

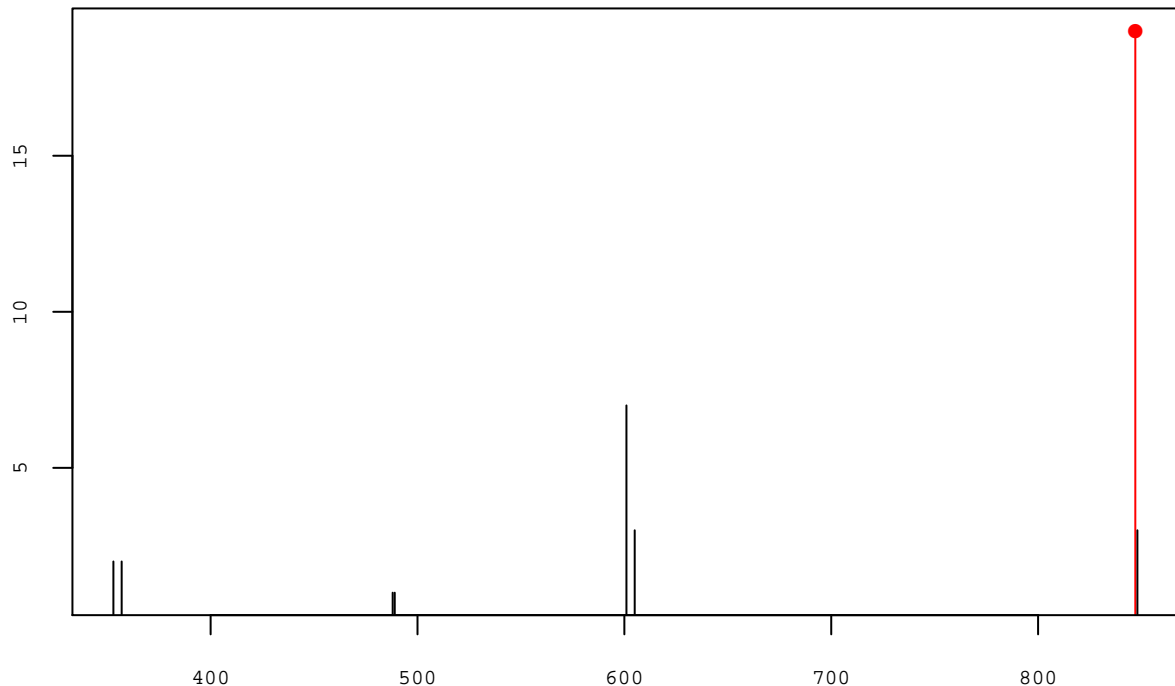
PggD5980\_c3\_g1\_i1

5' TATTTTAGTCCTAGAGTCTATAA-AAGTGCCAA '3

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3' GTCAGGGTTTCAGATATTGTT '5

Fragment Abundance



Cleavage site: 847 Tag abundance: 19 Weighted abundance: 1.9 Category: 0  
sRNA abundance: 16 Alignment score: 4 MFE ratio: 0.746 p-value: 0.005