

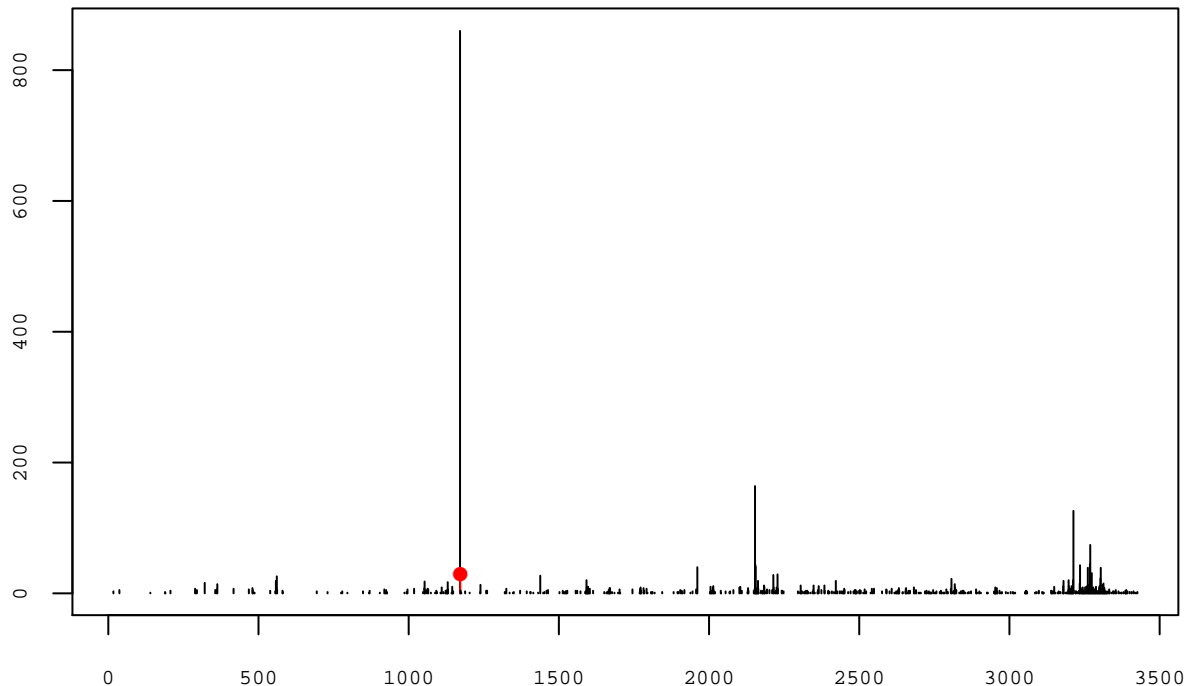
PggD9924\_c6\_g1\_i1

5' CTTGTTGGAATGAAGCCTGGTCCGGATTCCAT '3

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|||||

3' CCCTTACTTCGGACCAGGCTT '5

Fragment Abundance



Transcript position

Cleavage site: 1172 Tag abundance: 30 Weighted abundance: 30 Category: 2  
sRNA abundance: 240 Alignment score: 2 MFE ratio: 0.926 p-value: 0.021