

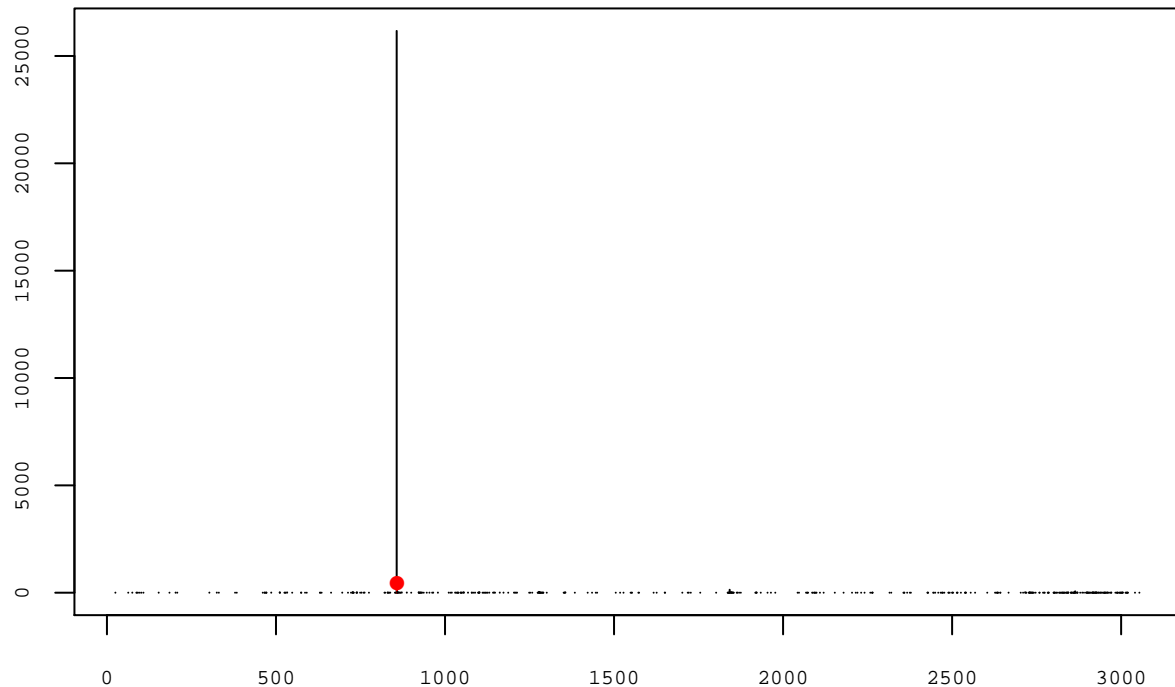
PggD11091_c20_g1_i1

5' ATGGTCGGGATGAAGCCTGGTCCGGATTCTAT '3

||o|||||||||||o|

3' CCCTTACTTCGGACCAGGCTT '5

Fragment Abundance



Transcript position

Cleavage site: 858 Tag abundance: 456 Weighted abundance: 228 Category: 2
sRNA abundance: 240 Alignment score: 2.5 MFE ratio: 0.921 p-value: 0.001