

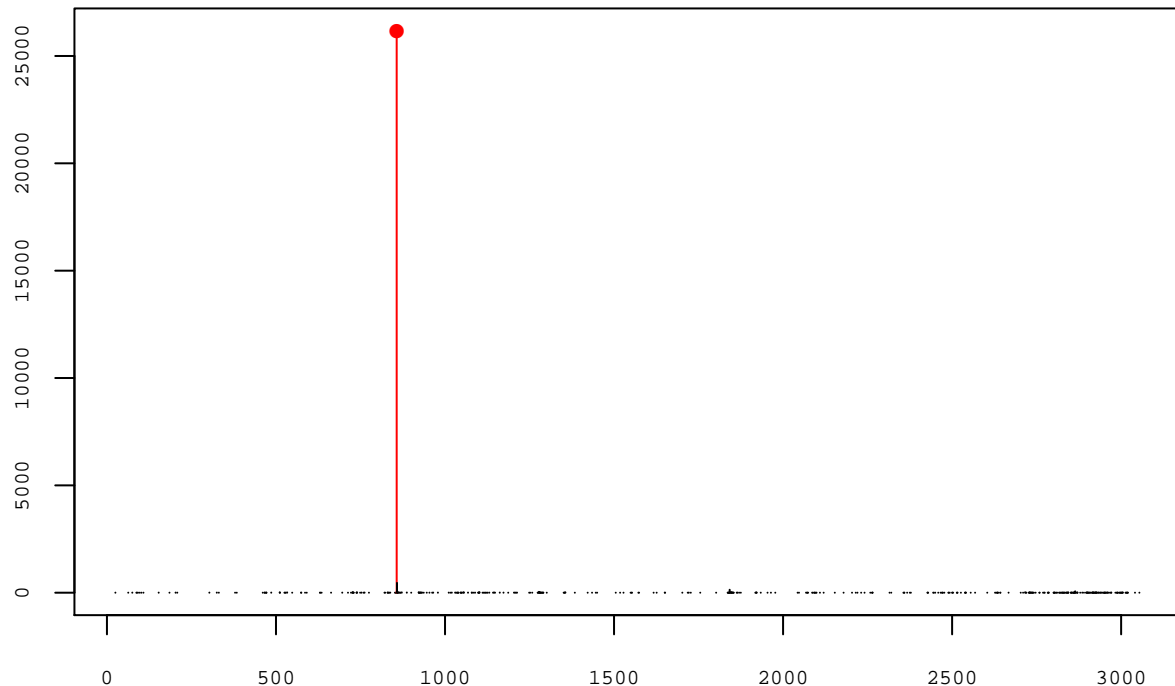
PggD11091_c20_g1_i1

5' GATGGTCGGGATGAAGCCTGGTCCGGATTCTA '3

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

3' TCCCTTACTTCGGACCAGGCT '5

Fragment Abundance



Transcript position

Cleavage site: 857 Tag abundance: 26168 Weighted abundance: 13084 Category: 0
sRNA abundance: 3011 Alignment score: 3.5 MFE ratio: 0.825 p-value: 0.001