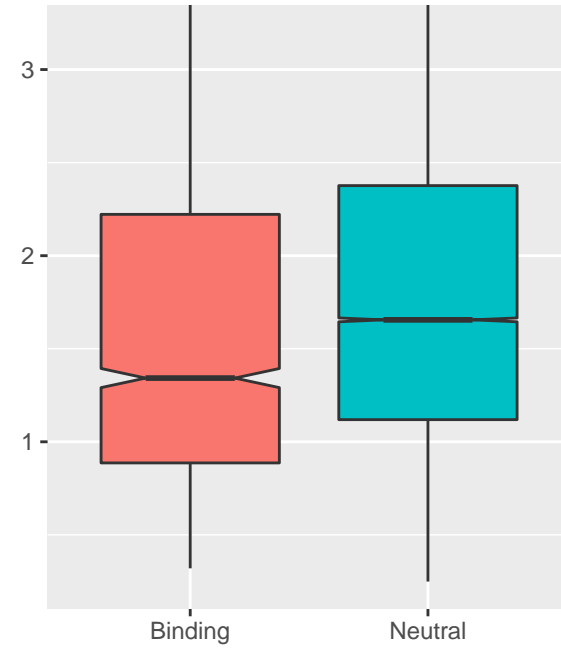
rnabackbone, jsd_100way_instanc#(binding)=458
p=2e-23



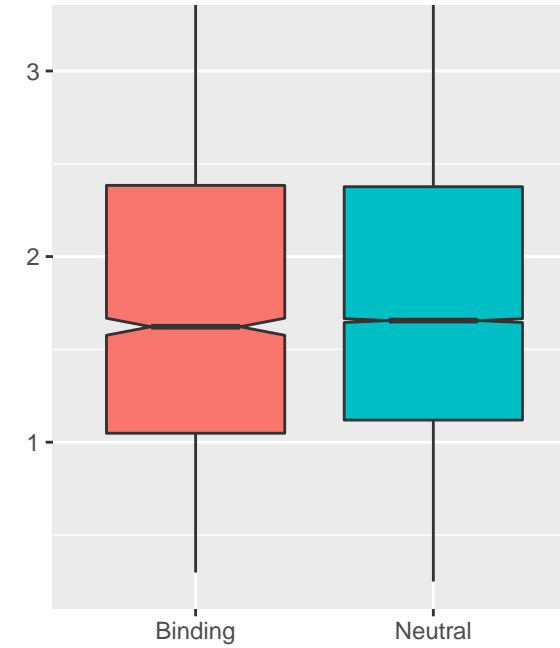
peptide, jsd_100way_instances_m#(binding)=2173
p=9.7e-14



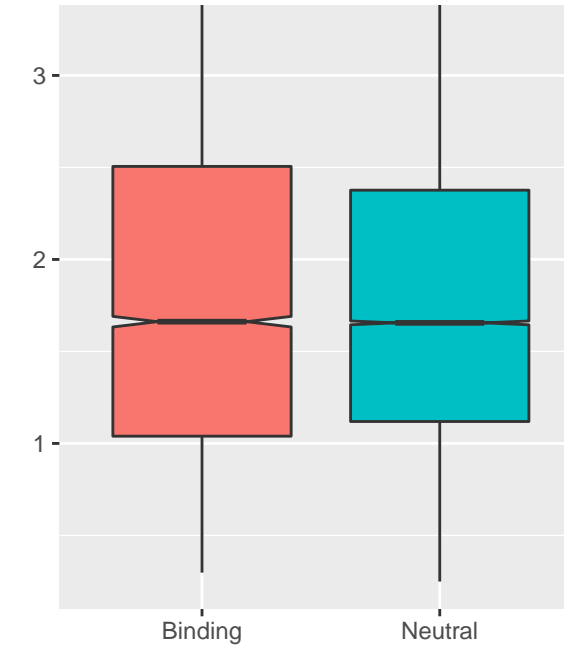
ion, jsd_100way_instances_major_#(binding)=1691
p=1



metabolite, jsd_100way_instances_#(binding)=2109
p=0.96



max, jsd_100way_instances_major_#(binding)=6486
p=0.41



factor(label) Binding Neutral