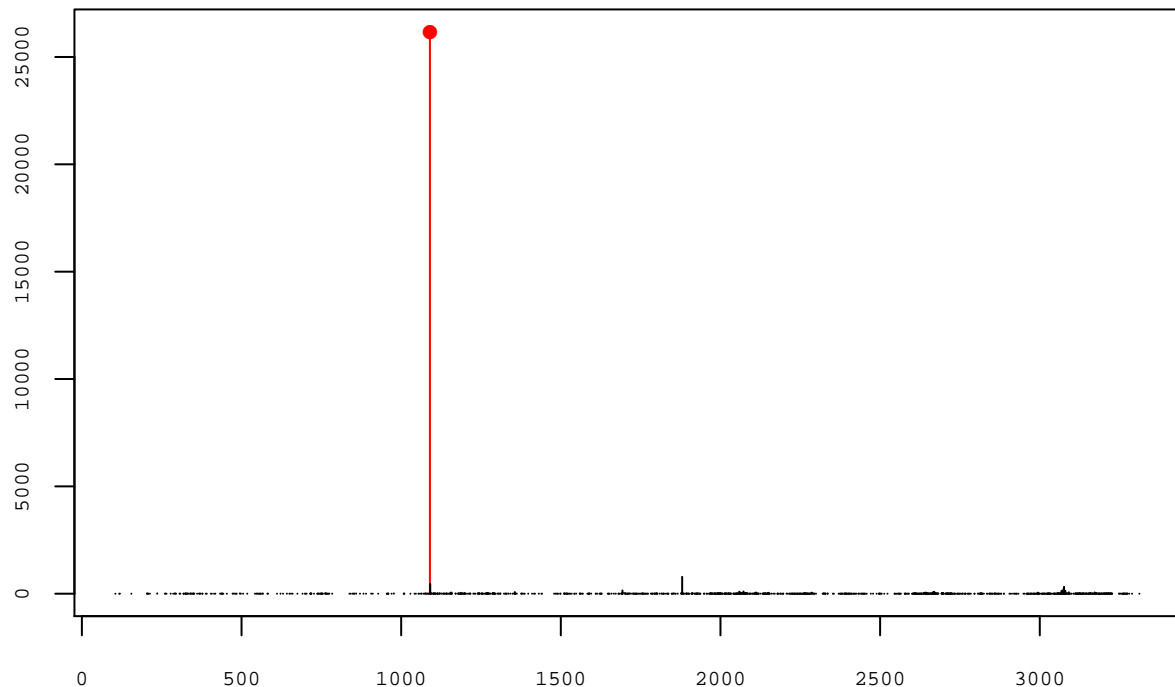


PggD7141\_c1\_g1\_i1

5' GATGGTTGGTATGAAGCCTGGTCCGGATTCTA '3  
|| |||||  
3' CCCTTACTTCGGACCAGGCT '5

Fragment Abundance



Transcript position

Cleavage site: 1090	Tag abundance: 26168	Weighted abundance: 13084	Category: 0
sRNA abundance: 730	Alignment score: 2.5	MFE ratio: 0.819	p-value: 0.001