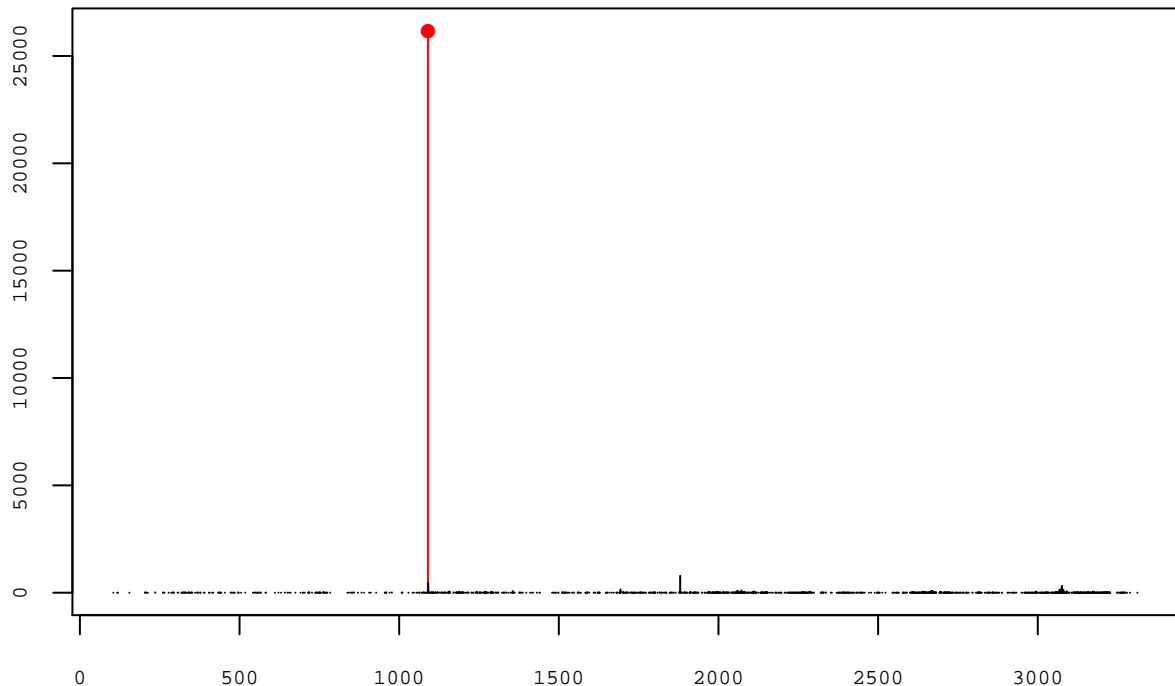


PggD7141_c1_g1_i1

5' GATGGTTGGTATGAAGCCTGGTCCGGATTCTA '3
o| || ||||| ||||| ||||| o
3' TCC-CCTTACTTCGGACCAGGCT '5

Fragment Abundance



Transcript position

Cleavage site: 1090 Tag abundance: 26168 Weighted abundance: 13084 Category: 0
sRNA abundance: 3011 Alignment score: 4 MFE ratio: 0.761 p-value: 0.001