

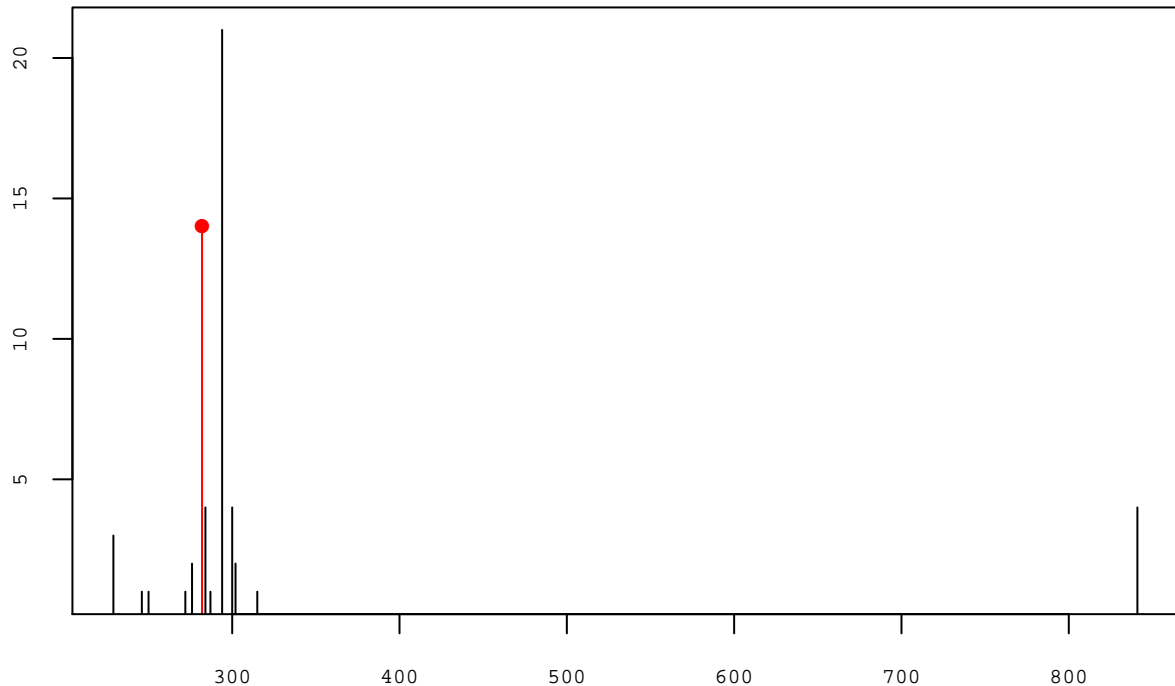
PggD6653\_c100\_g1\_i1

5' GACGACAGTTCAGGAAAGCTGTGGGAAGATAT '3

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3' GTCAAGTTCCTTCGACACCTT '5

Fragment Abundance



Cleavage site: 282  
sRNA abundance: 470

Tag abundance: 14  
Alignment score: 1.5

Weighted abundance: 4.667  
MFE ratio: 0.974

Category: 2  
p-value: 0.003