

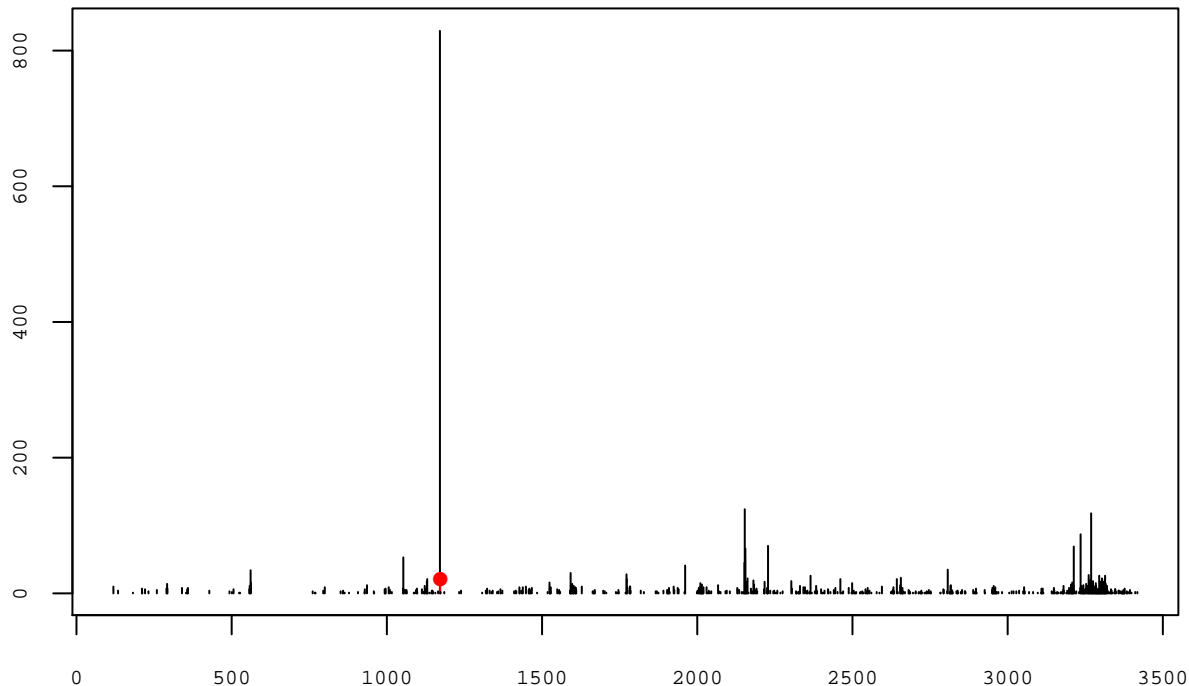
PggD9924\_c6\_g1\_i1

5' CTTGTTGGAATGAAGCCTGGTCCGGATTCCAT '3

|||||

3' CCCTTACTTCGGACCAGGCTT '5

Fragment Abundance



Transcript position

Cleavage site: 1172 Tag abundance: 21 Weighted abundance: 21 Category: 2  
sRNA abundance: 97 Alignment score: 2 MFE ratio: 0.926 p-value: 0.025