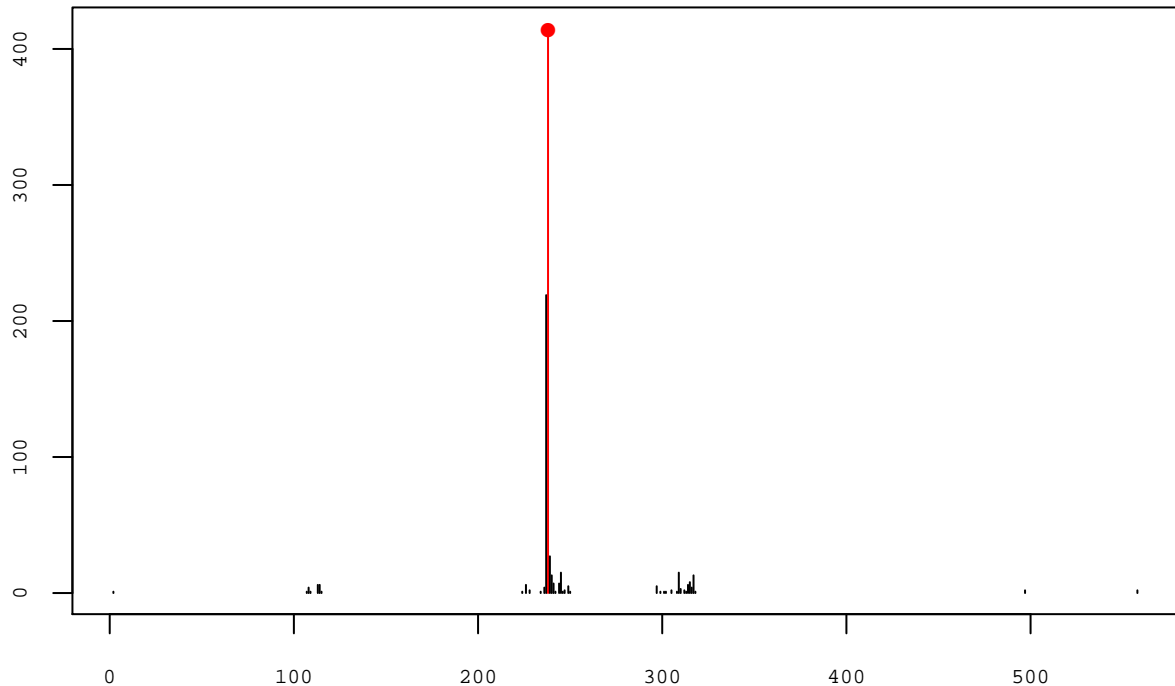


5' AATTTTAGGTGTTAAATATGTAAATACATAAG '3
|||o||||||||||||||||||
3' AAAGTCCACAATTATACATTAT '5

Fragment Abundance



Transcript position

Cleavage site: 238 Tag abundance: 414 Weighted abundance: 0.653 Category: 0
sRNA abundance: 5 Alignment score: 0.5 MFE ratio: 0.928 p-value: 0.004