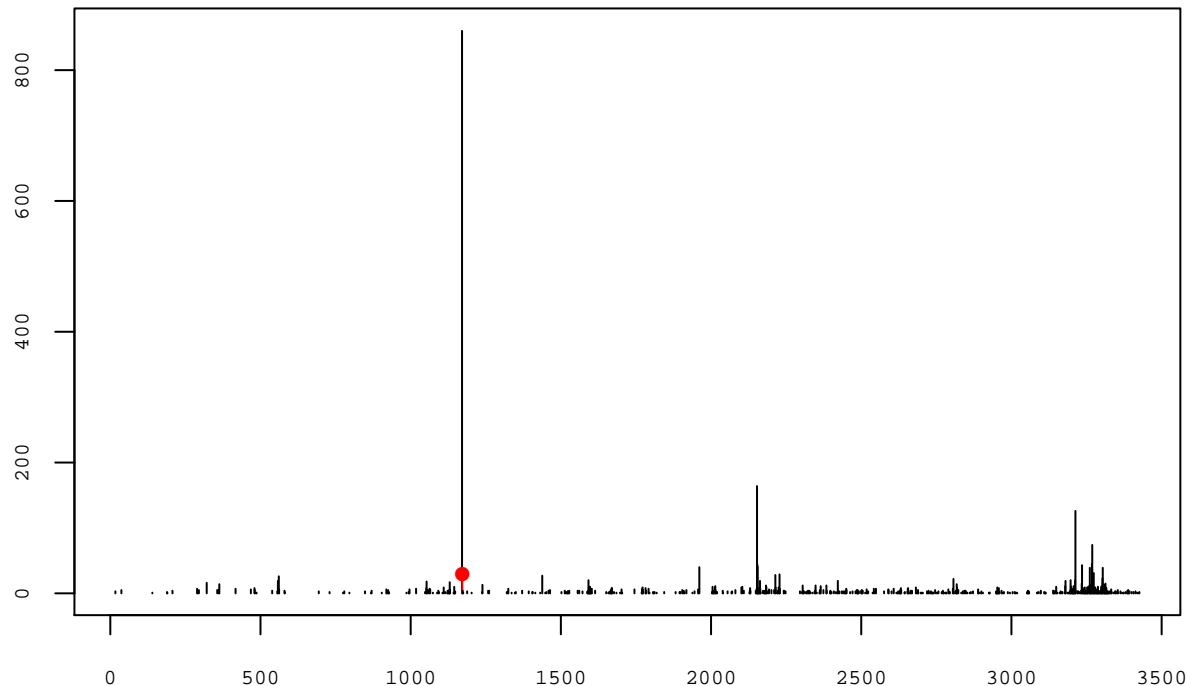


PggD9924_c6_g1_i1

5' CTTGTTGGAATGAAGCCTGGTCCG-GATTCCAT '3

|||||
3' CCCCTTACTTCGGACCAGGCTC '5

Fragment Abundance



Transcript position

Cleavage site: 1172 Tag abundance: 30 Weighted abundance: 30 Category: 2
sRNA abundance: 15 Alignment score: 3 MFE ratio: 0.79 p-value: 0.021