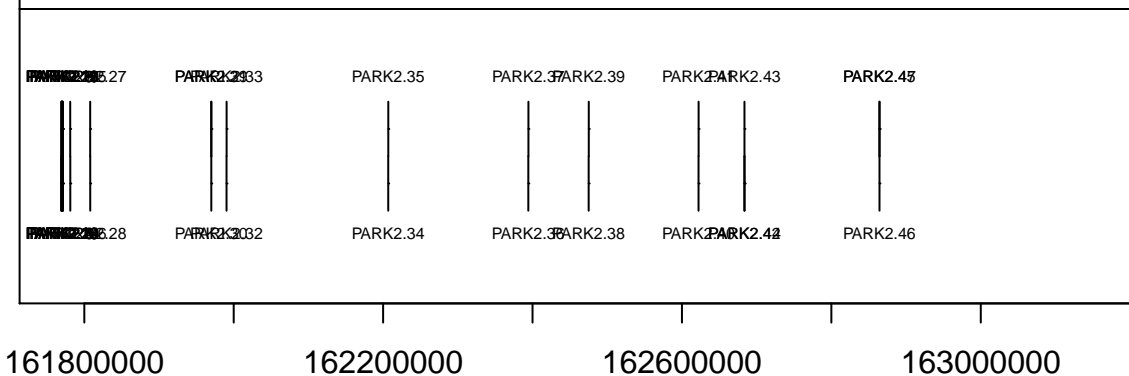
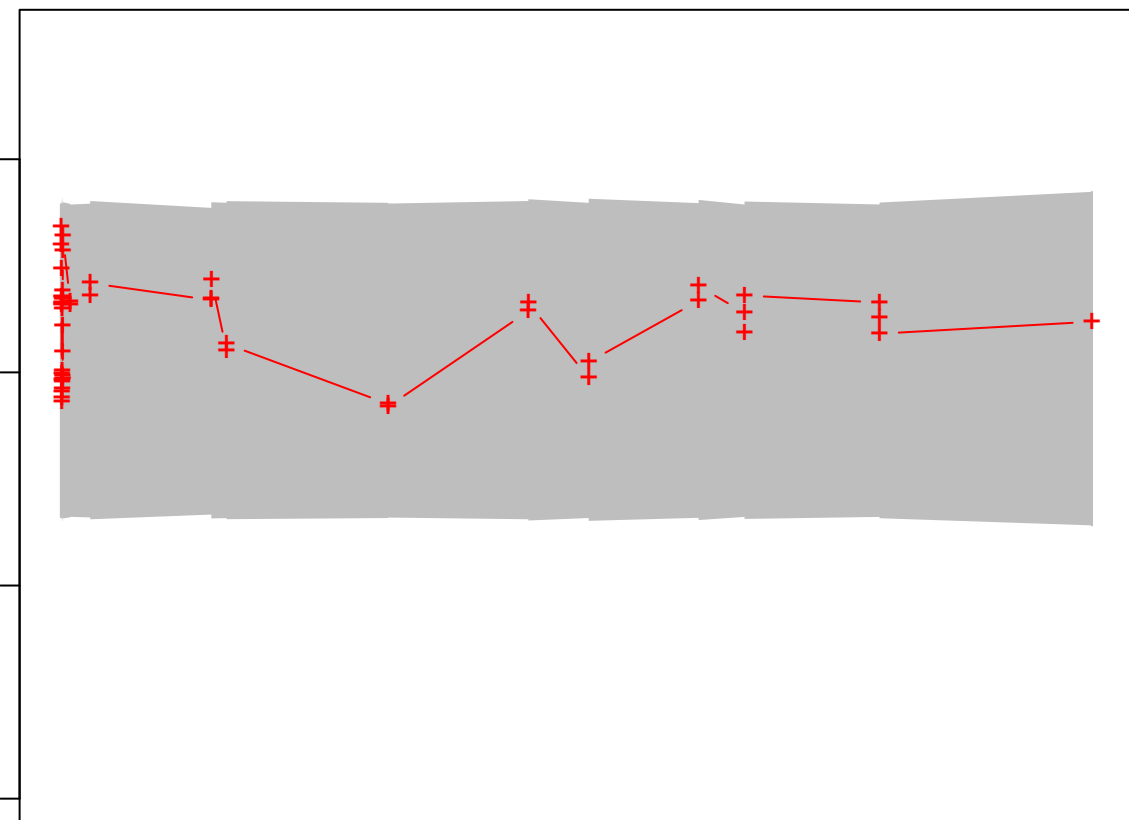


# PARK2 gene

Observed by expected read ratio

1.5  
1.0  
0.5  
0.0



161800000

162200000

162600000

163000000