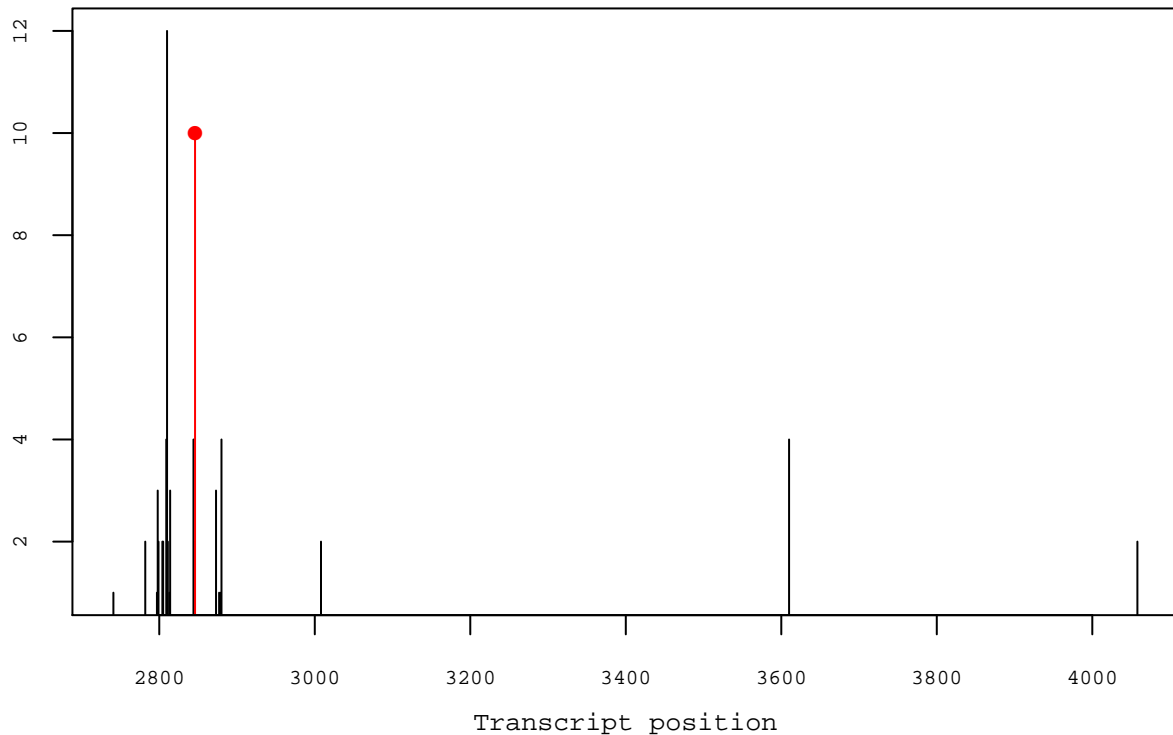


5' ACTGTGATCTACGGCCAGGATCGGACACCGCA '3  
o ||||| |||||  
3' GCACTAGATGCCAGTCCTAGCCTA '5

Fragment Abundance



Cleavage site: 2846 Tag abundance: 10 Weighted abundance: 2.5 Category: 2  
sRNA abundance: 111 Alignment score: 3.5 MFE ratio: 0.86 p-value: 0.003