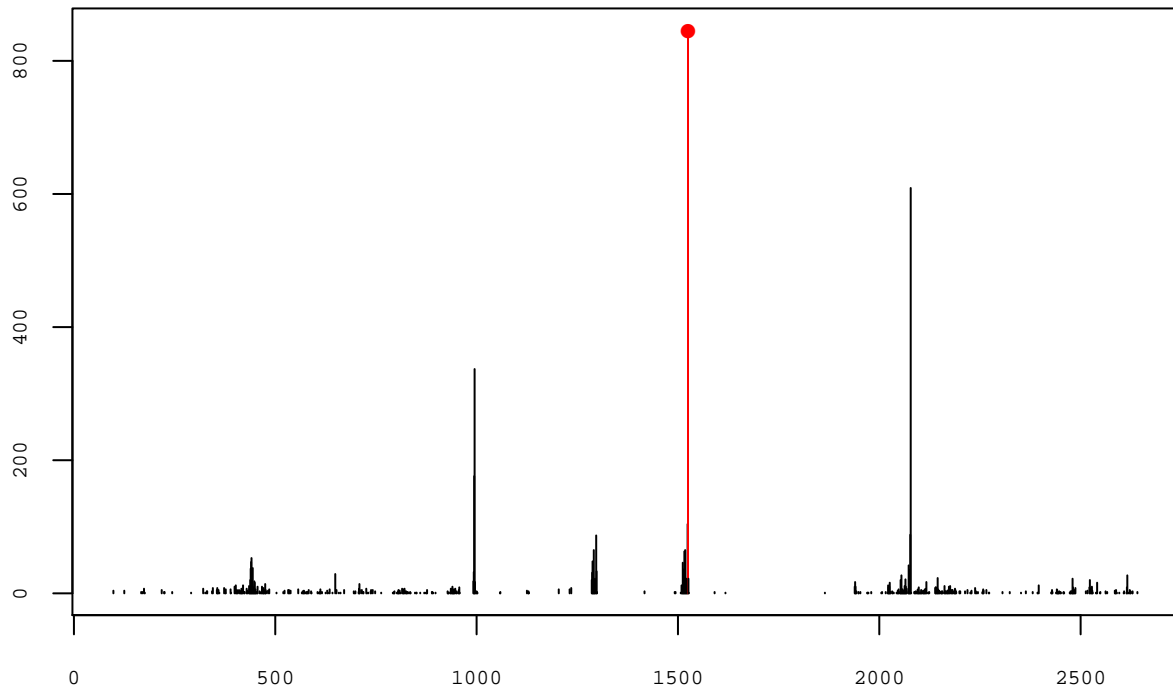


PggD14287_c19_g1_i1

5' AATTTTAGGTGTTAGATATGTAAATACATAGG '3
|||o|||||||o|||||||
3' AAAGTCCACAATTATACATTAT '5

Fragment Abundance



Transcript position

Cleavage site: 1525 Tag abundance: 845 Weighted abundance: 2.067 Category: 0
sRNA abundance: 5 Alignment score: 1.5 MFE ratio: 0.931 p-value: 0.031