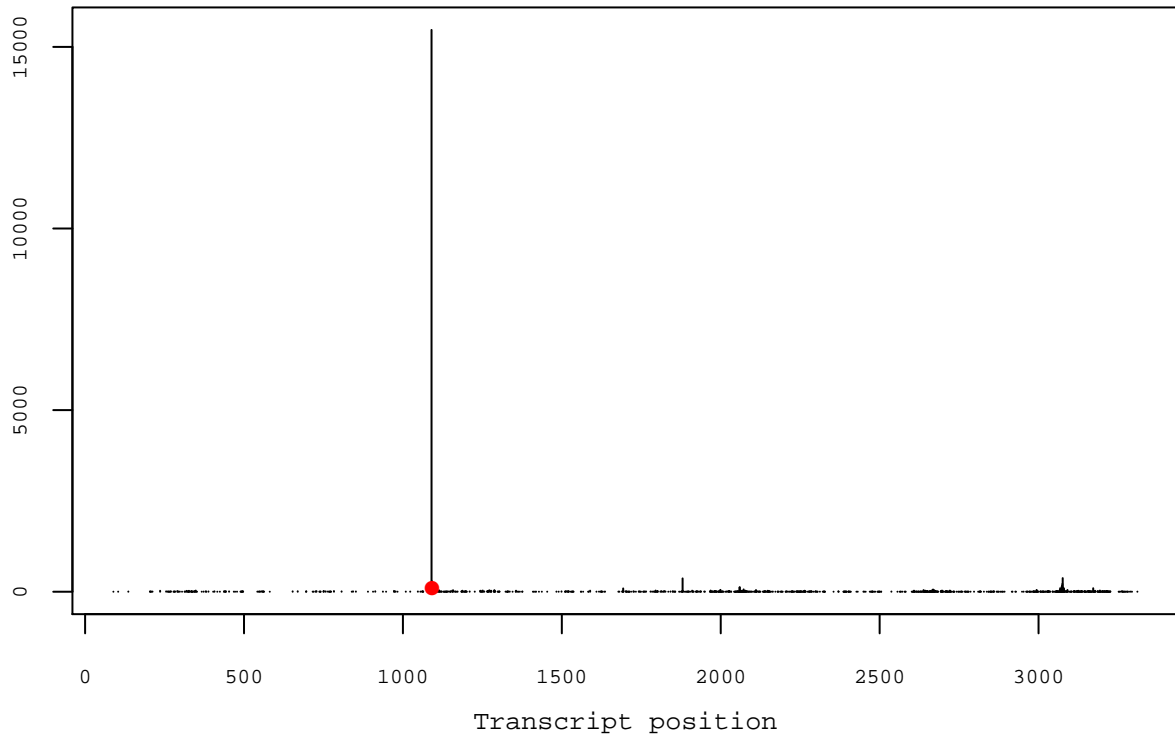


PggD7141_c1_g1_i1

5' TGGTTGGTATGAAGCCTGGTCCG-GATTCTATT '3
|| |||||
3' CCTTACTTCGGACCAGGCTCT '5

Fragment Abundance



Cleavage site: 1092 Tag abundance: 91 Weighted abundance: 45.5 Category: 2
sRNA abundance: 44 Alignment score: 3 MFE ratio: 0.774 p-value: 0.037