

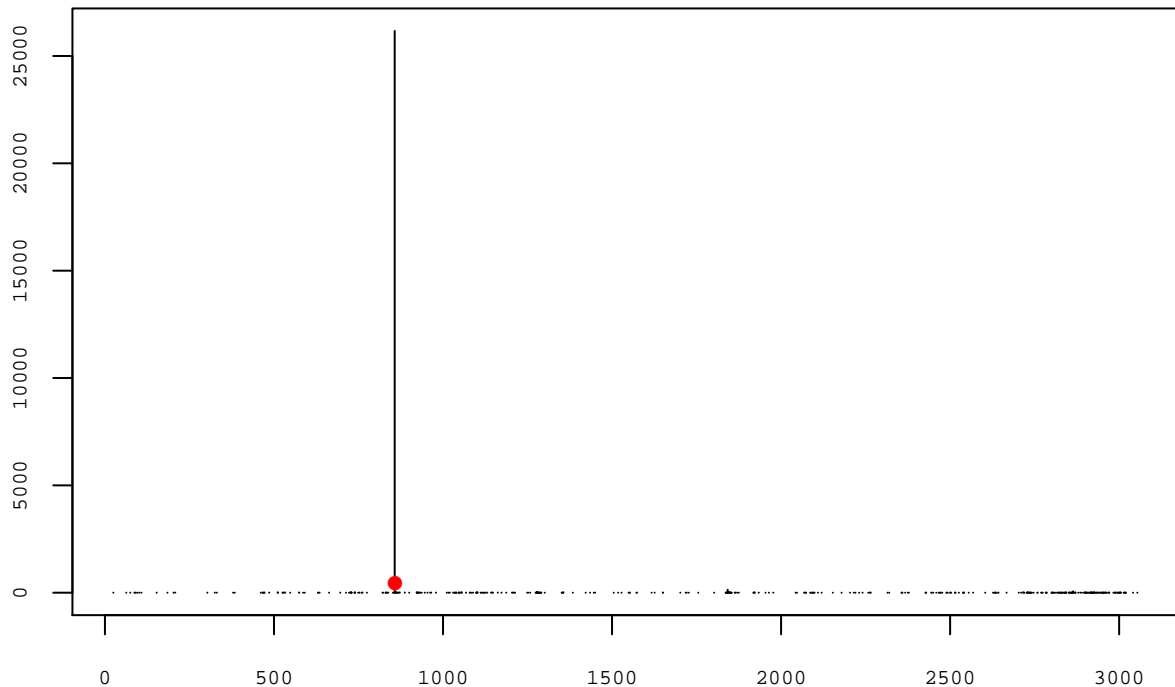
PggD11091\_c20\_g1\_i1

5' ATGGTCGGGATGAAGCCTGGTCCG-GATTCTAT '3

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3' CCCTTACTTCGGACCAGGCTC '5

Fragment Abundance



Transcript position

Cleavage site: 858 Tag abundance: 456 Weighted abundance: 228 Category: 2  
sRNA abundance: 371 Alignment score: 2.5 MFE ratio: 0.846 p-value: 0.001