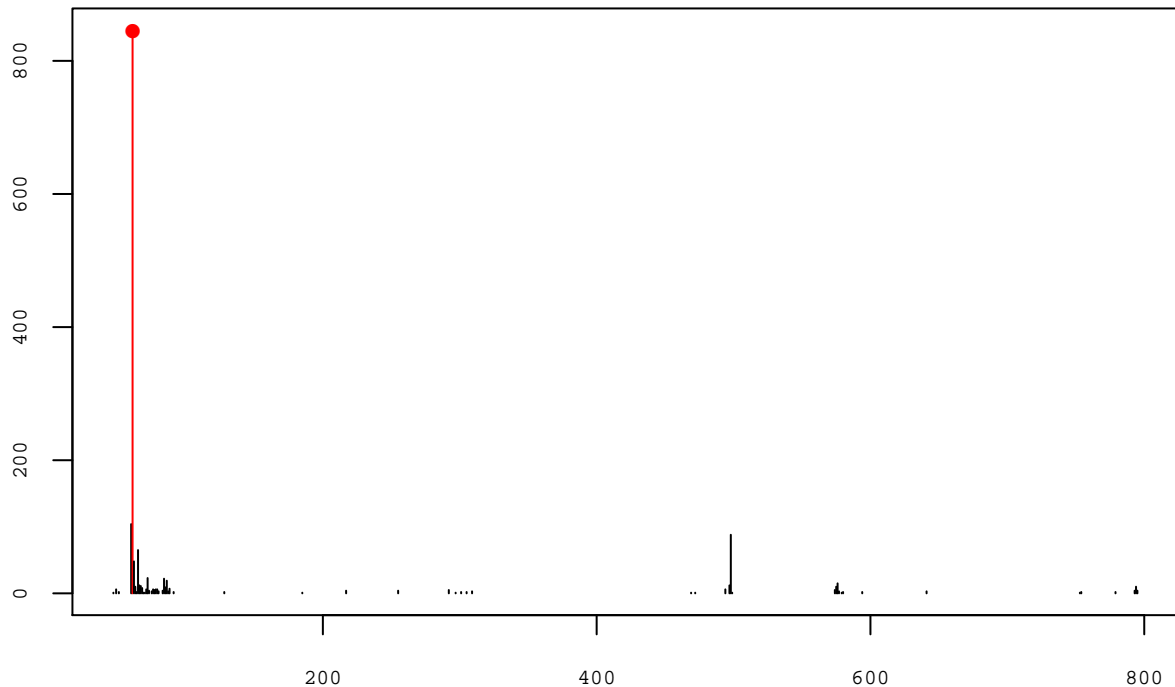


PggD12654_c92_g1_i1

5' AGTTT TAGGTGTTAAATATGTAAATACATAGG '3
|||o||||||||||||||||||
3' AAAGTCCACAATT TATACATTAT '5

Fragment Abundance



Transcript position

Cleavage site: 61 Tag abundance: 845 Weighted abundance: 2.067 Category: 0
sRNA abundance: 5 Alignment score: 0.5 MFE ratio: 0.928 p-value: 0.004