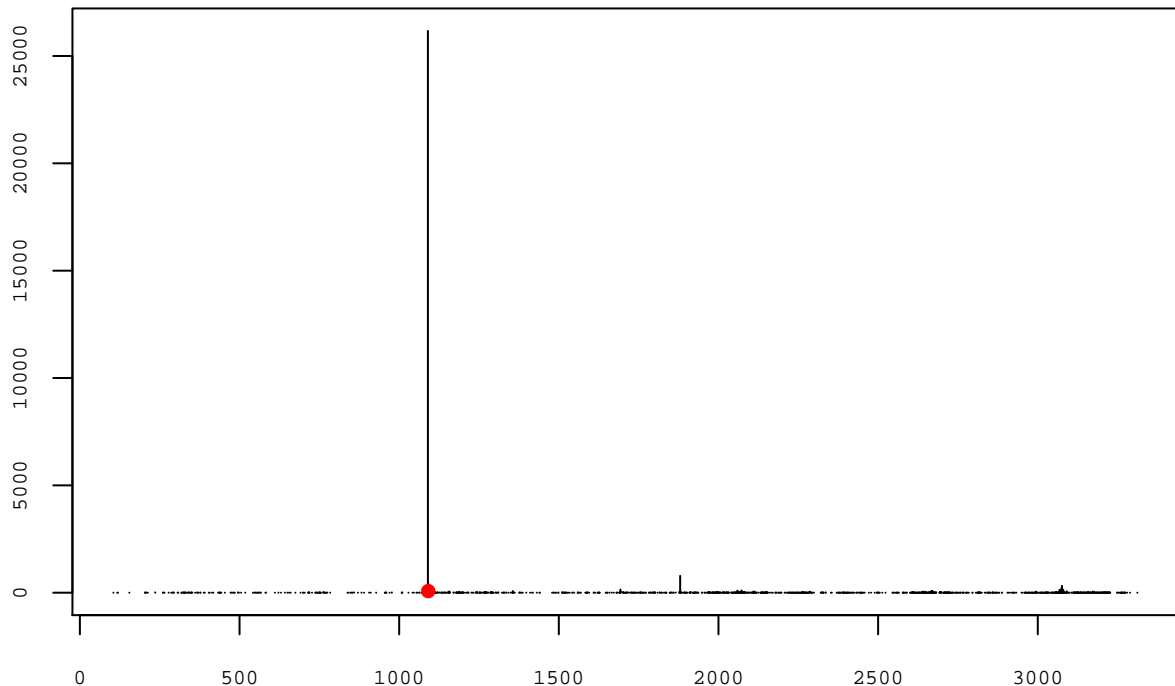


PggD7141\_c1\_g1\_i1

5' TGGTTGGTATGAAGCCTGGTCCG-GATTCTATT '3

3' CCTTACTTCGGACCAGGCTCT '5

Fragment Abundance



Transcript position

Cleavage site: 1092

Tag abundance: 57

Weighted abundance: 28.5

Category: 2

sRNA abundance: 143

Alignment score: 3

MFE ratio: 0.774

p-value: 0.038