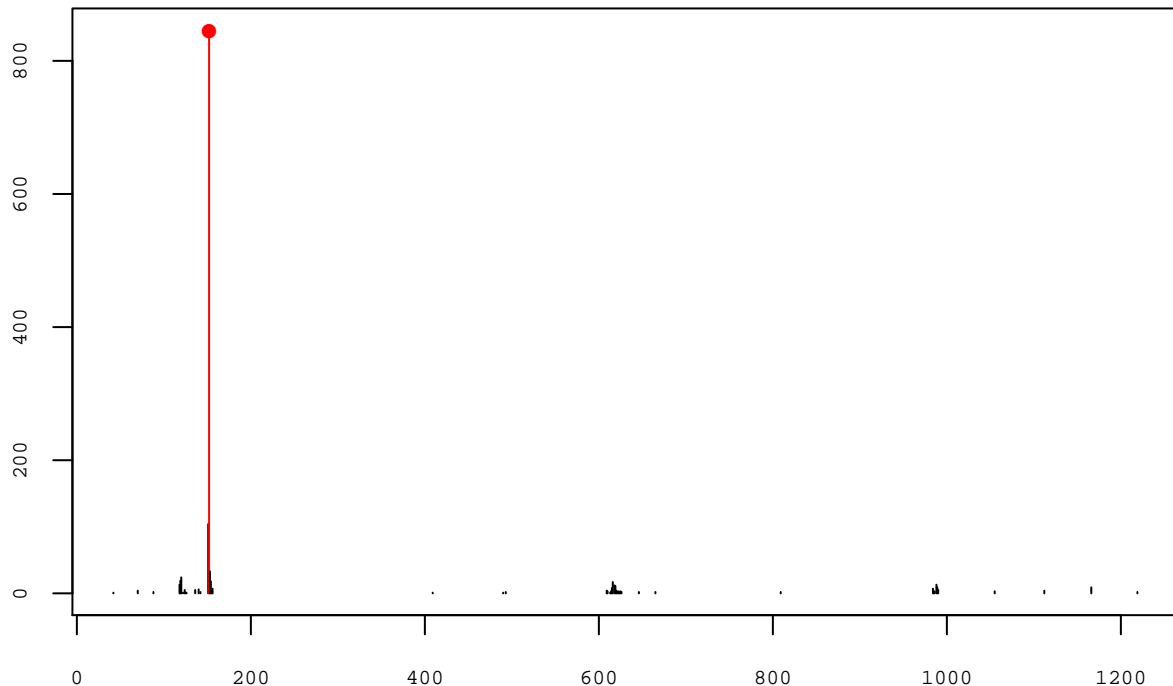


PggD6971_c107_g1_i1

5' AAATTAGGTGTTAAATATGTAAATACATAGG '3
||o||||||||||||||||||
3' AAAGTCCACAATTATACATTAT '5

Fragment Abundance



Transcript position

Cleavage site: 152 Tag abundance: 845 Weighted abundance: 2.067 Category: 0
sRNA abundance: 5 Alignment score: 1.5 MFE ratio: 0.921 p-value: 0.049