

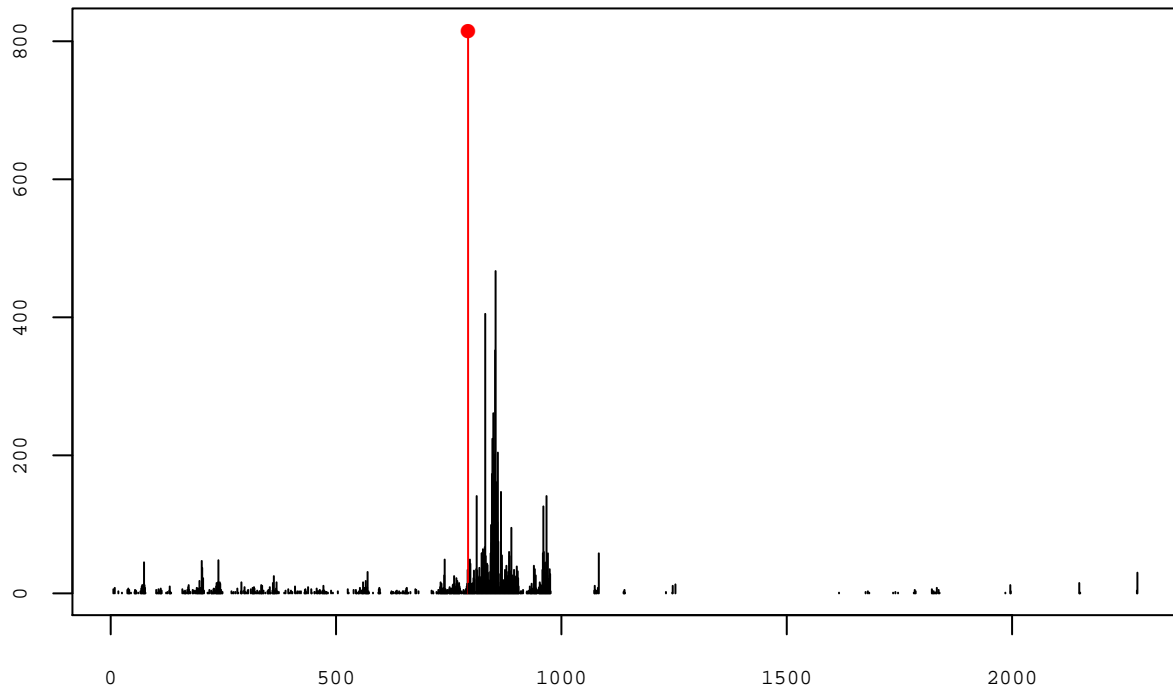
PggD6617_c1_g2_i1

5' TCAAGCATGCTTGCTCTCTTCTGTCCTAGAT 3'

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3' CACGAGTGAGAGAAGACAGT 5'

Fragment Abundance



Transcript position

Cleavage site: 793 Tag abundance: 815 Weighted abundance: 815 Category: 0
sRNA abundance: 44 Alignment score: 2 MFE ratio: 0.858 p-value: 0.008