

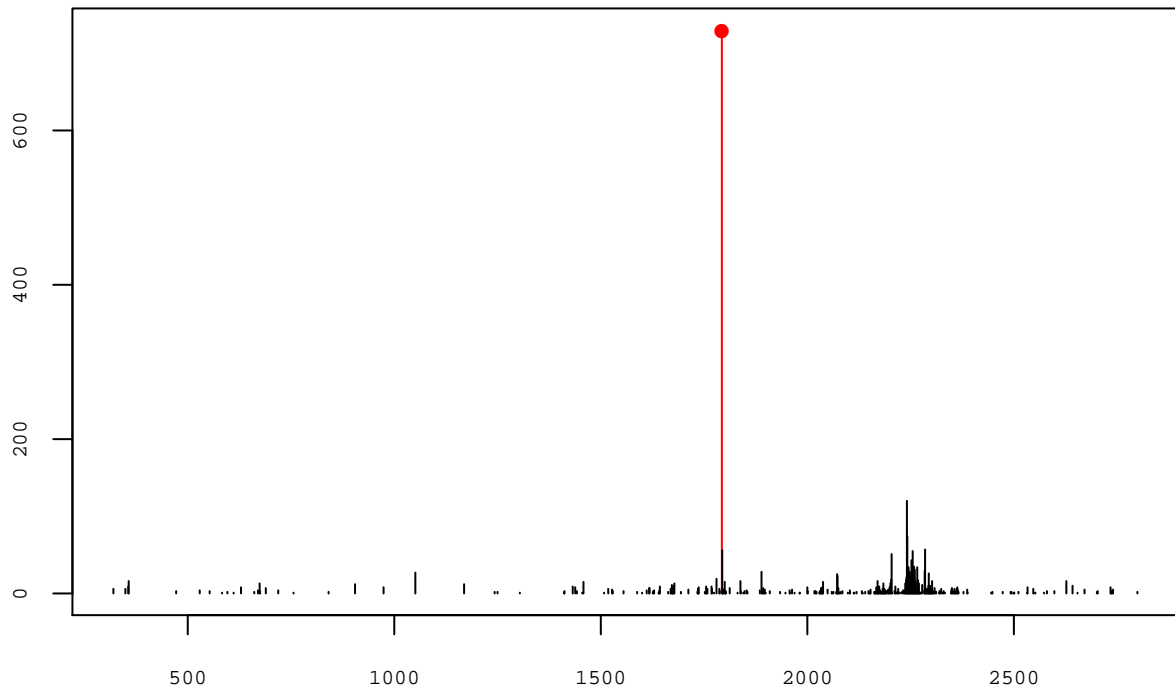
PggD2930_c0_g1_i1

5' ACACAGTGGGGAACCCTTCAGTCCAGTAACCC 3'

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3' CCTCGAGGGAAGTCAGGTT 5'

Fragment Abundance



Transcript position

Cleavage site: 1793	Tag abundance: 729	Weighted abundance: 182.25	Category: 0
sRNA abundance: 168	Alignment score: 3	MFE ratio: 0.768	p-value: 0.007