

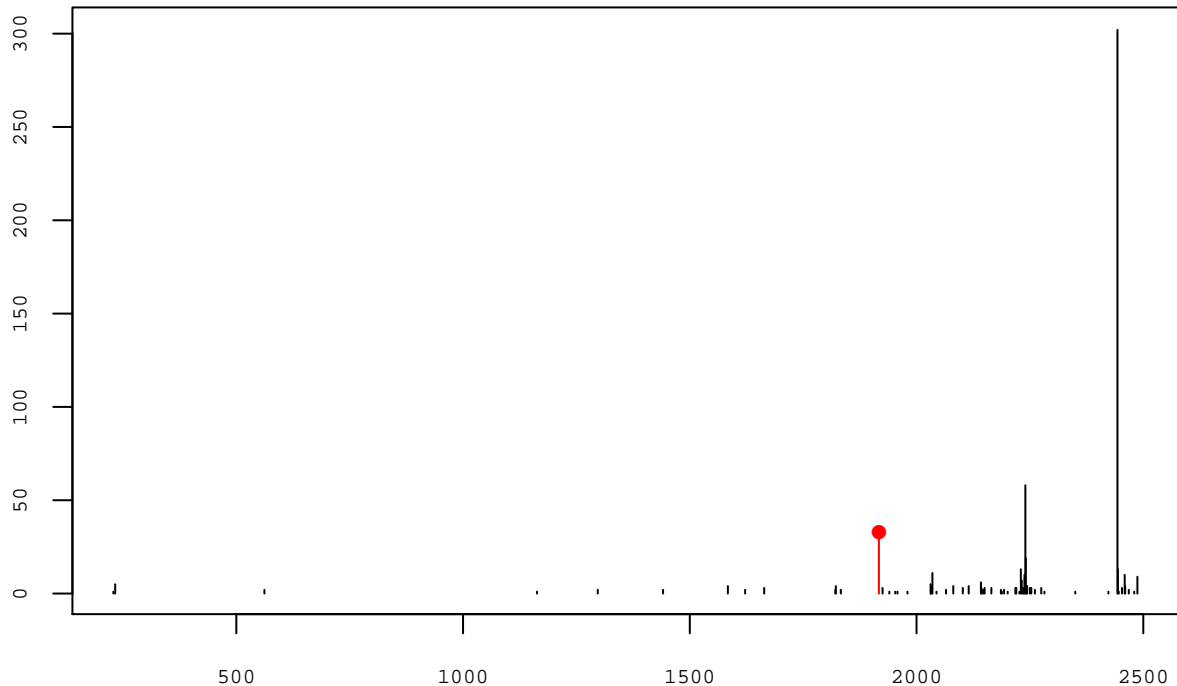
PggD12654_c9_g1_i1

5' CTCTTCTGCAGCATCATCAGGATTCTCAGCCT 3'

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3' TCCGTCGTAGTAGTTCTAAGA 5'

Fragment Abundance



Transcript position

Cleavage site: 1917 Tag abundance: 33 Weighted abundance: 11 Category: 2
sRNA abundance: 139 Alignment score: 3 MFE ratio: 0.879 p-value: 0.027