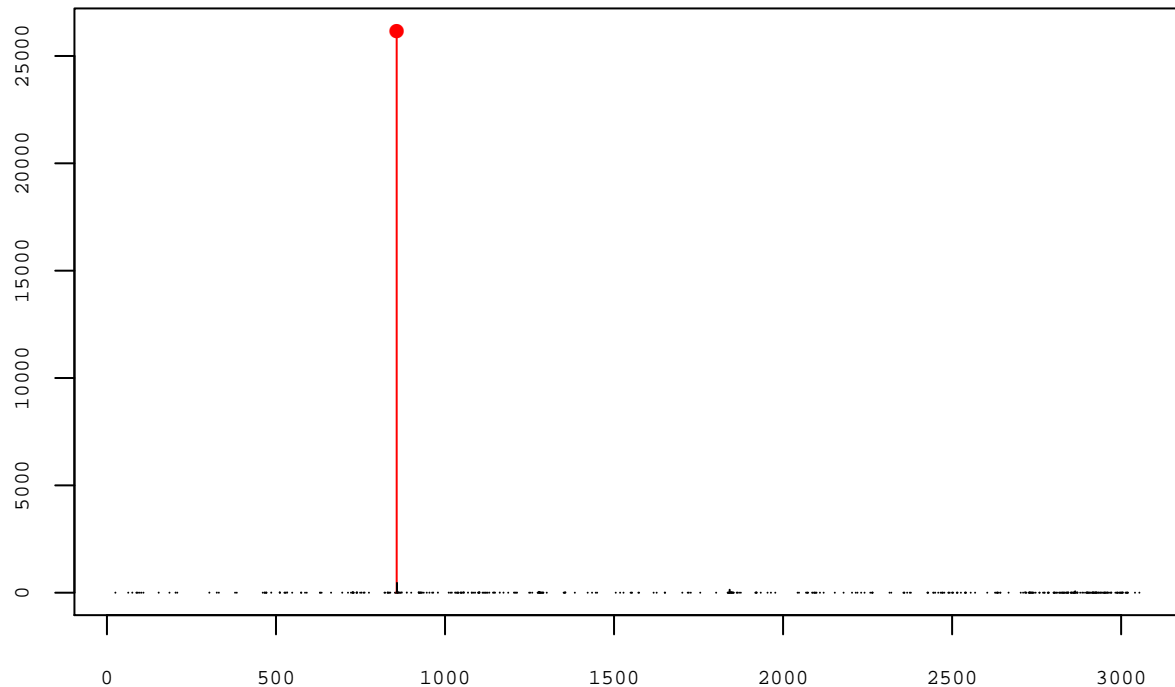


PggD11091_c20_g1_i1

5' GATGGTCGGGATGAAGCCTGGTCCGGATTCTA '3
||o|||||||||||o
3' CCCCTTACTTCGGACCAGGCT '5

Fragment Abundance



Transcript position

Cleavage site: 857 Tag abundance: 26168 Weighted abundance: 13084 Category: 0
sRNA abundance: 44559 Alignment score: 3 MFE ratio: 0.855 p-value: 0.001