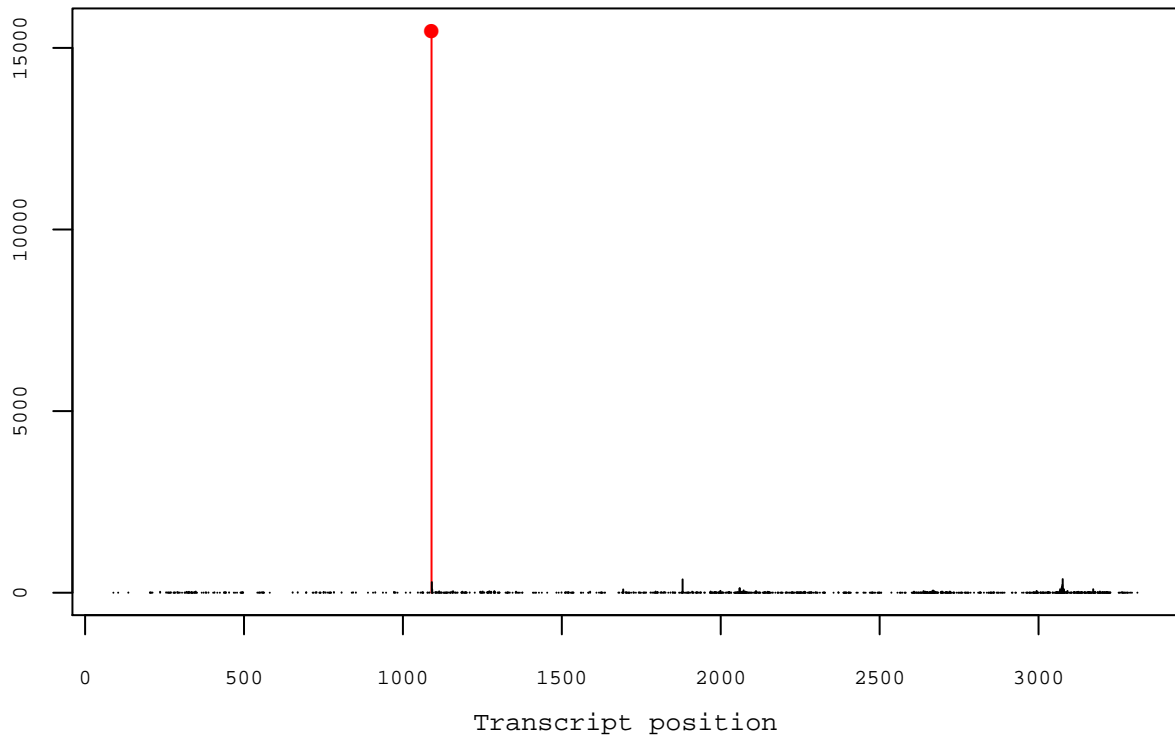


PggD7141_c1_g1_i1

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

Fragment Abundance



Cleavage site: 1090 Tag abundance: 15468 Weighted abundance: 7734 Category: 0
sRNA abundance: 696 Alignment score: 4 MFE ratio: 0.761 p-value: 0.001