

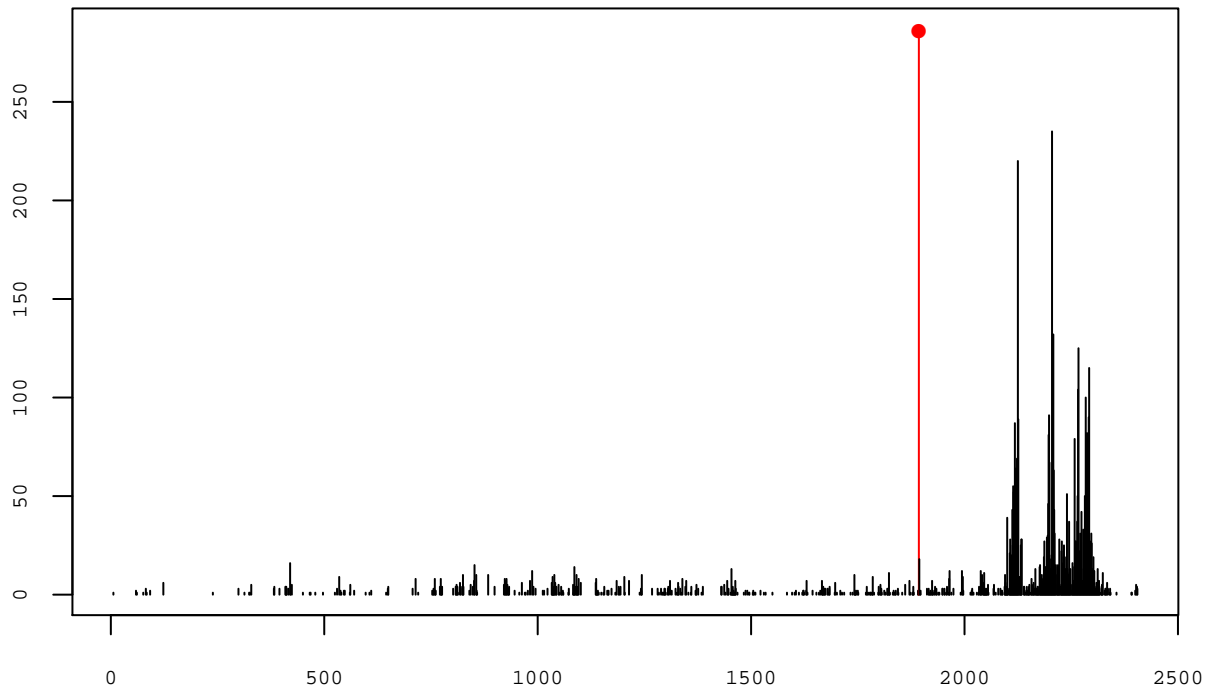
PggD9652_c11_g1_i1

5' TTTAATAGGGGGA-CCCTTCAGTCCAATTCACC '3

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3' CCCTCGAGGGAAGTCAGGTT '5

Fragment Abundance



Transcript position

Cleavage site: 1893	Tag abundance: 286	Weighted abundance: 71.5	Category: 0
sRNA abundance: 1709	Alignment score: 2.5	MFE ratio: 0.786	p-value: 0.003