

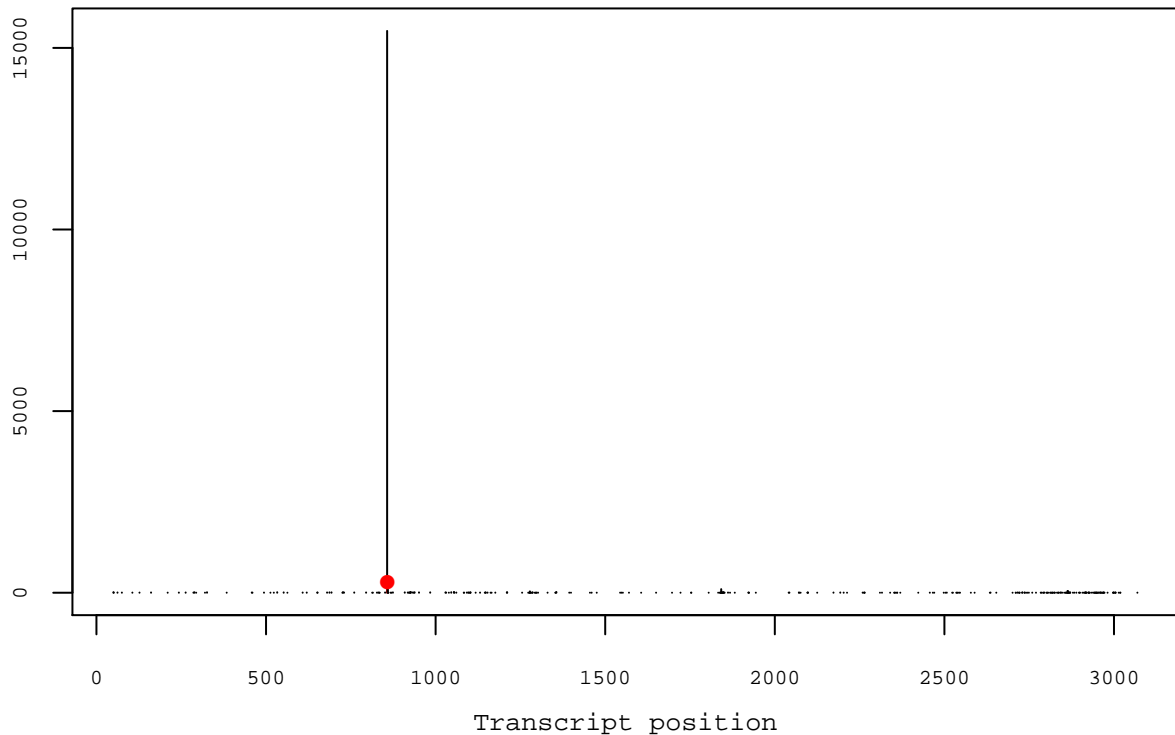
PggD11091_c20_g1_i1

5' ATGGTCGGGATGAAGCCTGGTCCG-GATTCTAT '3

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3' CCCTTACTTCGGACCAGGCTC '5

Fragment Abundance



Cleavage site: 858 Tag abundance: 291 Weighted abundance: 145.5 Category: 2
sRNA abundance: 146 Alignment score: 2.5 MFE ratio: 0.846 p-value: 0.001