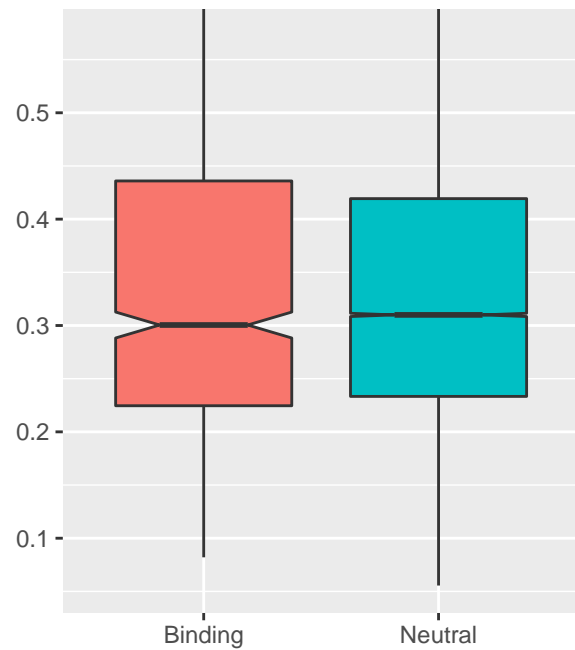
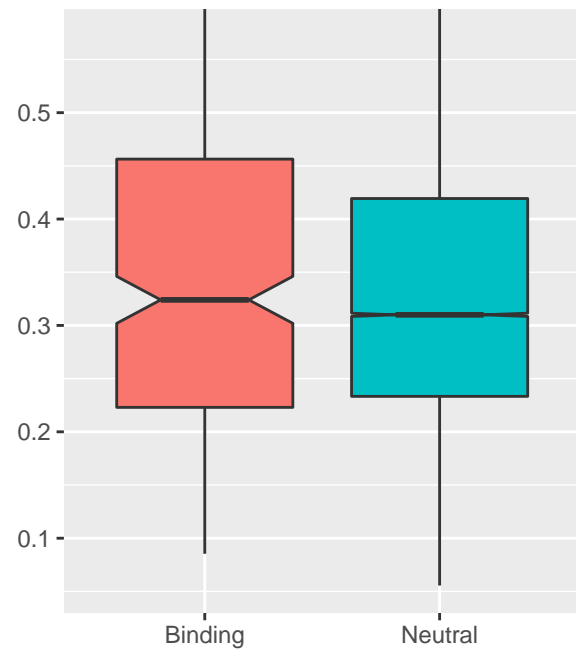


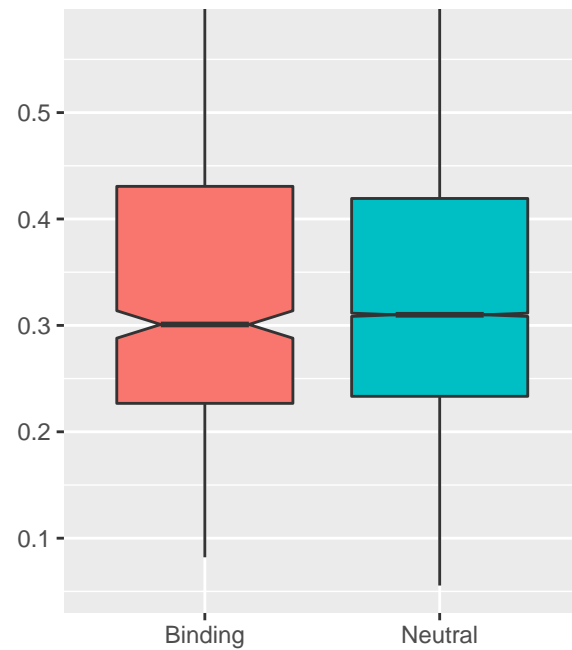
dna, jsd\_SE\_diff\_ratio  
#(binding)=749  
p=0.79



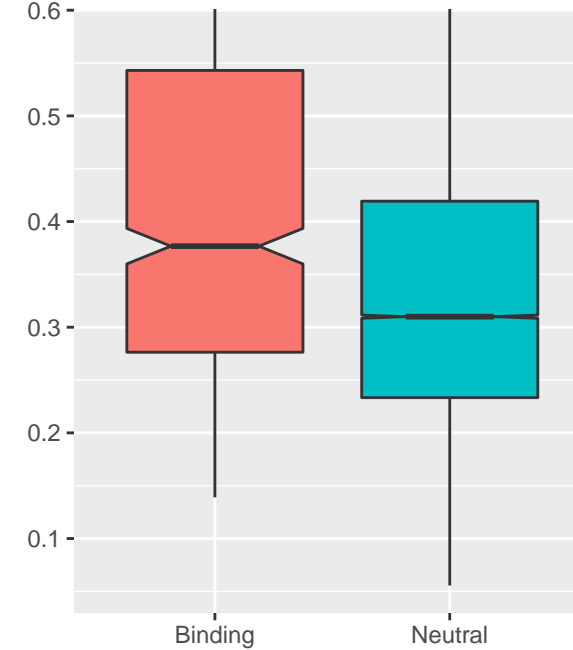
dnabase, jsd\_SE\_diff\_ratio  
#(binding)=279  
p=0.19



dnabackbone, jsd\_SE\_diff\_ratio  
#(binding)=625  
p=0.73



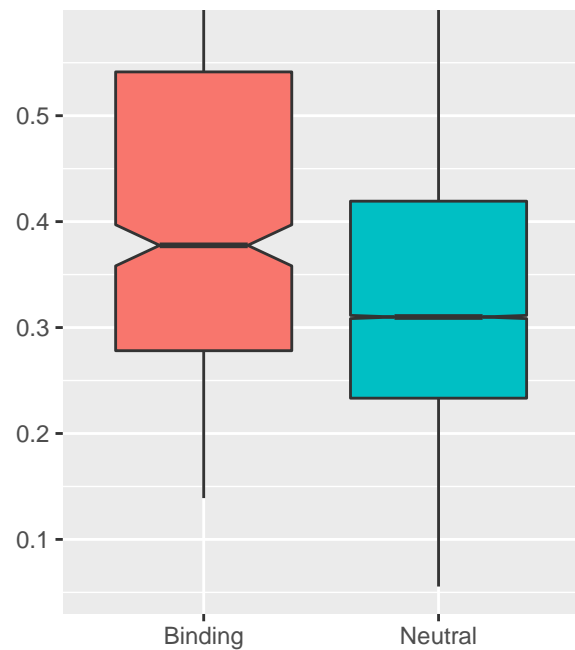
rna, jsd\_SE\_diff\_ratio  
#(binding)=629  
p=8.6e-28



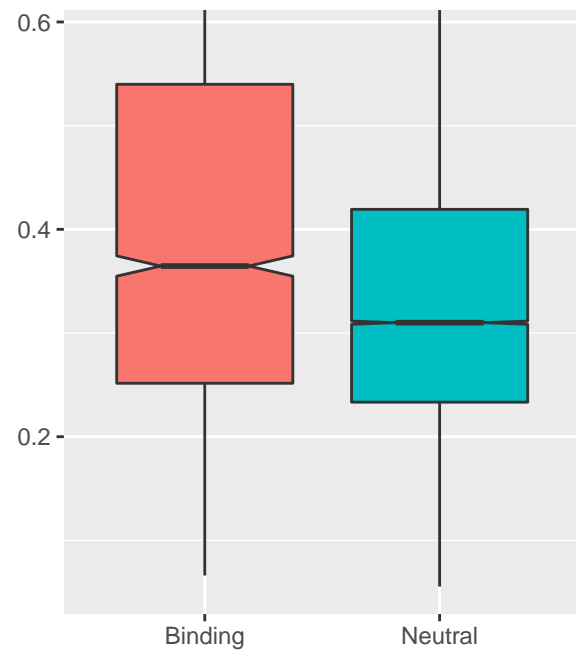
rnabase, jsd\_SE\_diff\_ratio  
#(binding)=322  
p=2.9e-17



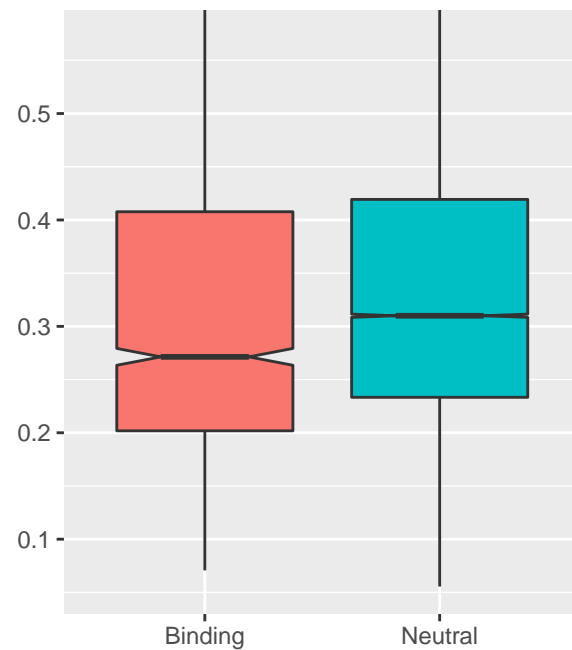
rnabackbone, jsd\_SE\_diff\_ratio  
#(binding)=458  
p=1.6e-21



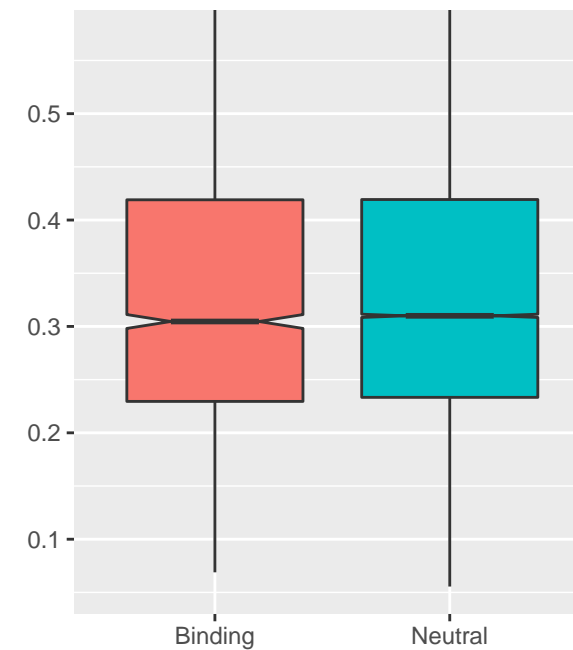
peptide, jsd\_SE\_diff\_ratio  
#(binding)=2173  
p=1.2e-41



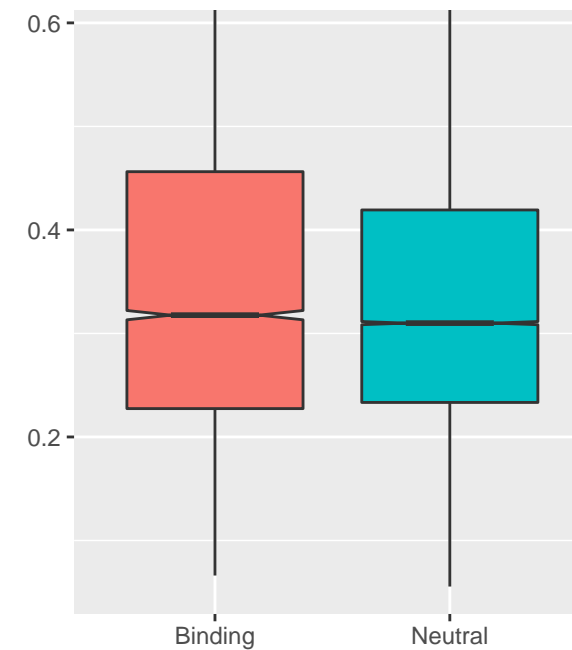
ion, jsd\_SE\_diff\_ratio  
#(binding)=1691  
p=1



metabolite, jsd\_SE\_diff\_ratio  
#(binding)=2109  
p=0.77



max, jsd\_SE\_diff\_ratio  
#(binding)=6486  
p=1.2e-06



factor(label)  Binding  Neutral