

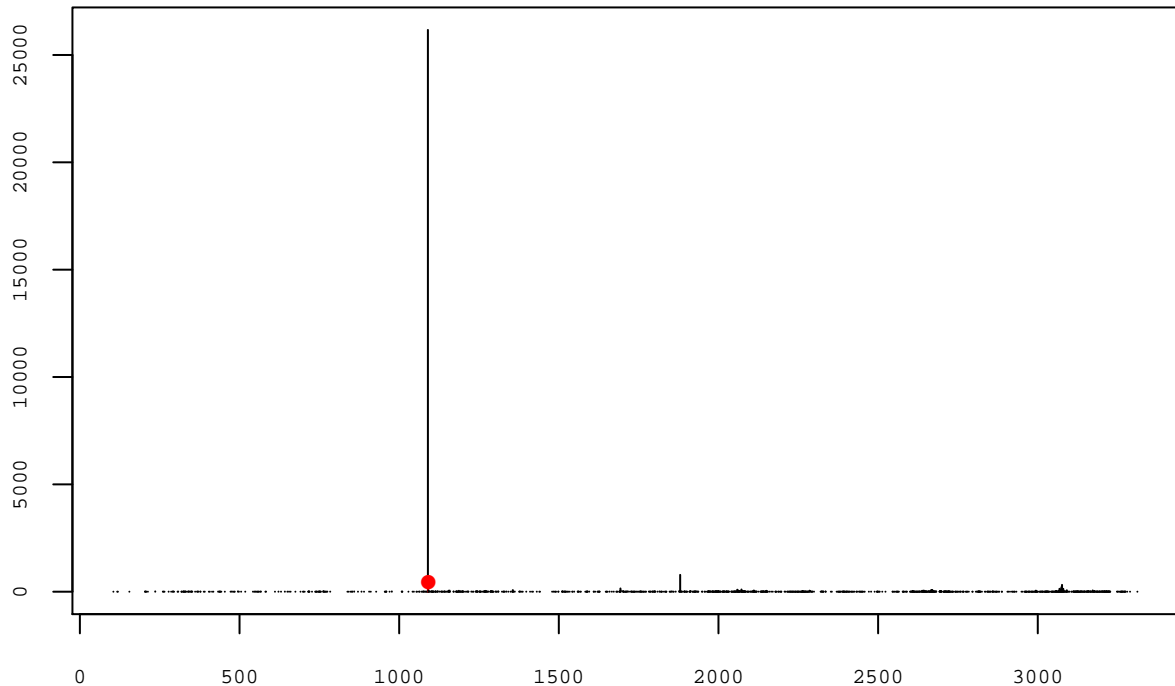
PggD7141_c1_g1_i1

5' ATGGTTGGTATGAAGCCTGGTCCGGATTCTAT '3

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3' CCCTTACTTCGGACCAGGCTT '5

Fragment Abundance



Transcript position

Cleavage site: 1091 Tag abundance: 456 Weighted abundance: 228 Category: 2
sRNA abundance: 240 Alignment score: 3 MFE ratio: 0.818 p-value: 0.038