

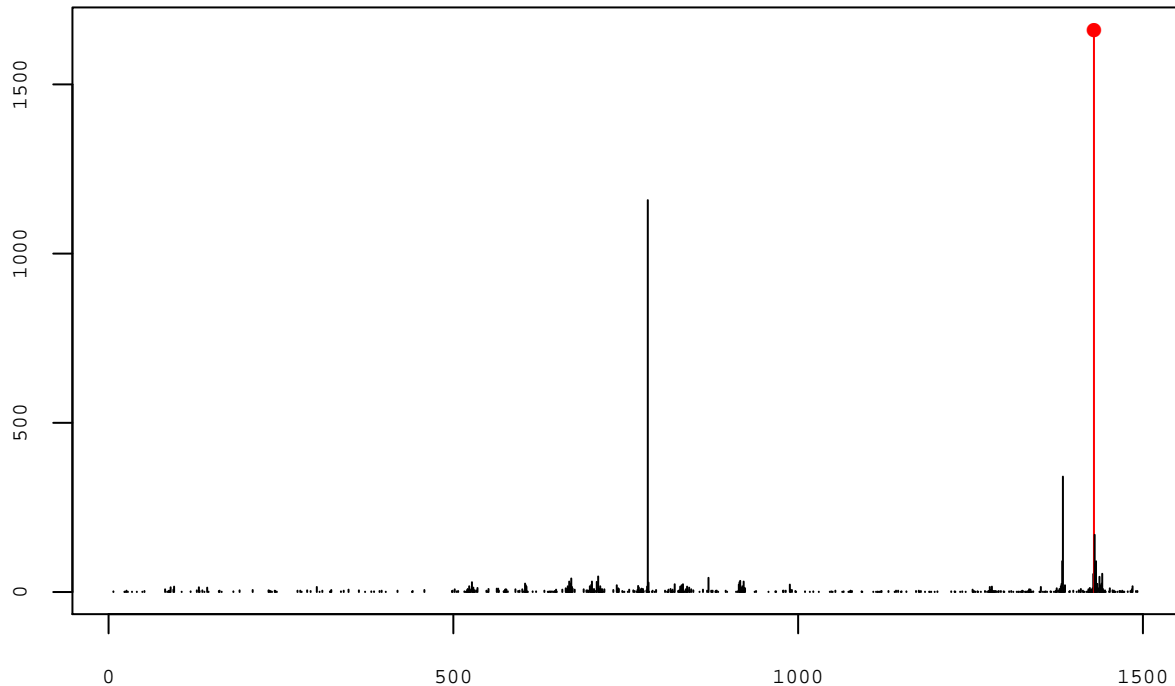
PggD5076_c3_g1_i1

5' CATTTCCTTTTAATTCCTTCT-GTAAATATCGCA '3

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3' AAAAATTAAGGAAGGTCATT '5

Fragment Abundance



Transcript position

Cleavage site: 1429	Tag abundance: 1661	Weighted abundance: 415.25	Category: 0
sRNA abundance: 8	Alignment score: 4	MFE ratio: 0.754	p-value: 0.003