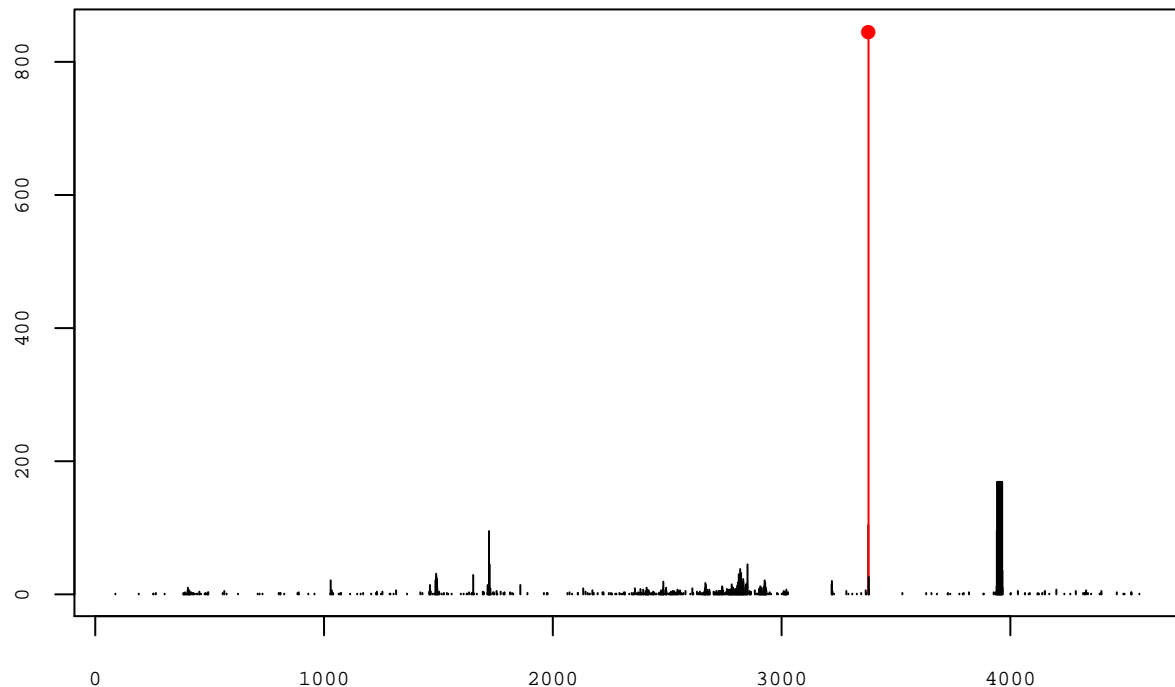


PggD7328_c7_g1_i1

5' ATGTTTAGGTGTTAAATATGTAAATACATAGG '3
||o||||||||||||||||||
3' AAAGTCCACAATTTATACATTAT '5

Fragment Abundance



Transcript position

Cleavage site: 3380 Tag abundance: 845 Weighted abundance: 2.067 Category: 0
sRNA abundance: 5 Alignment score: 1.5 MFE ratio: 0.921 p-value: 0.016