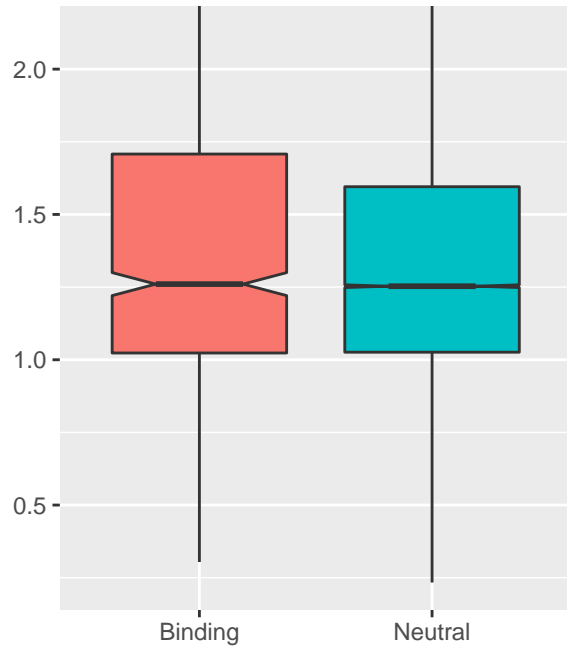
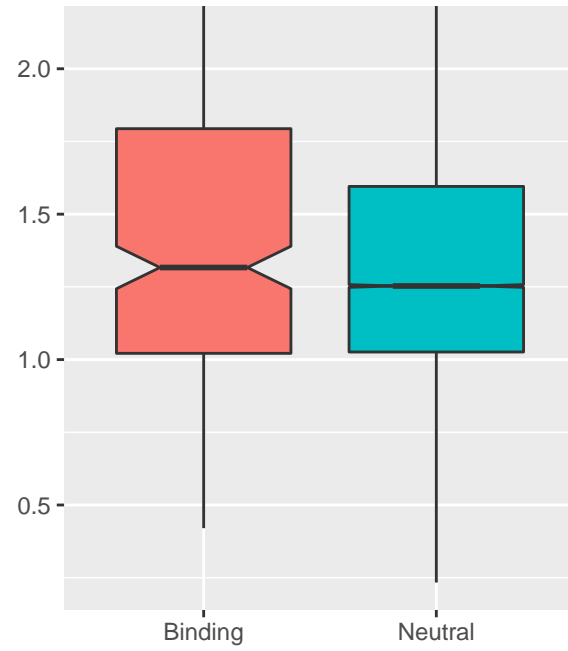


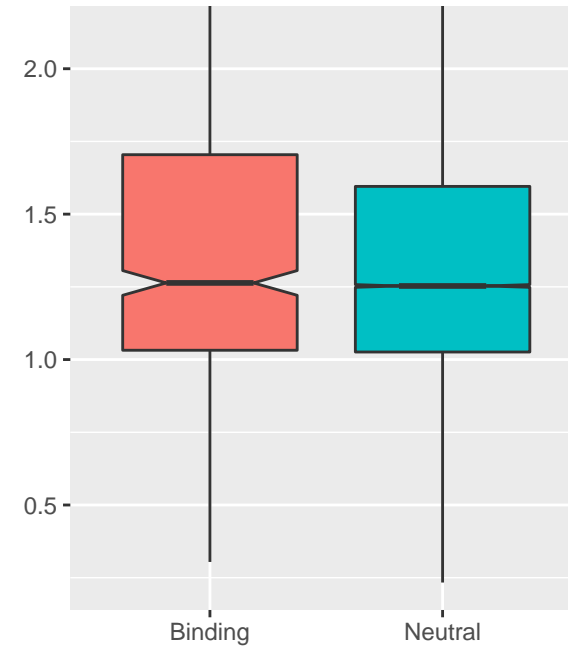
dna, jsds_ratio
#(binding)=749
p=0.074



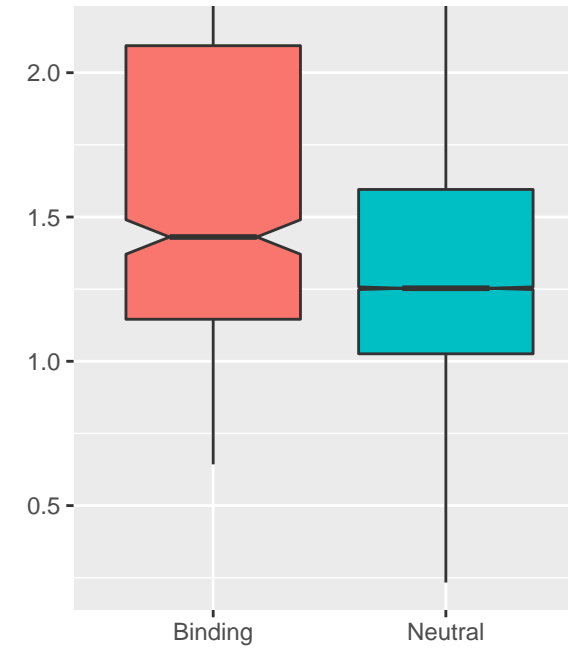
dnabase, jsds_ratio
#(binding)=279
p=0.0055



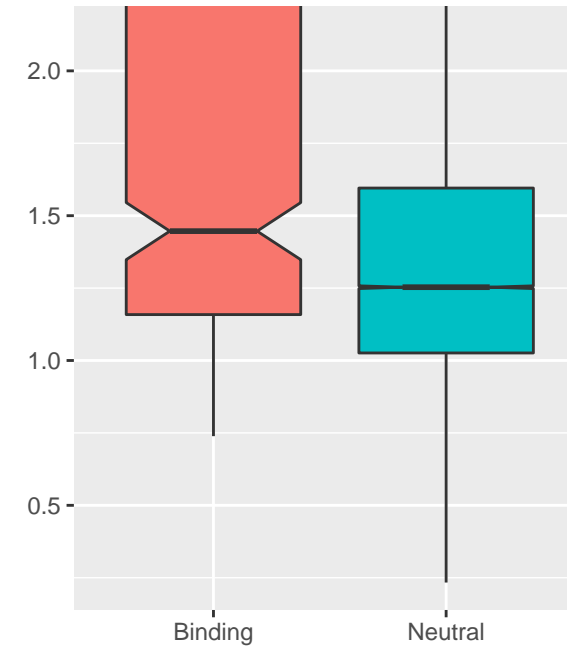
dnabackbone, jsds_ratio
#(binding)=625
p=0.063



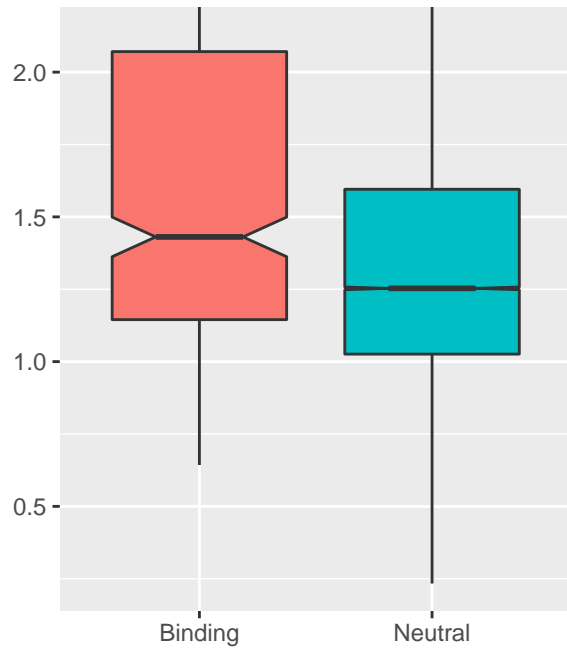
rna, jsds_ratio
#(binding)=629
p=2.9e-25



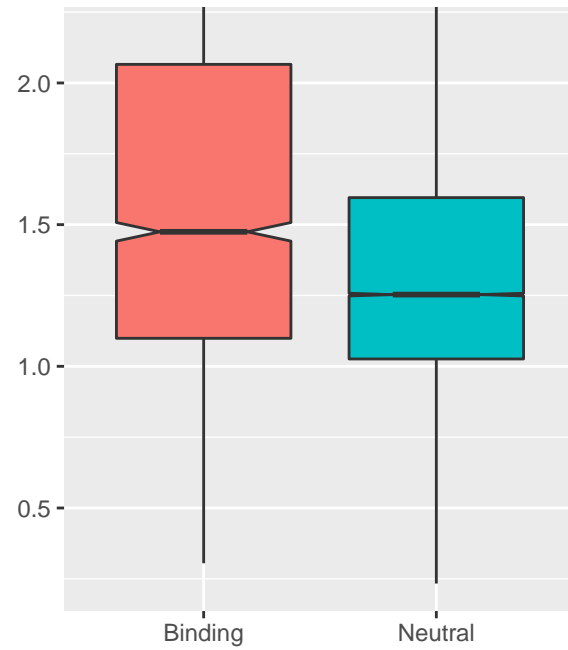
rnabase, jsds_ratio
#(binding)=322
p=7e-16



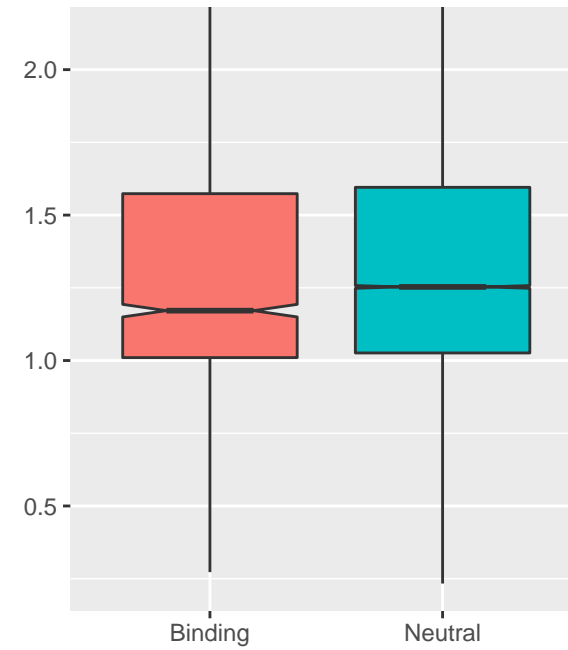
rnabackbone, jsds_ratio
#(binding)=458
p=2.5e-18



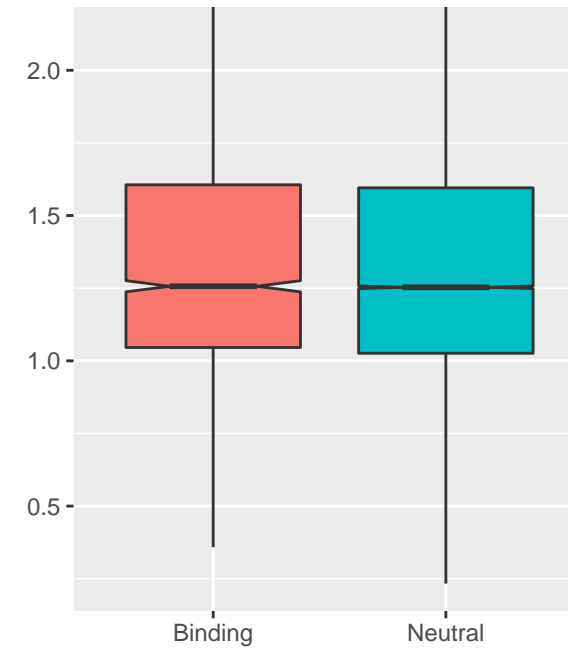
peptide, jsds_ratio
#(binding)=2173
p=2.1e-60



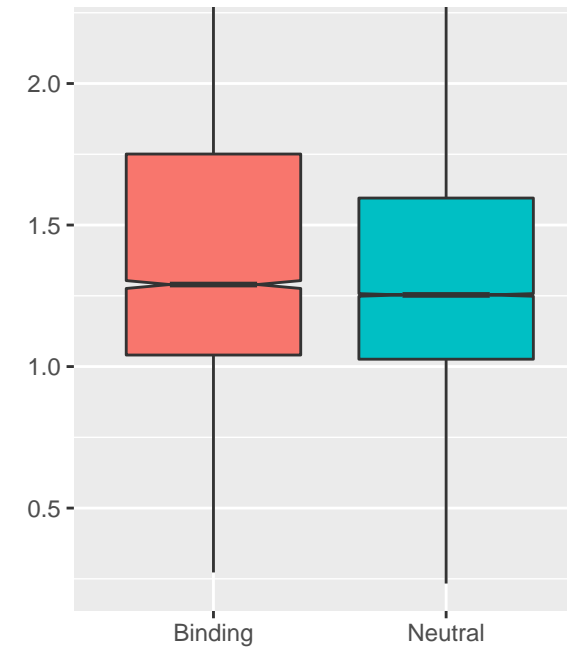
ion, jsds_ratio
#(binding)=1691
p=1



metabolite, jsds_ratio
#(binding)=2109
p=0.024



max, jsds_ratio
#(binding)=6486
p=1.8e-20



factor(label) Binding Neutral