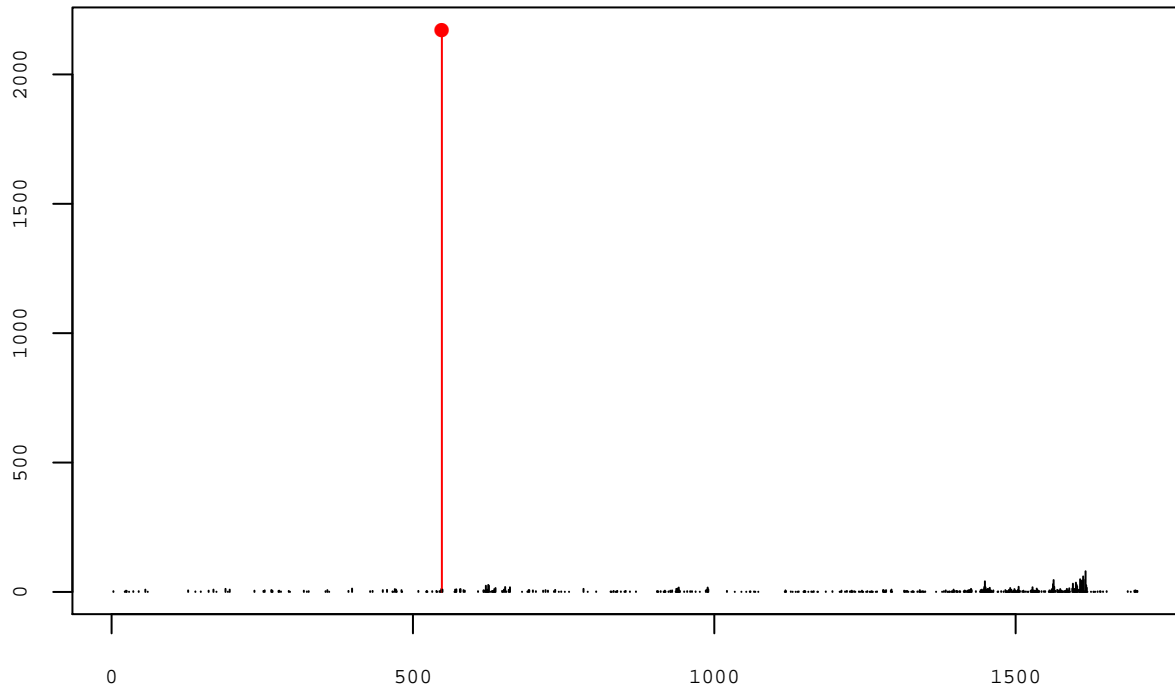


PggD9652_c39_g1_i1

5' CTGCGTTCAGTGTGCAAAAGATGGAGCTCTGT '3
 ||||| |||||○
3' GTCACACGTTCTCTACCTT '5

Fragment Abundance



Transcript position

Cleavage site: 548	Tag abundance: 2172	Weighted abundance: 2172	Category: 0
sRNA abundance: 14	Alignment score: 2.5	MFE ratio: 0.809	p-value: 0.001