

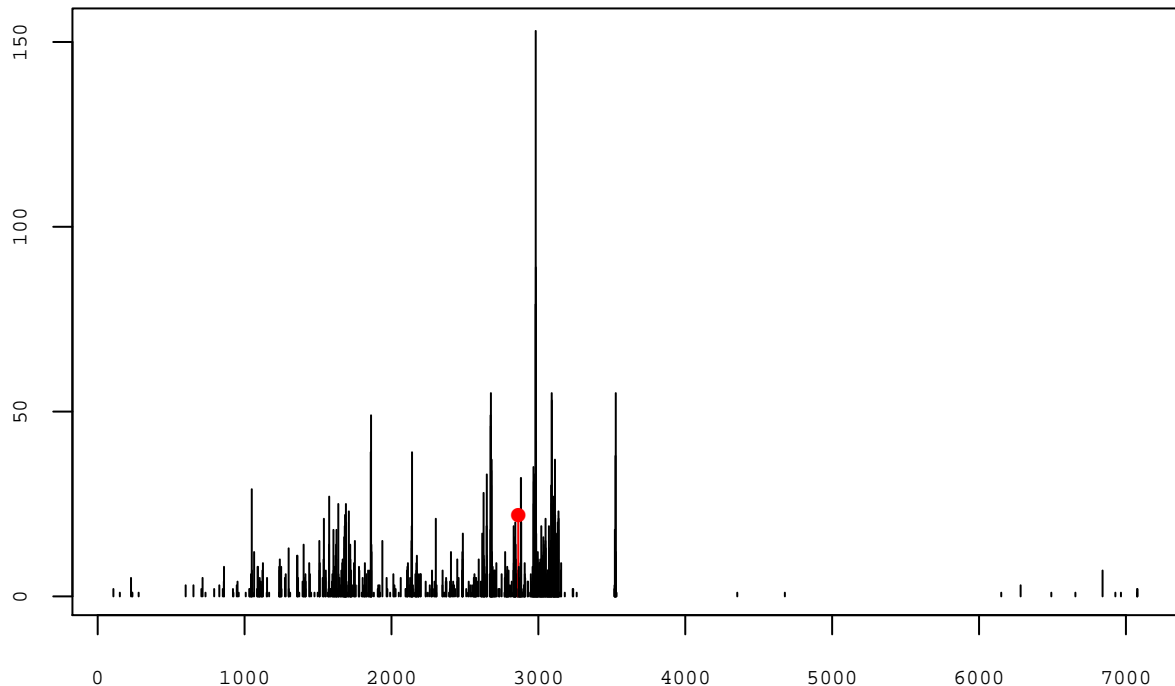
PggD15226_c4_g1_i2

5' TACAGGGTGGAGAAGAATGGTGGGAATGGTCAA '3

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3' CGTCTCTT-TTACCACCTTA '5

Fragment Abundance



Transcript position

Cleavage site: 2863 Tag abundance: 22 Weighted abundance: 22 Category: 2
sRNA abundance: 31 Alignment score: 3 MFE ratio: 0.804 p-value: 0.041