

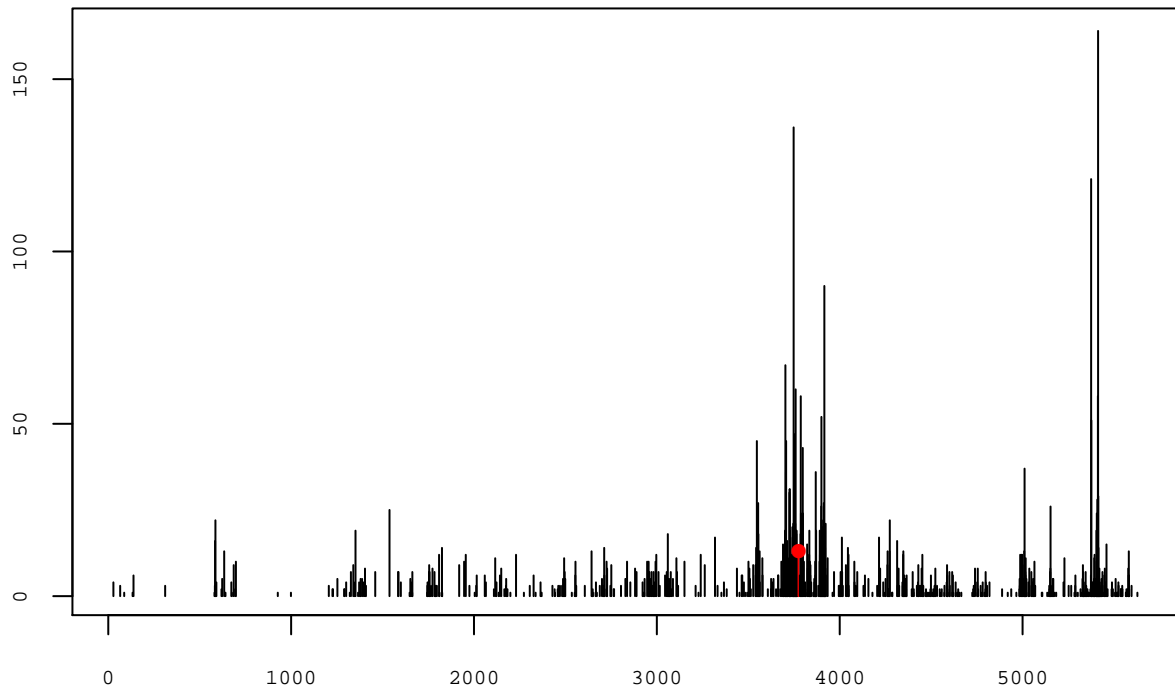
PggD9043_c6_g1_i1

5' GGTGAATGCTGTAAATTCTGAATCGACTCAAC '3

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3' ACGAATATTAAGATTAGCT '5

Fragment Abundance



Transcript position

Cleavage site: 3775 Tag abundance: 13 Weighted abundance: 13 Category: 2
sRNA abundance: 8 Alignment score: 4 MFE ratio: 0.734 p-value: 0.025