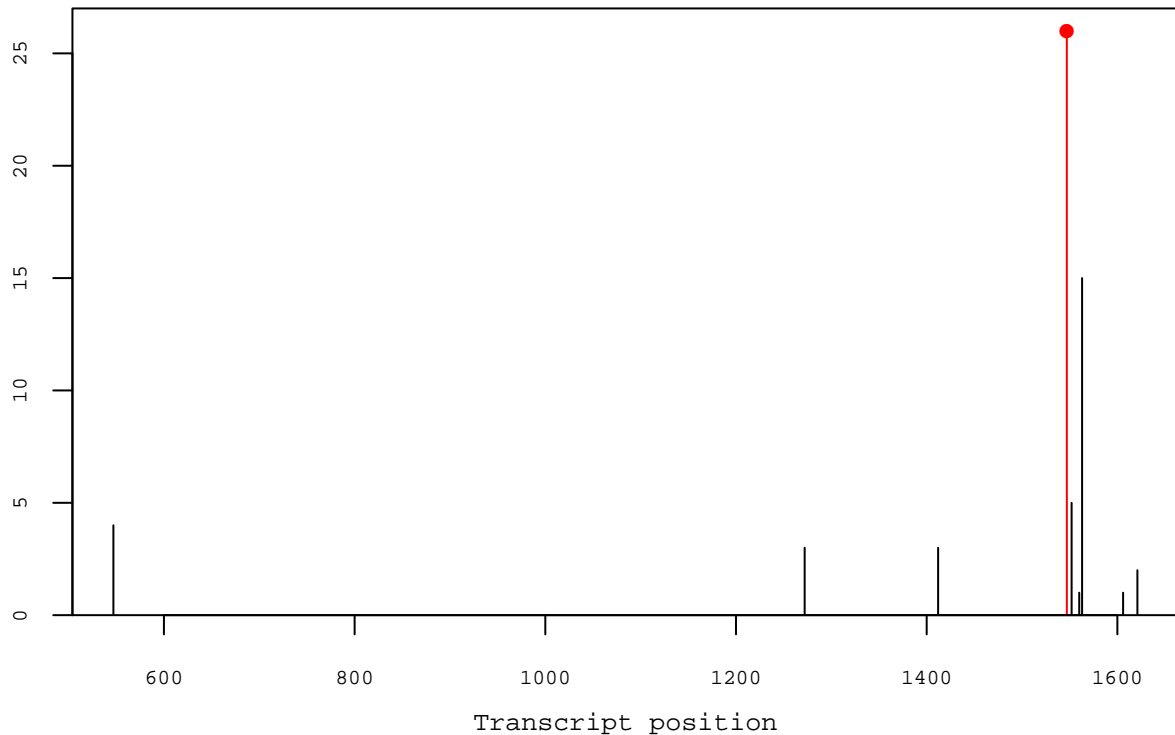


PggD7826_c63_g1_i1

5' CCTGTAACCTGTTTGCAGGTTACAGACGTACG '3
||| |||||
3' TTGTACAAACGTCCAATGTCT '5

Fragment Abundance



Cleavage site: 1547 Tag abundance: 26 Weighted abundance: 1.083 Category: 0
sRNA abundance: 10 Alignment score: 1 MFE ratio: 0.869 p-value: 0.007