

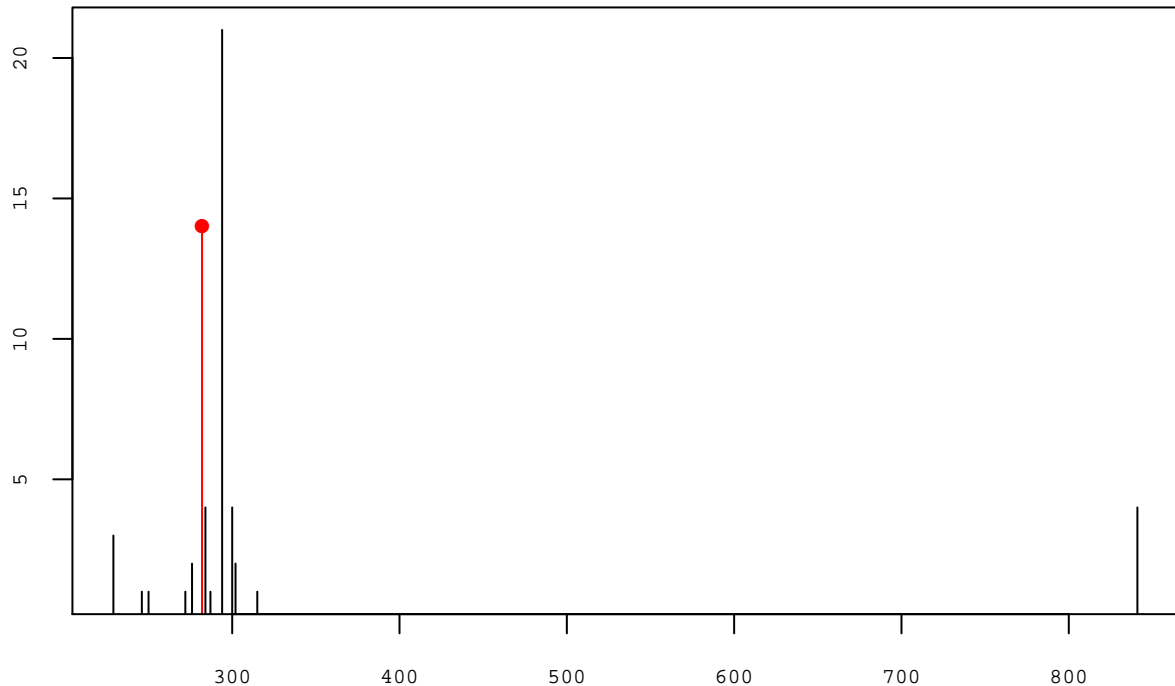
PggD6653_c100_g1_i1

5' GACGACAGTTCAGGAAAGCTGTGGGAAGATAT '3

|||||o|||||||o|

3' CAAGTTCTTTCGACACCTT '5

Fragment Abundance



Cleavage site: 282 Tag abundance: 14 Weighted abundance: 4.667 Category: 2
sRNA abundance: 80 Alignment score: 1.5 MFE ratio: 0.971 p-value: 0.003