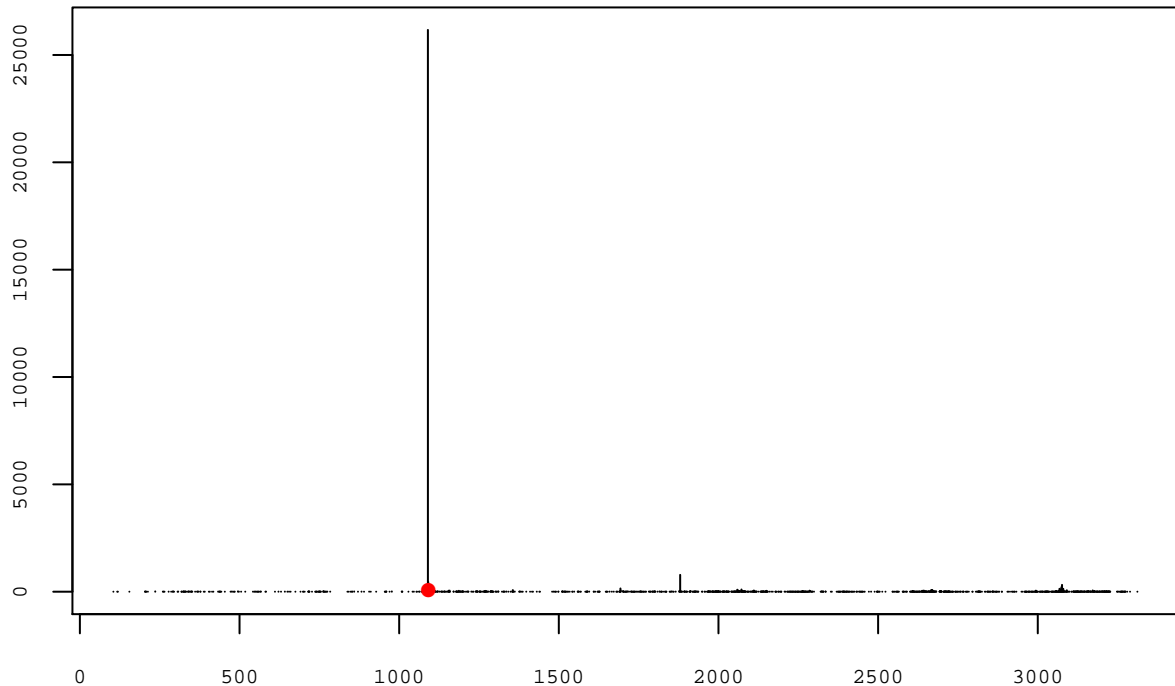


PggD7141_c1_g1_i1

5' TGGTTGGTATGAAGCCTGGTCCG-GATTCTATT '3
||| |||||
3' CCCTTACTTCGGACCAGGCTCT '5

Fragment Abundance



Transcript position

Cleavage site: 1092 Tag abundance: 57 Weighted abundance: 28.5 Category: 2
sRNA abundance: 23 Alignment score: 4 MFE ratio: 0.735 p-value: 0.038