

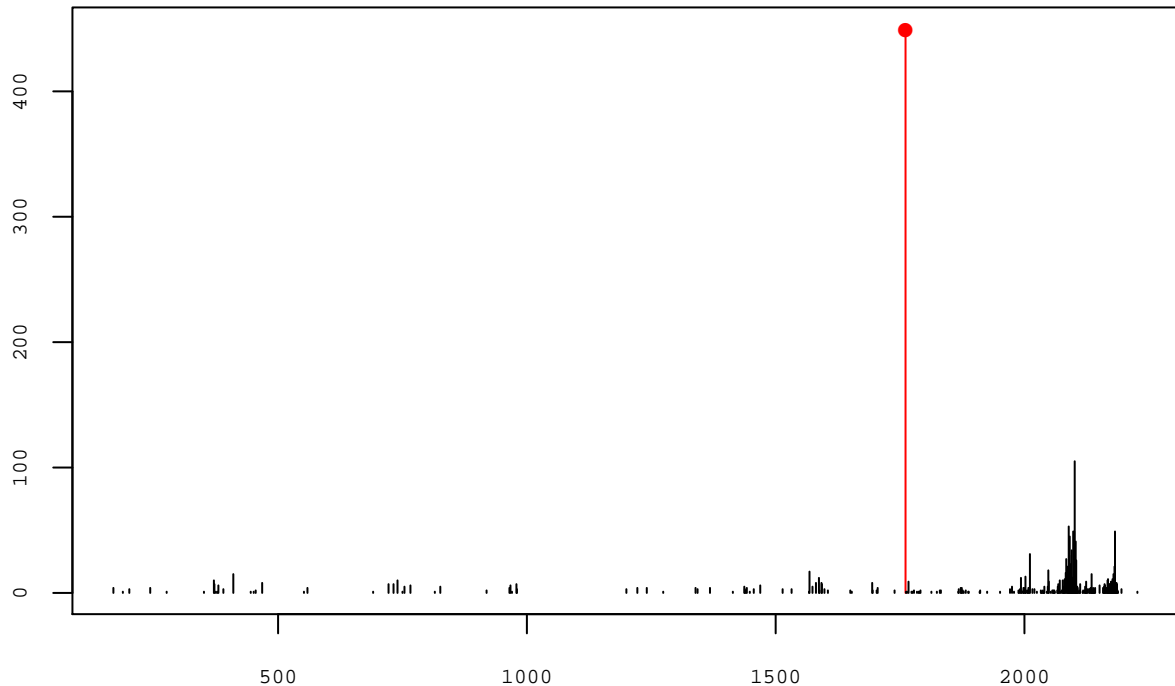
PggD5310_c30_g1_i1

5' TTCCACAGGGGGA-CCCTTCAGTCCAATCCATC '3

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3' CCCTCGAGGGAAGTCAGGTT '5

Fragment Abundance



Transcript position

Cleavage site: 1761	Tag abundance: 449	Weighted abundance: 224.5	Category: 0
sRNA abundance: 1130	Alignment score: 2.5	MFE ratio: 0.786	p-value: 0.004