

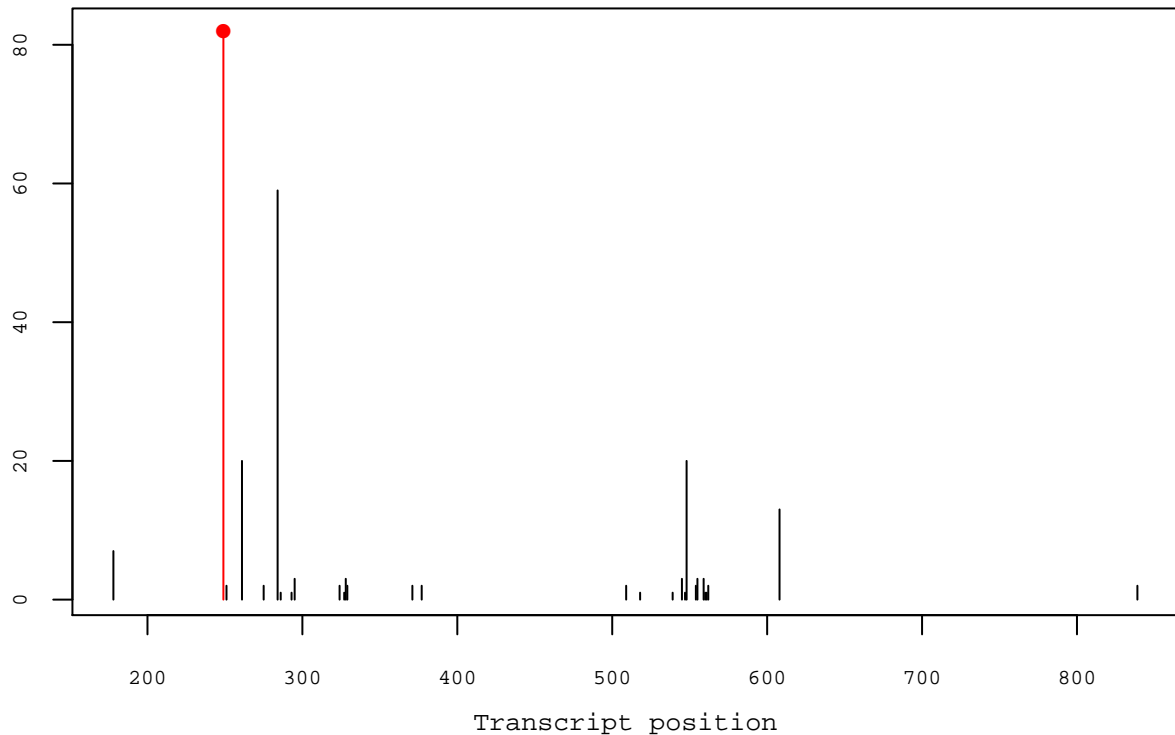
PggD11120_c28_g1_i1

5' GGTTCAGTTCAATAAAGCTGTGGGAAATTAC '3

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3' CAAGTTCTTTCGACACCTT '5

Fragment Abundance



Cleavage site: 249 Tag abundance: 82 Weighted abundance: 41 Category: 0
sRNA abundance: 56 Alignment score: 3 MFE ratio: 0.773 p-value: 0.017