

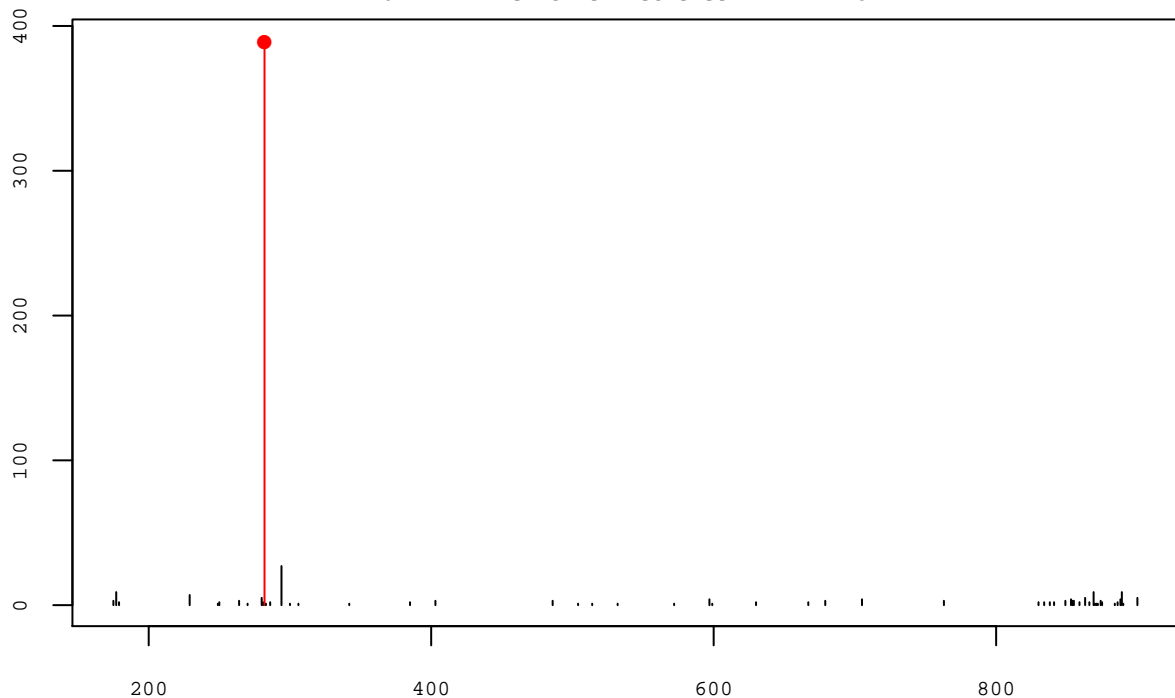
PggD6653_c100_g1_i1

5' GACGACAGTTCAGGAAAGCTGTGGGAAGATAT '3

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3' TCAAGTTCTTTCGACACCTT '5

Fragment Abundance



Transcript position

Cleavage site: 282 Tag abundance: 389 Weighted abundance: 129.667 Category: 0
sRNA abundance: 117 Alignment score: 1.5 MFE ratio: 0.972 p-value: 0.003