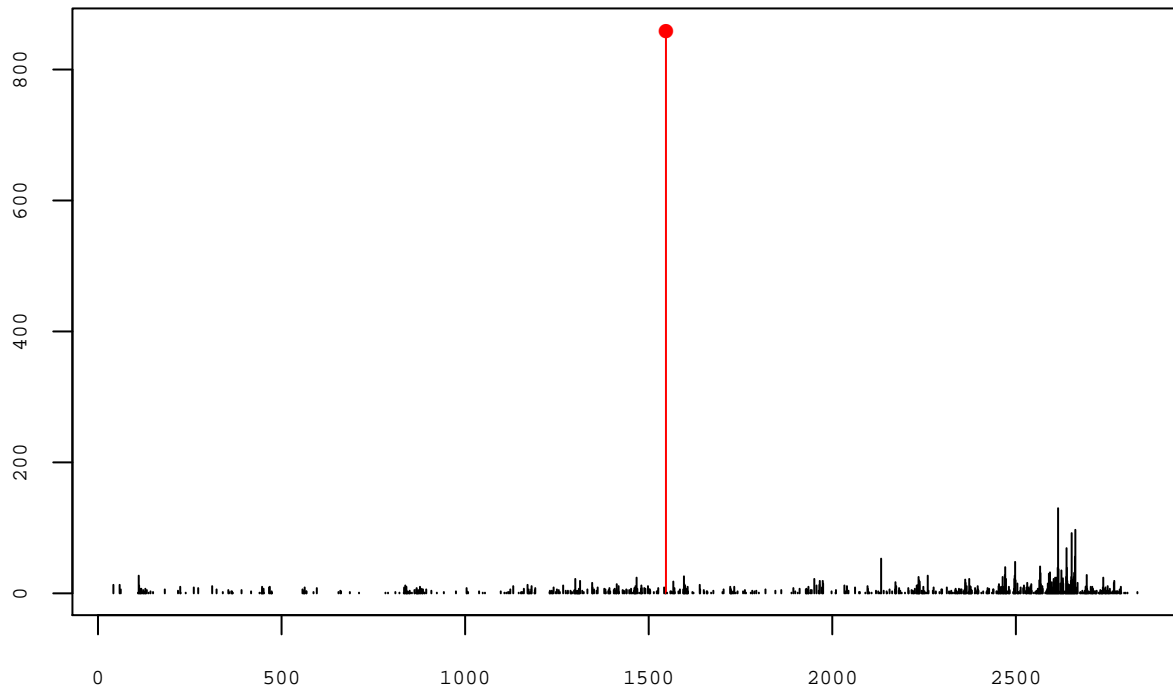


PggD4130\_c0\_g1\_i1

5' GTTCCCTCCAGTA-GCAAGAGATGGACAGATGG '3  
||||| |||||  
3' GTCACACGTTCTCTACCTT '5

Fragment Abundance



Transcript position

Cleavage site: 1547	Tag abundance: 859	Weighted abundance: 429.5	Category: 0
sRNA abundance: 14	Alignment score: 3	MFE ratio: 0.728	p-value: 0.001