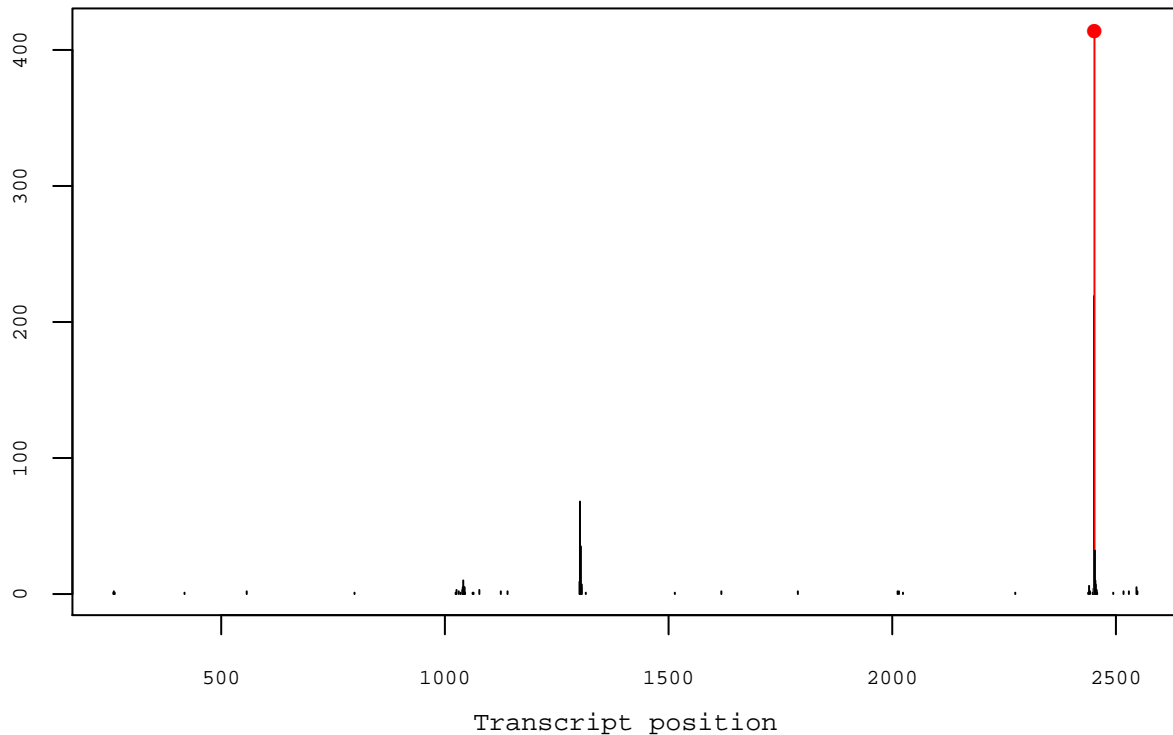


5' ATTTT TAGGTGTTAAATATGTAAATACATAAG '3
|||o|||||||
3' AAAGTCCACAATT TATACATTAT '5

Fragment Abundance



Cleavage site: 2452 Tag abundance: 414 Weighted abundance: 0.653 Category: 0
sRNA abundance: 5 Alignment score: 0.5 MFE ratio: 0.928 p-value: 0.001