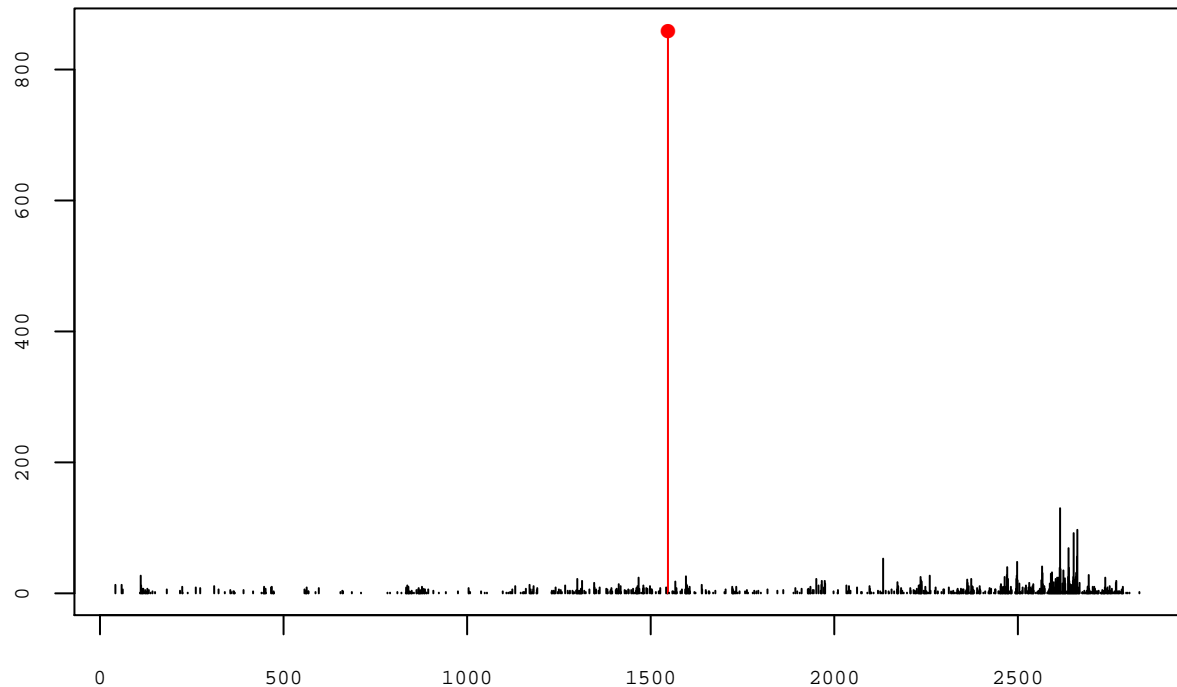


PggD4130_c0_g1_i1

5' GTTCCCTCCAGTA-GCAAGAGATGGACAGATGG '3
||||| |||||
3' GGTACACGTTCTCTACCTT '5

Fragment Abundance



Transcript position

Cleavage site: 1547	Tag abundance: 859	Weighted abundance: 429.5	Category: 0
sRNA abundance: 134	Alignment score: 3	MFE ratio: 0.753	p-value: 0.001