

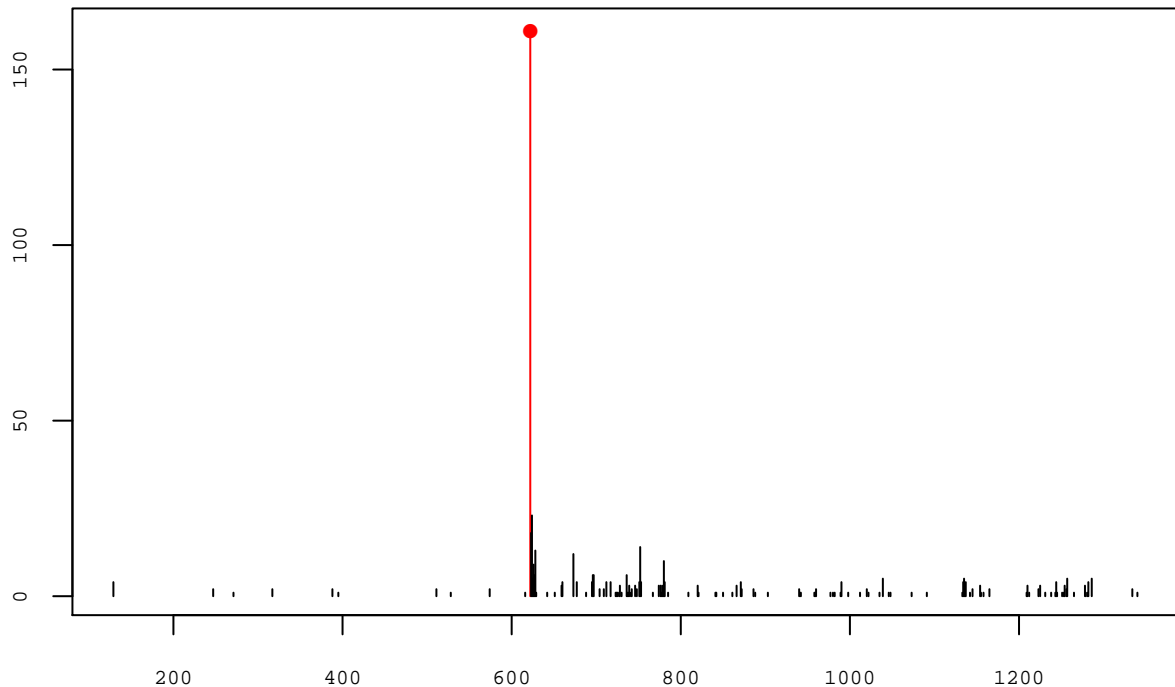
PggD13991_c0_g1_i1

5' GAGCTGATACATTTGTGATATCTTAATTTGGA '3

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3' ATGTAAATGCTATAGAATT '5

Fragment Abundance



Transcript position

Cleavage site: 622 Tag abundance: 161 Weighted abundance: 79.833 Category: 0
sRNA abundance: 9 Alignment score: 2 MFE ratio: 0.79 p-value: 0.001