

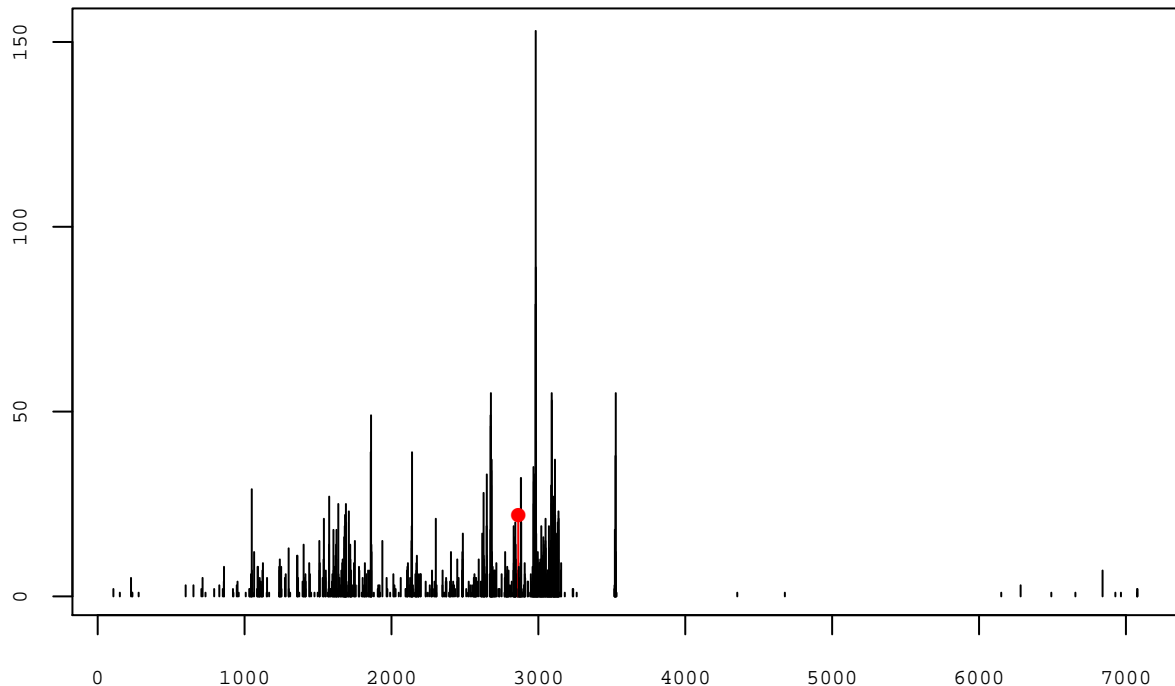
PggD15226_c4_g1_i2

5' TACAGGGTGGAGA-AGAATGGTGAATGGTCAA '3

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3' AACGTCTCTTTTACCACCTTA '5

Fragment Abundance



Transcript position

Cleavage site: 2863 Tag abundance: 22 Weighted abundance: 22 Category: 2
sRNA abundance: 408 Alignment score: 4 MFE ratio: 0.748 p-value: 0.041