

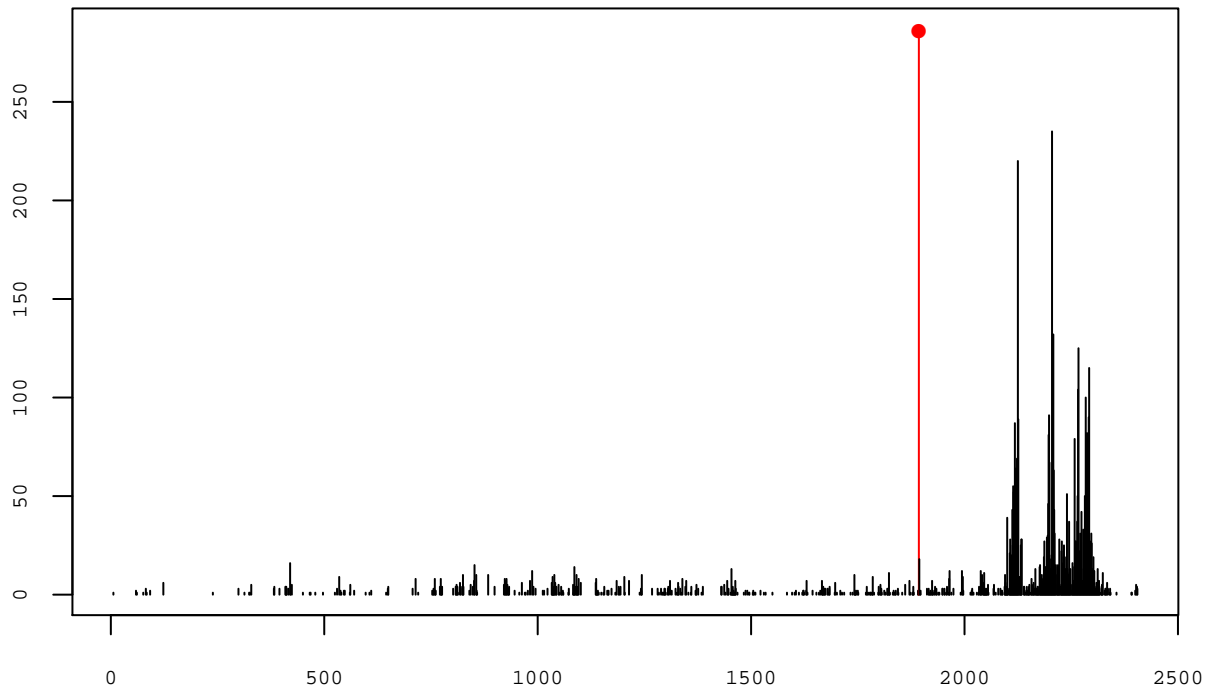
PggD9652_c11_g1_i1

5' TTTAATAGGGGGA-CCCTTCAGTCCAATTCACC '3

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3' TCCCTCGAGGGAAGTCAGGTT '5

Fragment Abundance



Transcript position

Cleavage site: 1893

Tag abundance: 286

Weighted abundance: 71.5

Category: 0

sRNA abundance: 13228

Alignment score: 2.5

MFE ratio: 0.794

p-value: 0.003