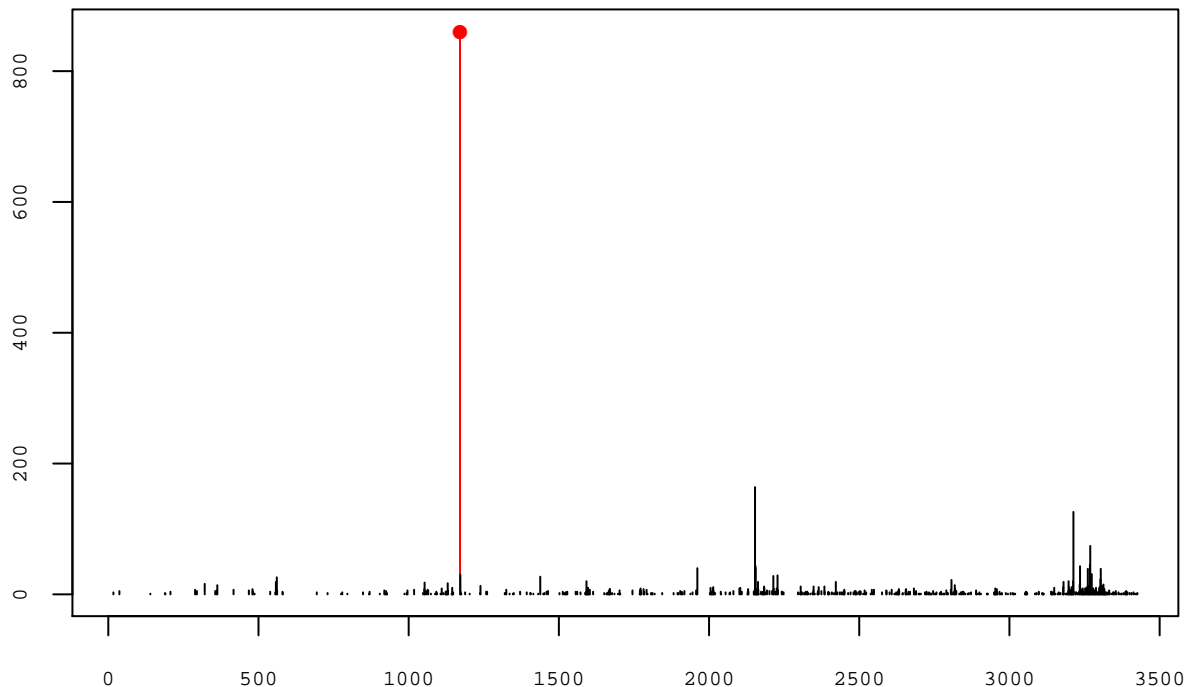


PggD9924\_c6\_g1\_i1

5' GCTTGTGGAATGAAGCCTGGTCCGGATTCCA '3  
|||||  
3' CCCTTACTTCGGACCAGGCT '5

Fragment Abundance



Transcript position

Cleavage site: 1171	Tag abundance: 860	Weighted abundance: 860	Category: 0
sRNA abundance: 730	Alignment score: 1.5	MFE ratio: 0.929	p-value: 0