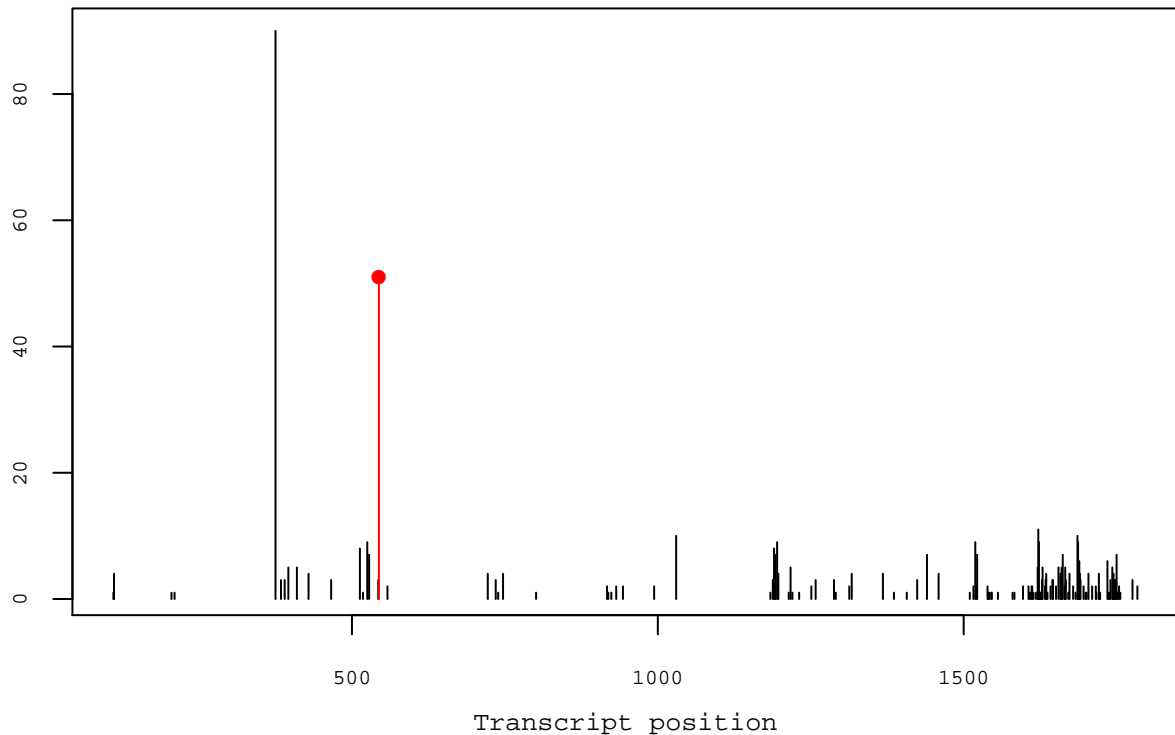


PggD7691_c15_g1_i1

5' TCCCTCTCGATATTGGCTCGGCTCAATTCCT 3'
 |||||
 3' GCTATAACCGTGCCGAGTT 5'



Cleavage site: 544 Tag abundance: 51 Weighted abundance: 51 Category: 2
sRNA abundance: 68 Alignment score: 2 MFE ratio: 0.878 p-value: 0.04