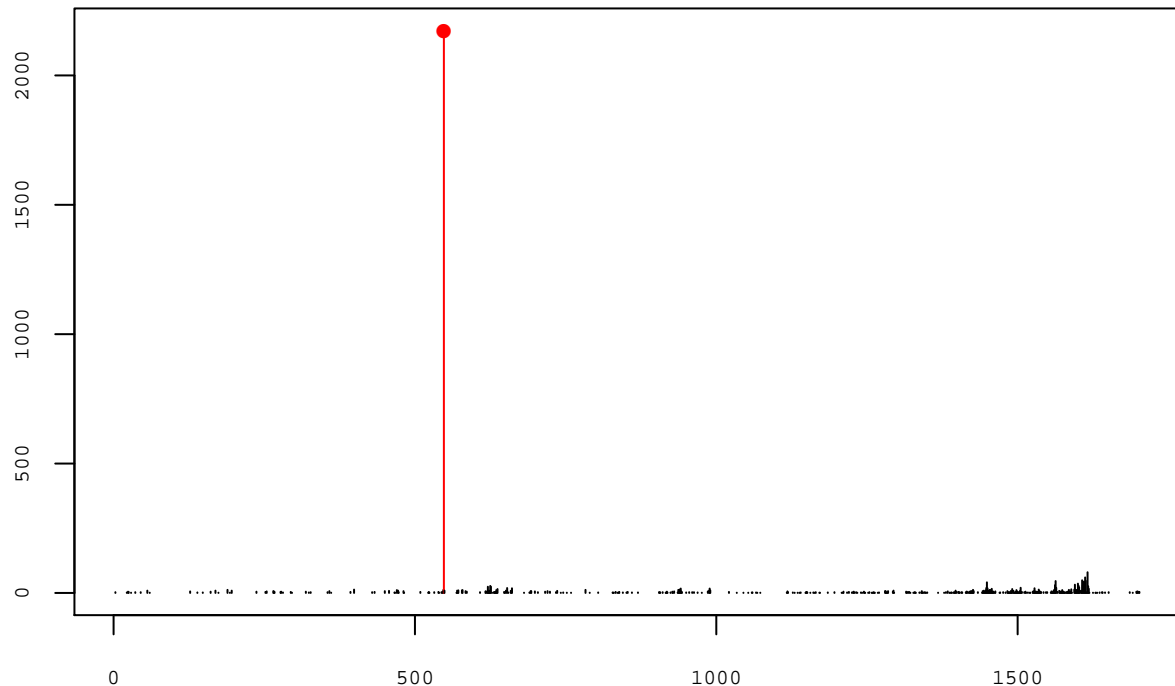


PggD9652_c39_g1_i1

5' CTGCGTTTCAGTGTGCAAAAGATGGAGCTCTGT '3
|o||||||||| |||||o
3' AGGTCACACGTTCTCTACCTT '5

Fragment Abundance



Transcript position

Cleavage site: 548 Tag abundance: 2172 Weighted abundance: 2172 Category: 0
sRNA abundance: 473 Alignment score: 3 MFE ratio: 0.76 p-value: 0.002