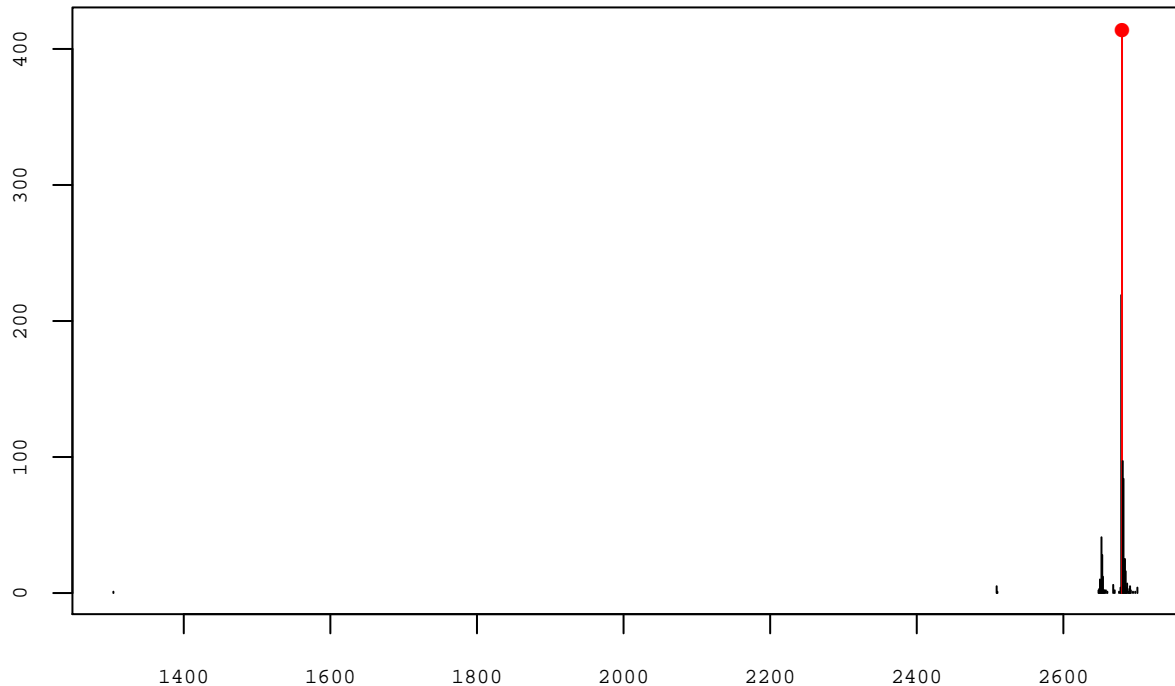


PggD6793\_c58\_g2\_i1

5' AAGTTTAGGTGTTAAATATGTAAATACATAAG '3  
||o||||||||||||||||||  
3' AAAGTCCACAATTATACATTAT '5

Fragment Abundance



Transcript position

Cleavage site: 2680 Tag abundance: 414 Weighted abundance: 0.653 Category: 0  
sRNA abundance: 5 Alignment score: 1.5 MFE ratio: 0.921 p-value: 0.03