

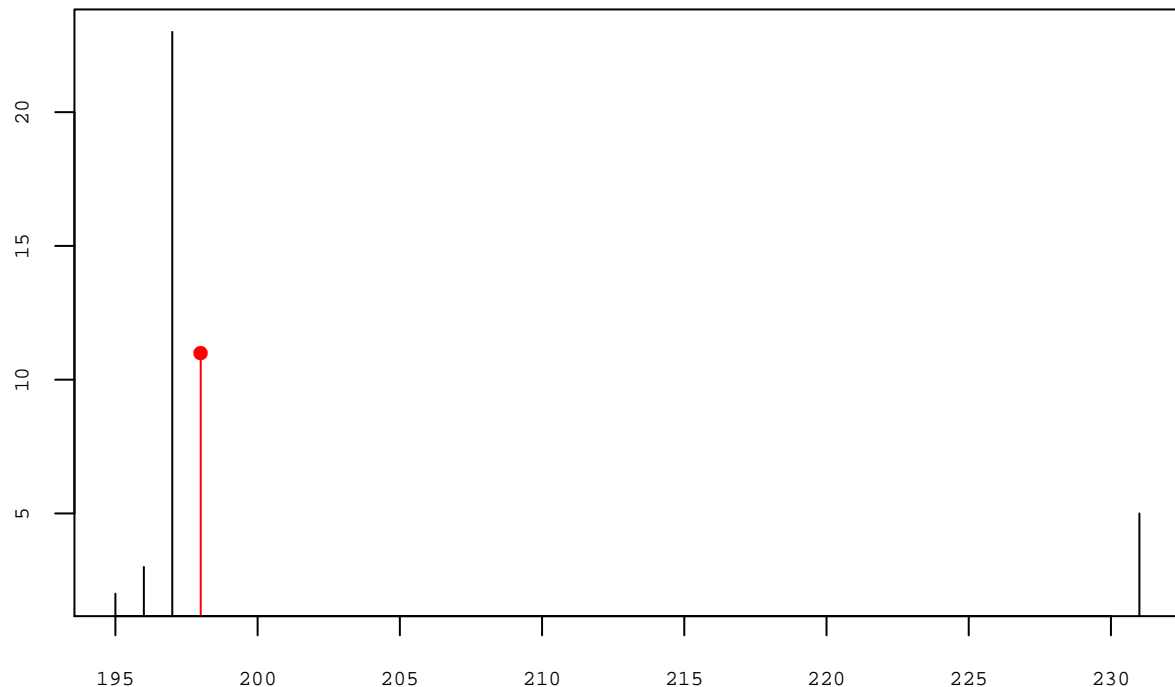
PggD7237_c562_g1_i1

5' TGTACCCTCCAAATAACCTAATAT-ATATGCAA '3

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3' TGGGAGGCTTATTGGATTATACA '5

Fragment Abundance



Transcript position

Cleavage site: 198 Tag abundance: 11 Weighted abundance: 5.5 Category: 2
sRNA abundance: 9 Alignment score: 3 MFE ratio: 0.745 p-value: 0.005