

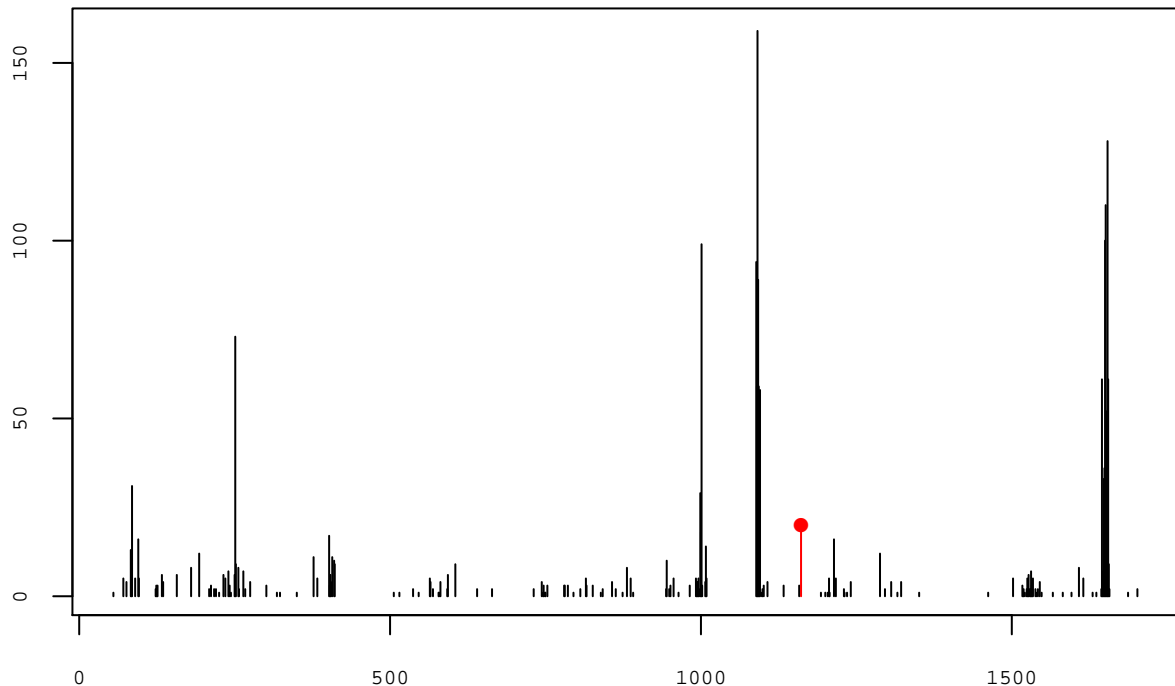
PggD11514\_c11\_g1\_i5

5' TCACCTGATTGACGGTCAGGATCGGACATCAC '3

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3' ACTAGATGCCAGTCCTAACCT '5

Fragment Abundance



Transcript position

Cleavage site: 1161 Tag abundance: 20 Weighted abundance: 20 Category: 2  
sRNA abundance: 6 Alignment score: 3.5 MFE ratio: 0.725 p-value: 0.028