

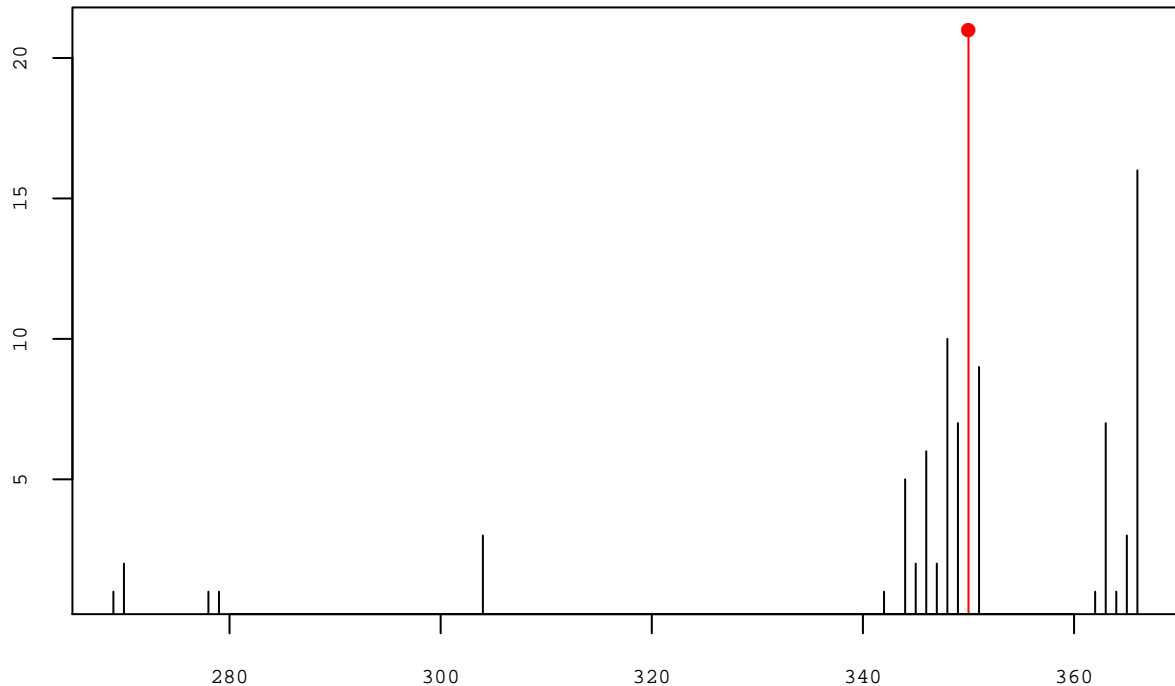
PggD14795_c465_g1_i1

5' GAACAAGAGTTGTTACGAACAAA-GAGTTGTT '3

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3' TGTGCTCGACAAGTGCTTGTATTAT '5

Fragment Abundance



Cleavage site: 350 Tag abundance: 21 Weighted abundance: 5.25 Category: 0
sRNA abundance: 23 Alignment score: 3 MFE ratio: 0.749 p-value: 0.028