

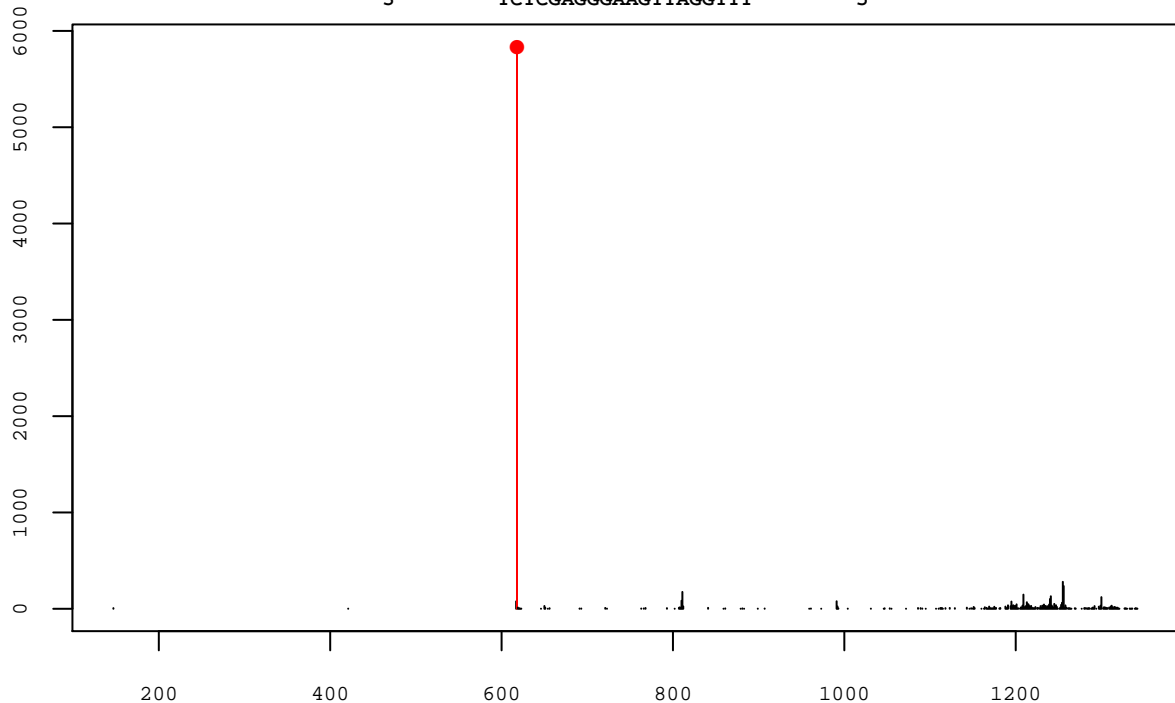
PggD6412_c14_g2_i1

5' GGAAATGGAGCTCCCTTCAAACCAAAGCCATC '3

o |||||

3' TCTCGAGGGAAGTTAGGTTT '5

Fragment Abundance



Transcript position

Cleavage site: 618 Tag abundance: 5834 Weighted abundance: 5834 Category: 0
sRNA abundance: 8438 Alignment score: 2.5 MFE ratio: 0.866 p-value: 0.001