

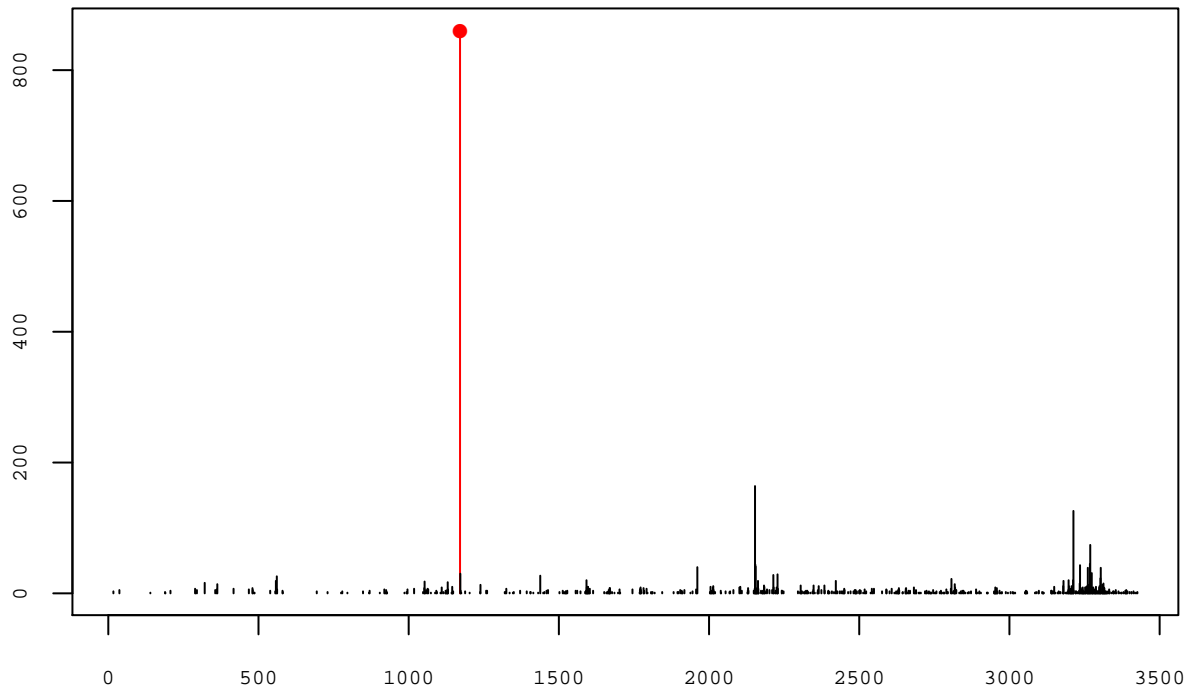
PggD9924_c6_g1_i1

5' GCTTGTGGAATGAAGCCTGGTCCGGATTCCA '3

|||||

3' CCCCTTACTTCGGACCAGGCT '5

Fragment Abundance



Transcript position

Cleavage site: 1171 Tag abundance: 860 Weighted abundance: 860 Category: 0
sRNA abundance: 44559 Alignment score: 2.5 MFE ratio: 0.859 p-value: 0