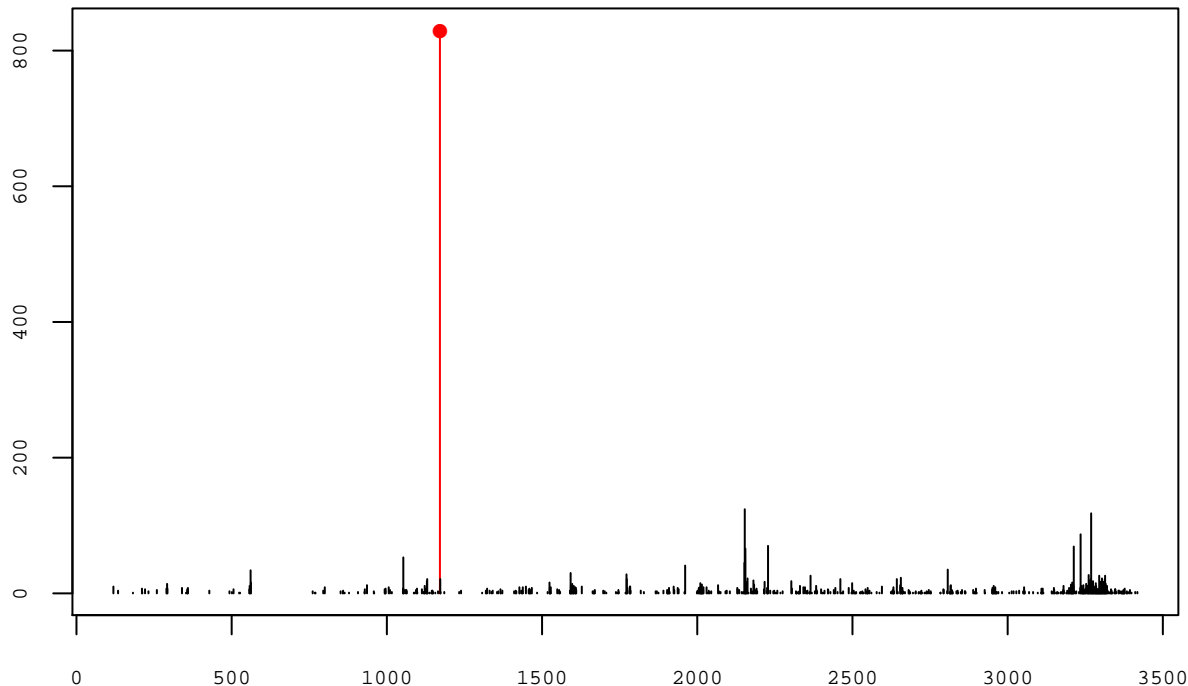


PggD9924_c6_g1_i1

5' GCTTGTGGGAATGAAGCCTGGTCCGGATTCCA '3
|||||
3' CCCCTTACTTCGGACCAGGCT '5

Fragment Abundance



Transcript position

Cleavage site: 1171 Tag abundance: 829 Weighted abundance: 829 Category: 0
sRNA abundance: 19192 Alignment score: 2.5 MFE ratio: 0.859 p-value: 0