

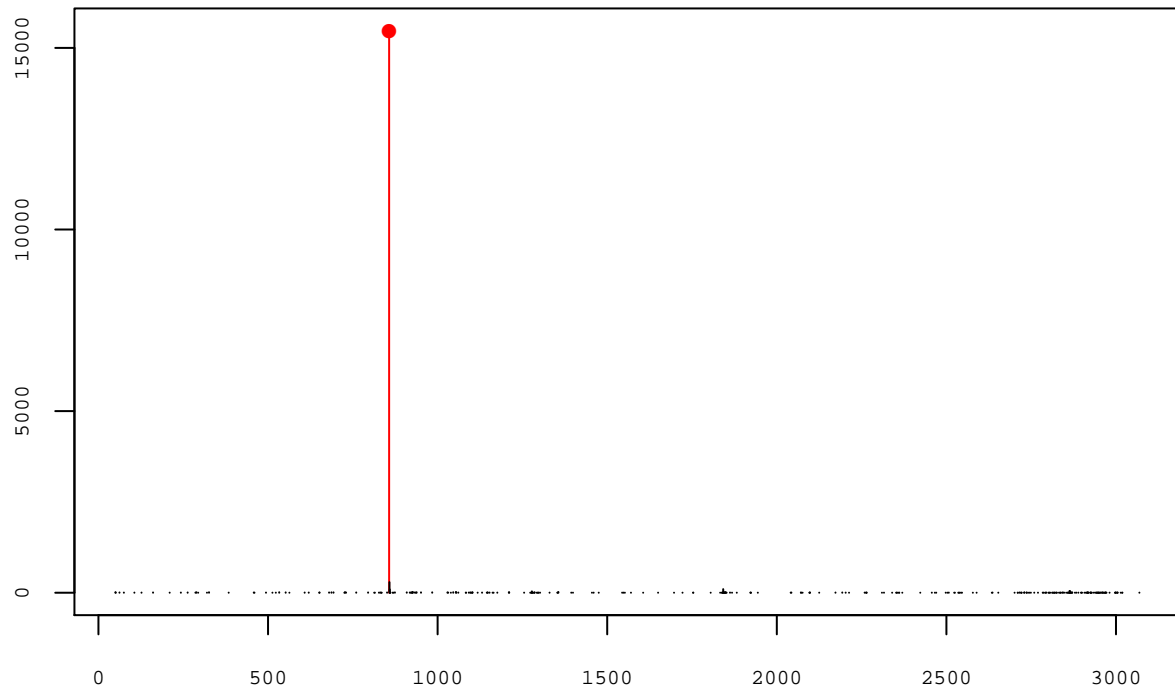
PggD11091_c20_g1_i1

5' GATGGTCGGGATGAAGCCTGGTCCGGATTCTA '3

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3' CCCTTACTTCGGACCAGGCT '5

Fragment Abundance



Transcript position

Cleavage site: 857 Tag abundance: 15468 Weighted abundance: 7734 Category: 0
sRNA abundance: 417 Alignment score: 2 MFE ratio: 0.924 p-value: 0.001