

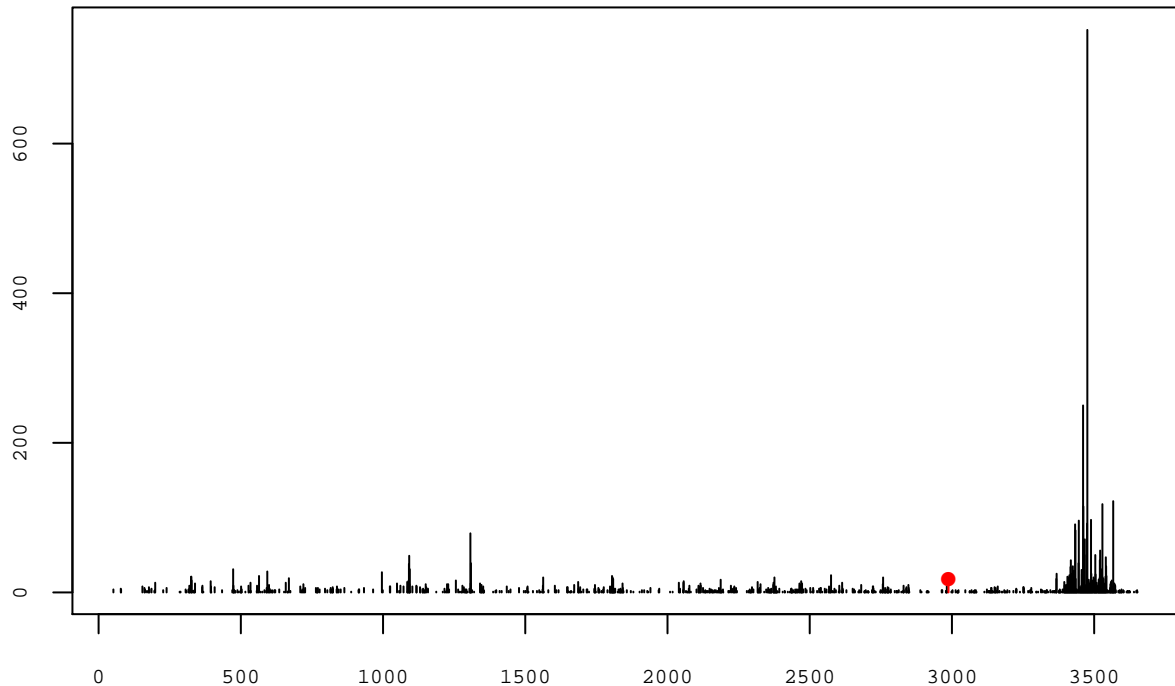
PggD11872_c0_g1_i2

5' CCTCTTAGATCAGGCTGGCAGCTT-GTATTTGT '3

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3' GTCTAGTACGACCGTCGAAGT '5

Fragment Abundance



Transcript position

Cleavage site: 2988 Tag abundance: 18 Weighted abundance: 18 Category: 2
sRNA abundance: 64 Alignment score: 3 MFE ratio: 0.756 p-value: 0.035