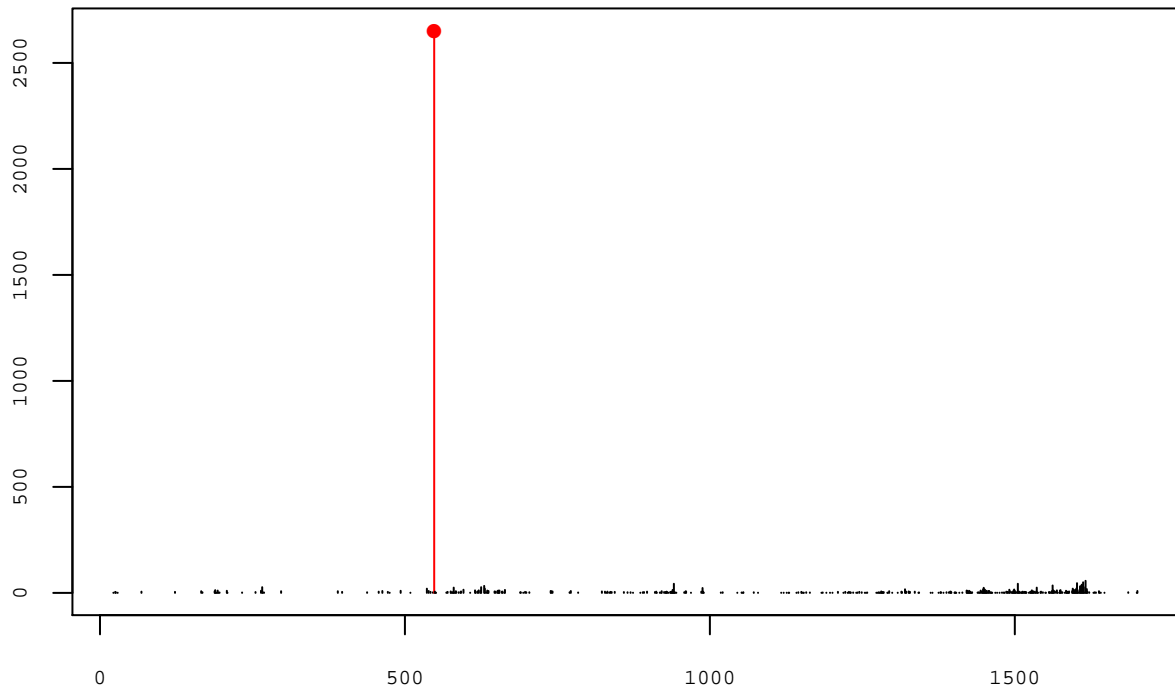


PggD9652_c39_g1_i1

5' CTGCGTTTCAGTGTGCAAAAGATGGAGCTCTGT '3
 ||||| |||||○
3' GTCACACGTTCTCTACCTT '5

Fragment Abundance



Transcript position

Cleavage site: 548 Tag abundance: 2651 Weighted abundance: 2651 Category: 0
sRNA abundance: 8 Alignment score: 2.5 MFE ratio: 0.809 p-value: 0.001