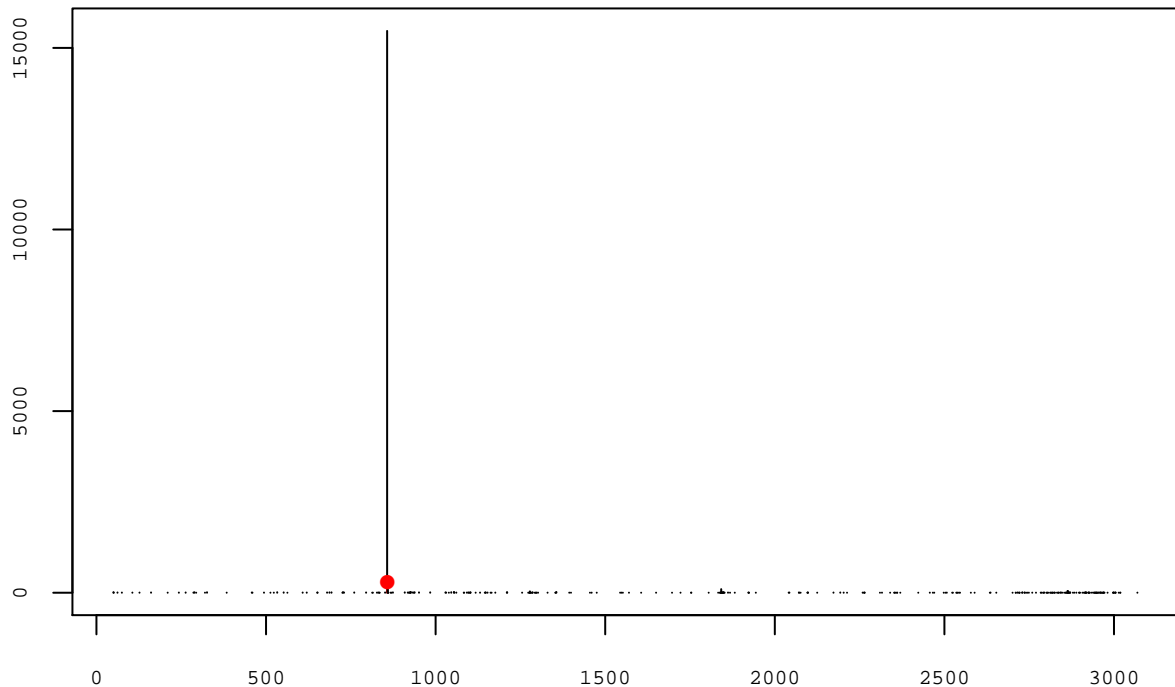


PggD11091_c20_g1_i1

5' ATGGTCGGGATGAAGCCTGGTCCG-GATTCTAT '3

||o|||||||||||||||
3' CCCCTTACTTCGGACCAGGCTC '5

Fragment Abundance



Transcript position

Cleavage site: 858 Tag abundance: 291 Weighted abundance: 145.5 Category: 2
sRNA abundance: 7 Alignment score: 3.5 MFE ratio: 0.786 p-value: 0.002