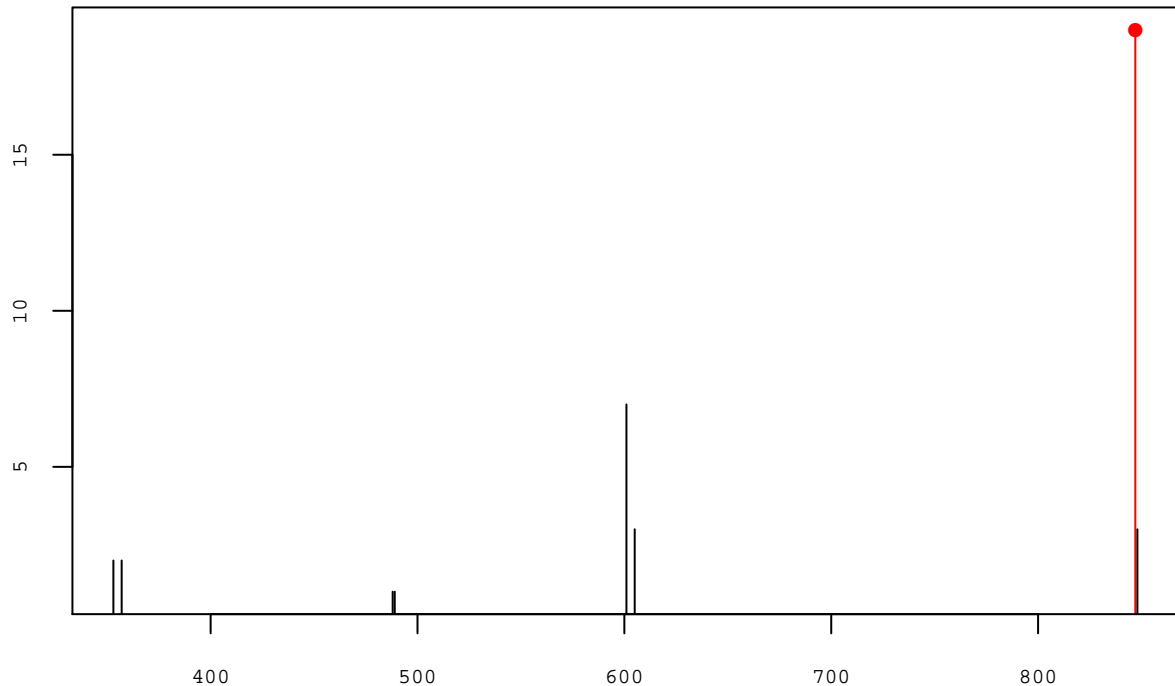


PggD5980_c3_g1_i1

5' TATTTTAGTCCTAGAGTCTATAA-AAGTGCCAA '3

|||||o|o||||||| ||
3' TCAGGGTTTCAGATATTGTT '5

Fragment Abundance



Transcript position

Cleavage site: 847 Tag abundance: 19 Weighted abundance: 1.9 Category: 0
sRNA abundance: 6 Alignment score: 3.5 MFE ratio: 0.776 p-value: 0.005