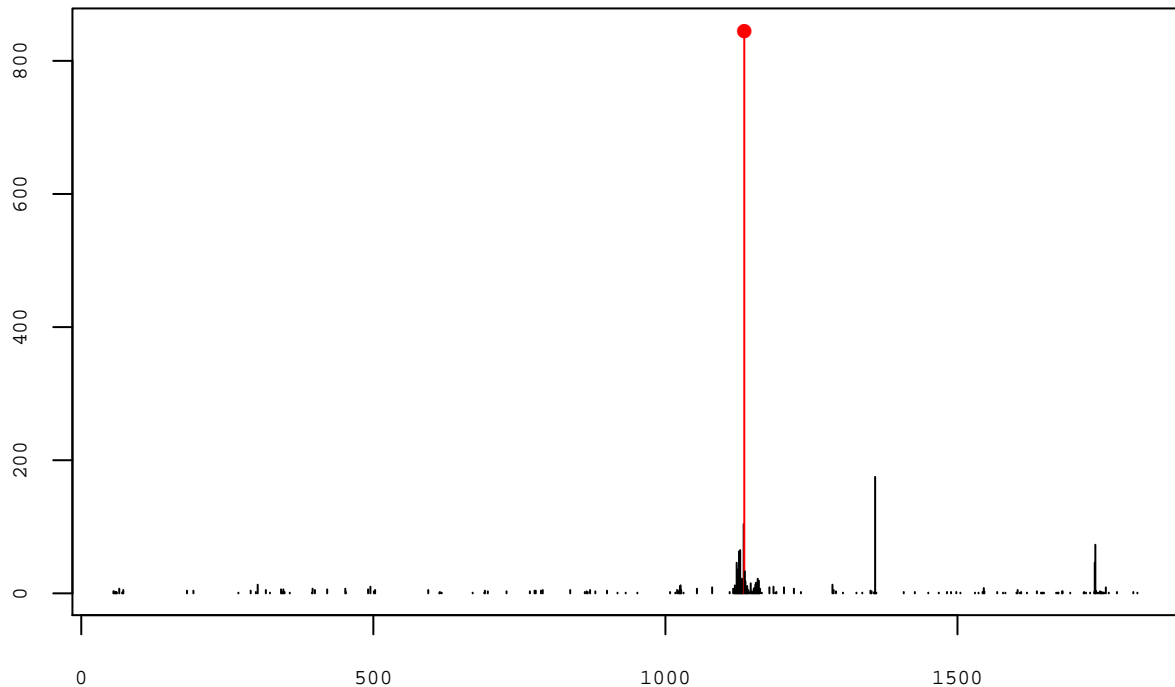


PggD7328\_c12\_g1\_i1

5' AATTTTAGGTGTTAGATATGTAAATACATAGG '3  
|||o|||||||o|||||||  
3' AAAGTCCACAATTTATACATTAT '5

Fragment Abundance



Transcript position

Cleavage site: 1135 Tag abundance: 845 Weighted abundance: 2.067 Category: 0  
sRNA abundance: 5 Alignment score: 1.5 MFE ratio: 0.931 p-value: 0.046