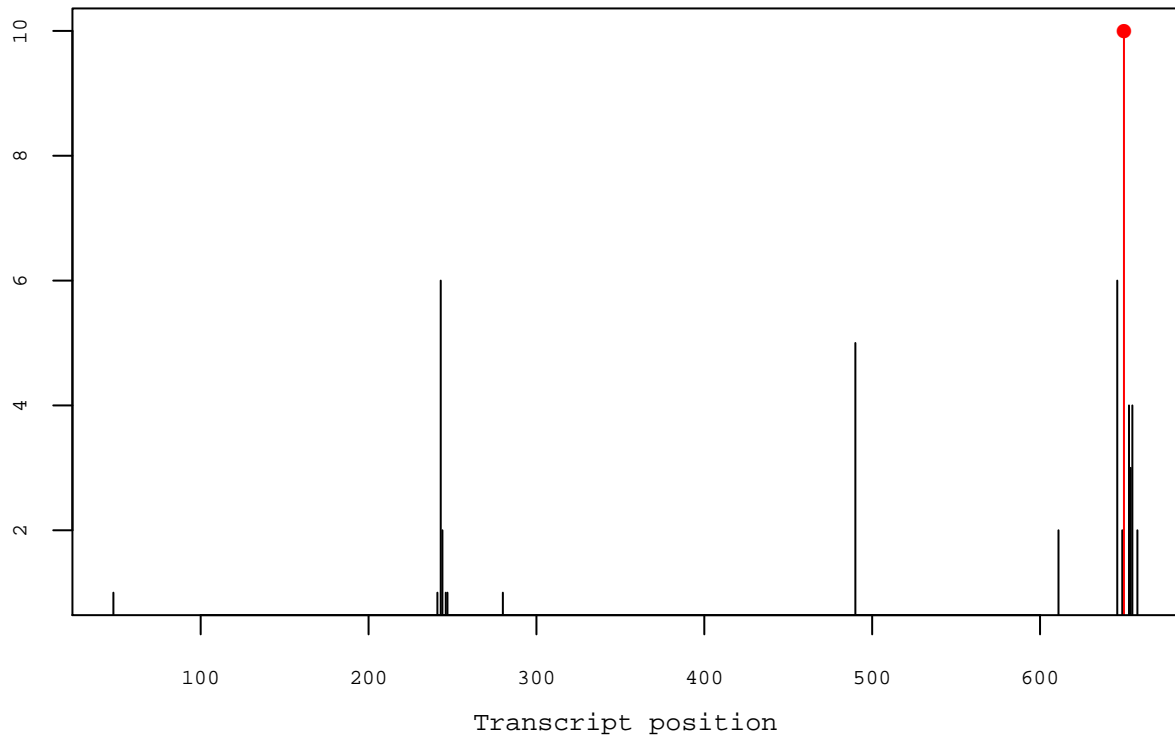


PggD770_c9_g1_i1

5' TACGGACACTGCATGCATCTCTGCATGCGGTG '3
||| ||| o ||| ||| ||| ||| o |||
3' GCCTGTGGCGTACGTAGAGATGTA '5

Fragment Abundance



Cleavage site: 650 Tag abundance: 10 Weighted abundance: 3.333 Category: 0
sRNA abundance: 13 Alignment score: 1.5 MFE ratio: 0.953 p-value: 0.009