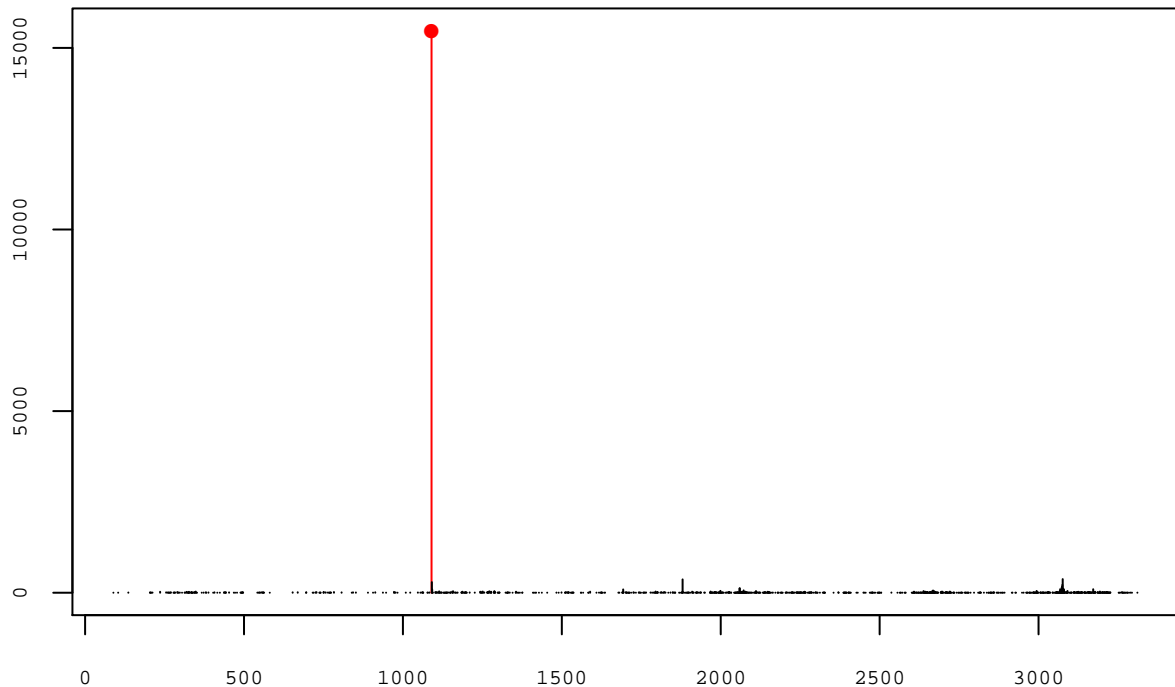


PggD7141_c1_g1_i1

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

Fragment Abundance



Transcript position

Cleavage site: 1090 Tag abundance: 15468 Weighted abundance: 7734 Category: 0
sRNA abundance: 19192 Alignment score: 3.5 MFE ratio: 0.757 p-value: 0.001