

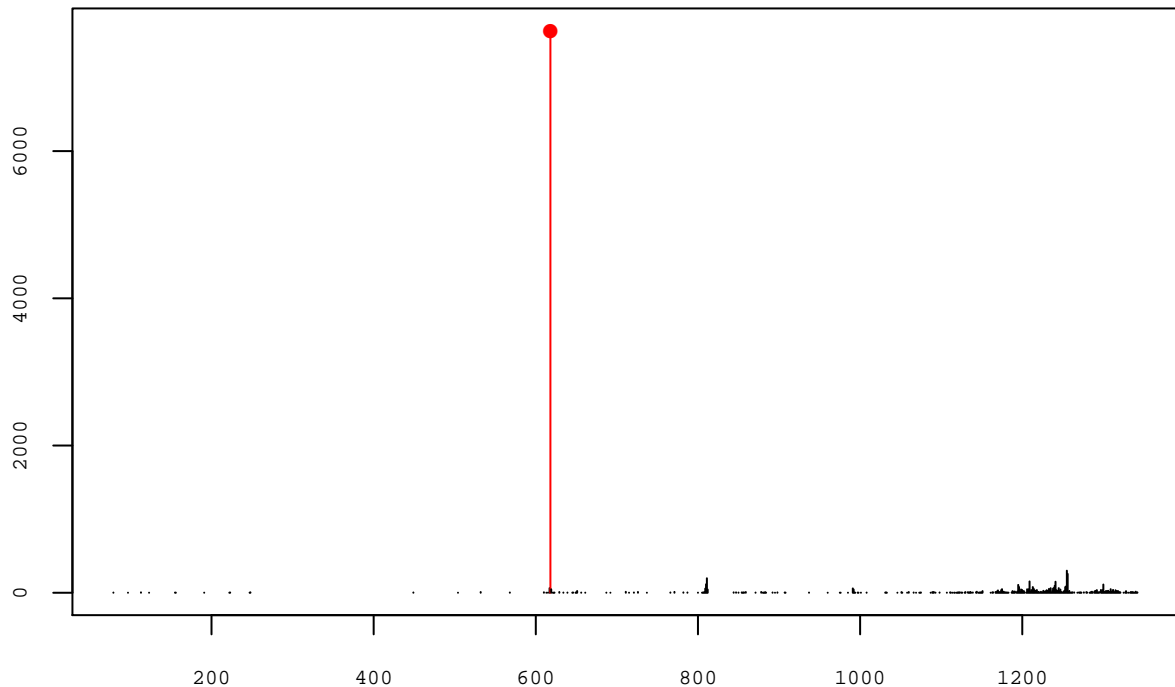
PggD6412_c14_g2_i1

5' GGAAATGGAGCTCCCTTCAAACCAAAGCCATC '3

oo|||||

3' GTCTCGAGGGAAGTTAGGTTT '5

Fragment Abundance



Transcript position

Cleavage site: 618 Tag abundance: 7633 Weighted abundance: 7633 Category: 0
sRNA abundance: 8957 Alignment score: 3 MFE ratio: 0.828 p-value: 0.001