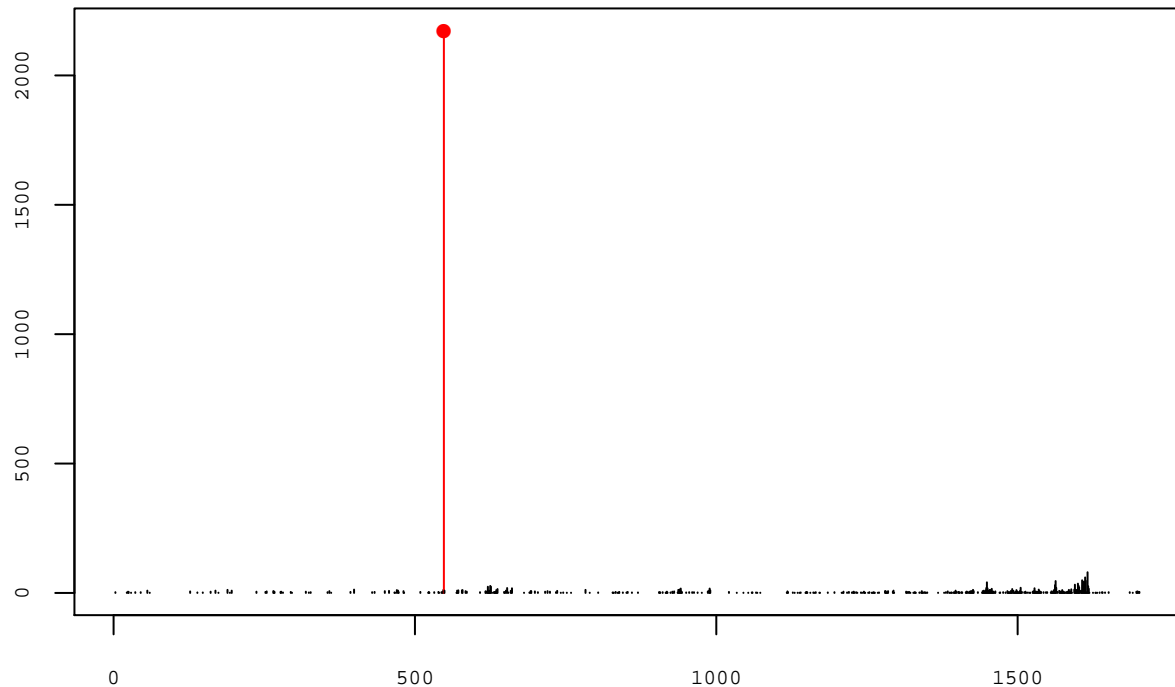


PggD9652\_c39\_g1\_i1

5' CTGCGTTTCAGTGTGCAAAAGATGGAGCTCTGT '3  
o||||||| | | | | | o  
3' GGTACACAGTTCTCTACCTT '5

Fragment Abundance



Transcript position

Cleavage site: 548 Tag abundance: 2172 Weighted abundance: 2172 Category: 0  
sRNA abundance: 134 Alignment score: 3 MFE ratio: 0.777 p-value: 0.002