

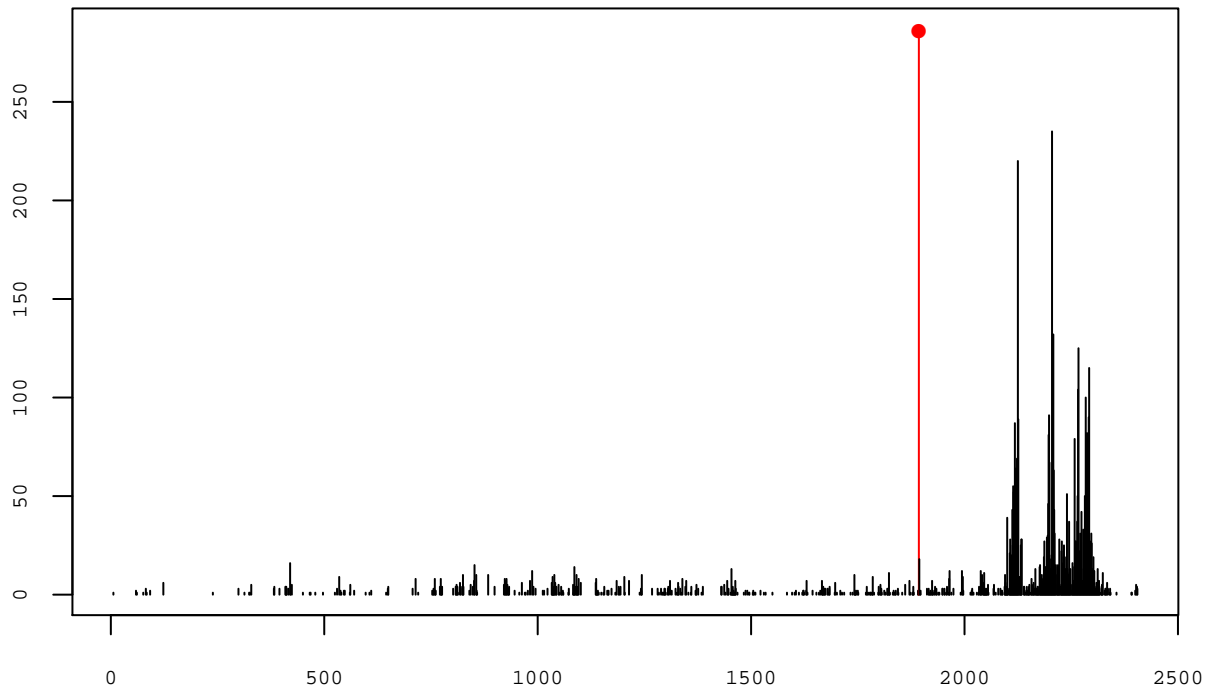
PggD9652_c11_g1_i1

5' TTTAATAGGGGGA-CCCTTCAGTCCAATTCACC '3

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3' CCTCGAGGGAAGTCAGGTT '5

Fragment Abundance



Transcript position

Cleavage site: 1893 Tag abundance: 286 Weighted abundance: 71.5 Category: 0
sRNA abundance: 282 Alignment score: 2.5 MFE ratio: 0.768 p-value: 0.005