

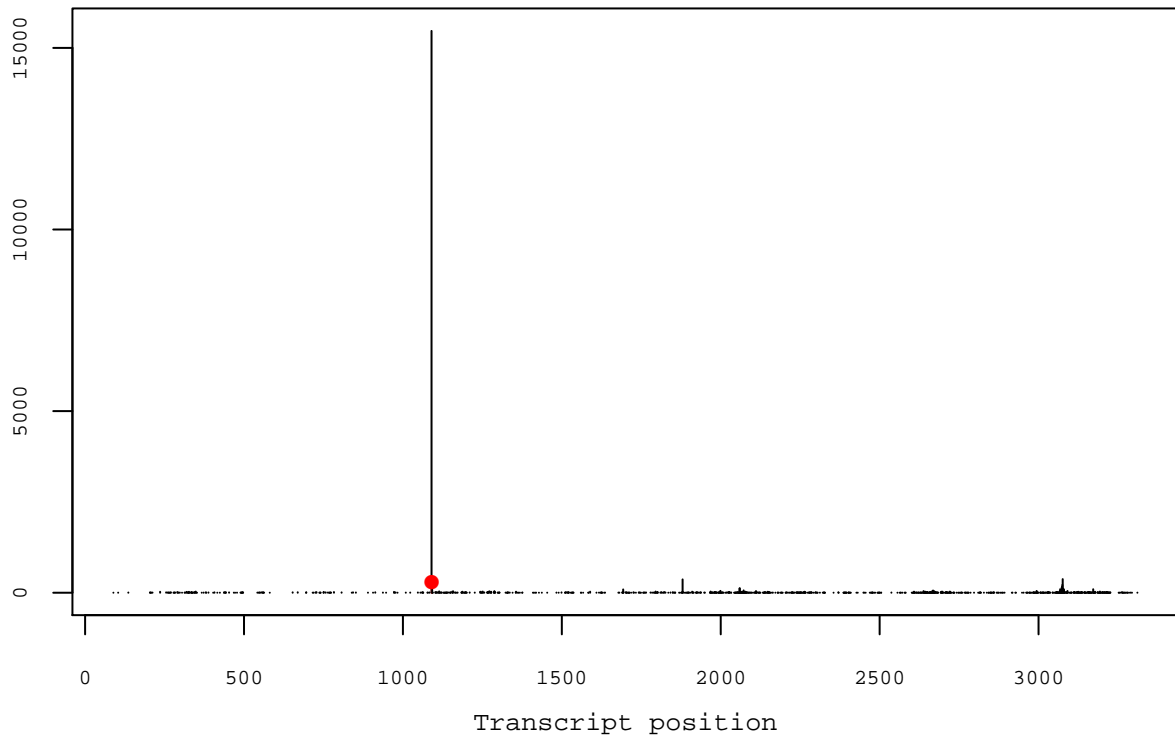
PggD7141_c1_g1_i1

5' ATGGTTGGTATGAAGCCTGGTCCGGATTCTAT '3

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3' CCCTTACTTCGGACCAGGCTT '5

Fragment Abundance



Cleavage site: 1091	Tag abundance: 291	Weighted abundance: 145.5	Category: 2
sRNA abundance: 97	Alignment score: 3	MFE ratio: 0.818	p-value: 0.037