

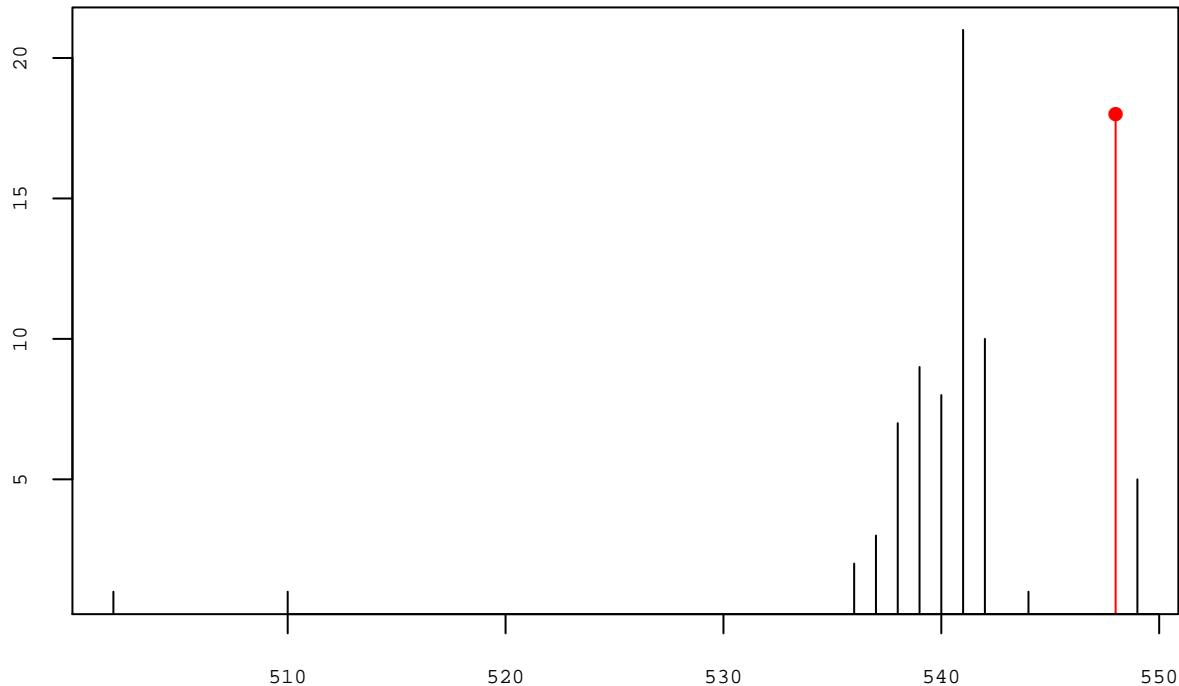
PggD794_c44_g1_i1

5' AATAGCTCGTGTTCGACTCGTTAAGAAGCTCG '3

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3' GTCGAGCACAAAGCTGAGCAATTCG '5

Fragment Abundance



Transcript position

Cleavage site: 548 Tag abundance: 18 Weighted abundance: 0.947 Category: 2
sRNA abundance: 20 Alignment score: 1.5 MFE ratio: 0.918 p-value: 0.007