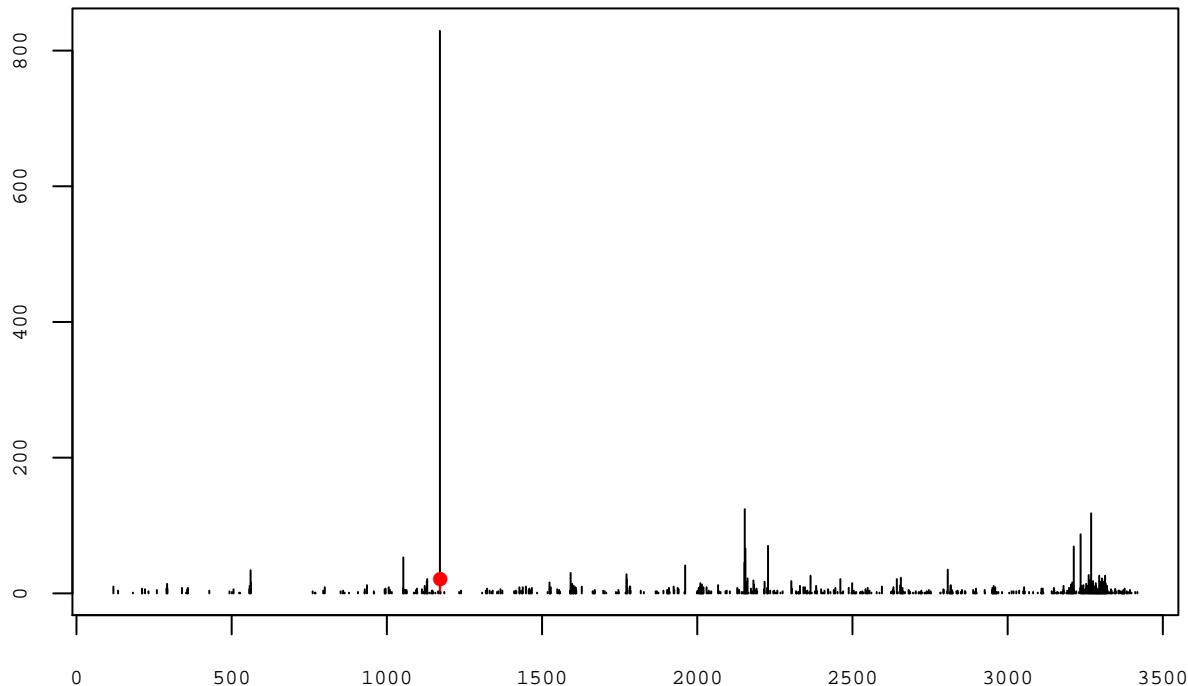


PggD9924\_c6\_g1\_i1

5' CTTGTTGGAATGAAGCCTGGTCCG-GATTCCAT '3

|||||  
3' CCCCTTACTTCGGACCAGGCTC '5

Fragment Abundance



Transcript position

Cleavage site: 1172 Tag abundance: 21 Weighted abundance: 21 Category: 2  
sRNA abundance: 7 Alignment score: 3 MFE ratio: 0.79 p-value: 0.025