

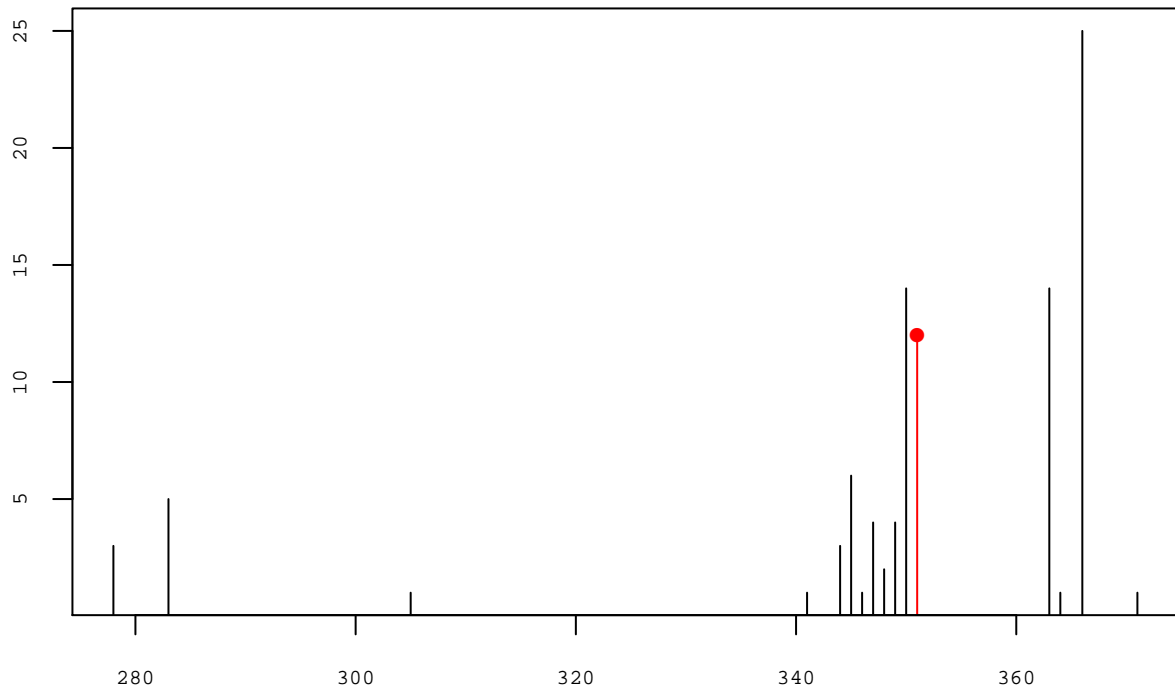
PggD14795\_c465\_g1\_i1

5' AACAAAGAGTTGTTTCACGAACAAAGAGTTGTTT '3

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3' GTGCTCGACAAGTGCTTGTGTTTATC '5

Fragment Abundance



Transcript position

Cleavage site: 351 Tag abundance: 12 Weighted abundance: 2.4 Category: 2  
sRNA abundance: 15 Alignment score: 3.5 MFE ratio: 0.708 p-value: 0.038