

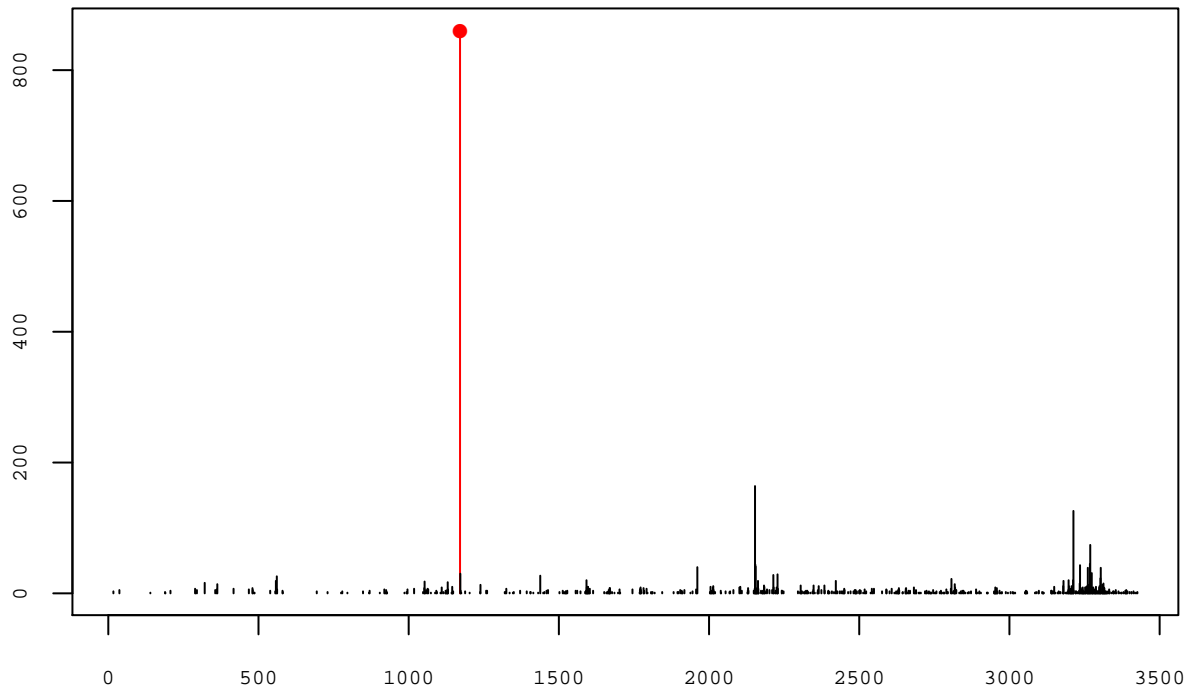
PggD9924\_c6\_g1\_i1

5' GCTTGTGGAATGAAGCCTGGTCCGGATTCCA '3

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3' TCCCCTTACTTCGGACCAGGCT '5

Fragment Abundance



Transcript position

Cleavage site: 1171 Tag abundance: 860 Weighted abundance: 860 Category: 0  
sRNA abundance: 3011 Alignment score: 3 MFE ratio: 0.829 p-value: 0