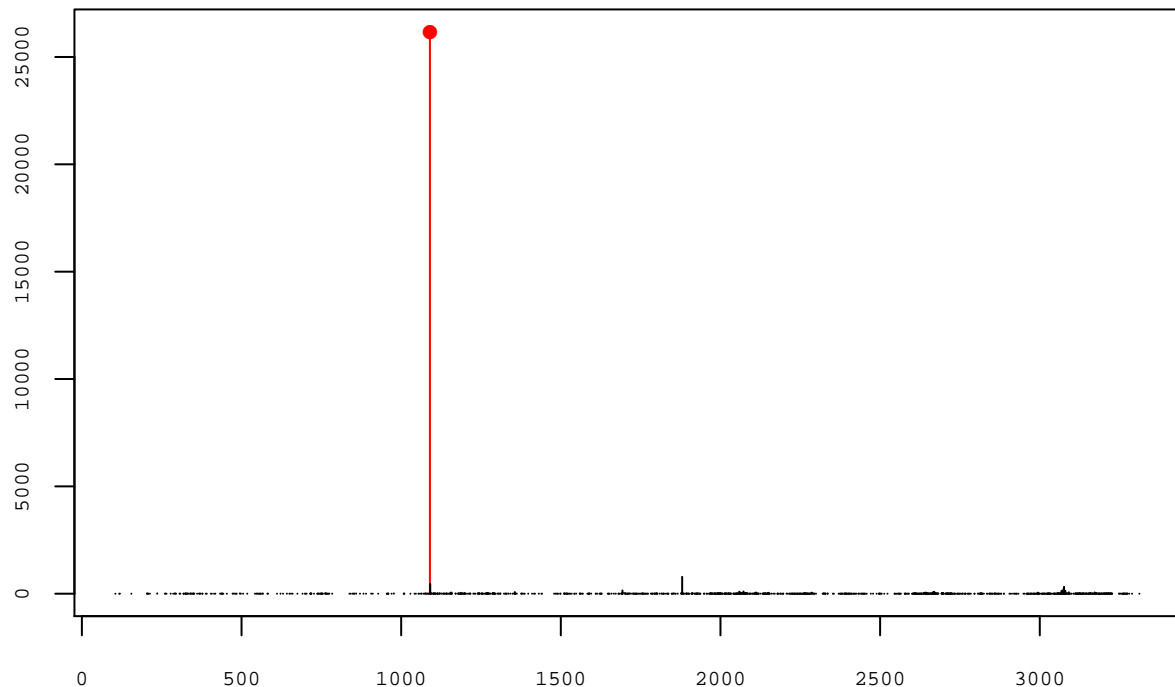


PggD7141\_c1\_g1\_i1

5' GATGGTTGGTATGAAGCCTGGTCCGGATTCTA '3  
|| ||||| ||||| |||||  
3' CCCCTTACTTCGGACCAGGCT '5

Fragment Abundance



Transcript position

Cleavage site: 1090 Tag abundance: 26168 Weighted abundance: 13084 Category: 0  
sRNA abundance: 44559 Alignment score: 3.5 MFE ratio: 0.757 p-value: 0.001