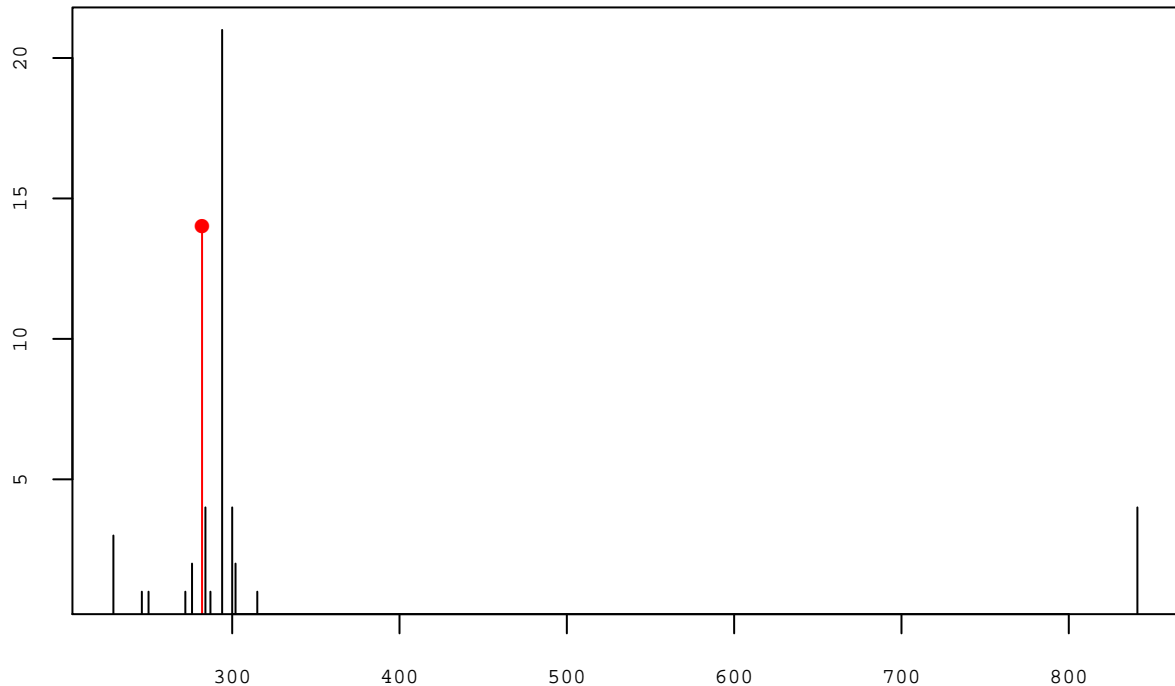


PggD6653_c100_g1_i1

5' GACGACAGTTCAGGAAAGCTGTGGGAAGATAT '3
|||||o|||||||o|
3' TCAAGTTCTTTCGACACCTT '5

Fragment Abundance



Transcript position

Cleavage site: 282
sRNA abundance: 196

Tag abundance: 14
Alignment score: 1.5

Weighted abundance: 4.667
MFE ratio: 0.972

Category: 2
p-value: 0.003