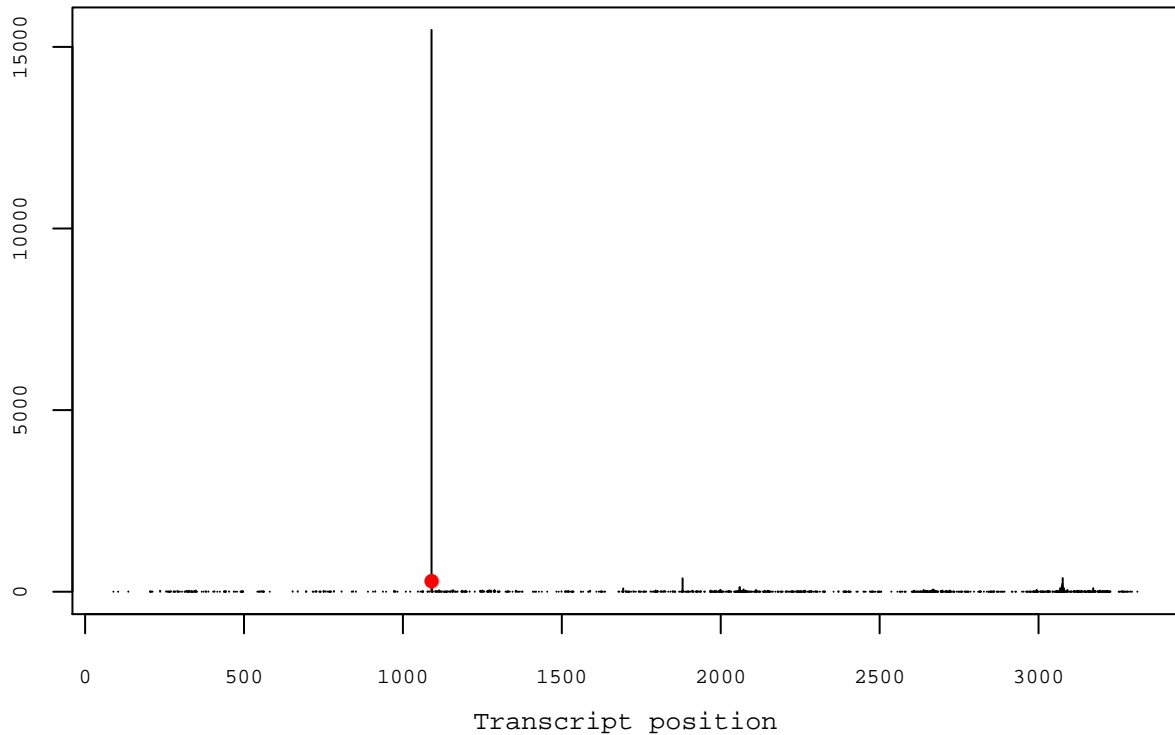


PggD7141_c1_g1_i1

5' ATGGTTGGTATGAAGCCTGGTCCG-GATTCTAT '3

3' CCCTTACTTCGGACCAGGCTC '5

Fragment Abundance



Cleavage site: 1091 Tag abundance: 291 Weighted abundance: 145.5 Category: 2
sRNA abundance: 146 Alignment score: 3 MFE ratio: 0.747 p-value: 0.037