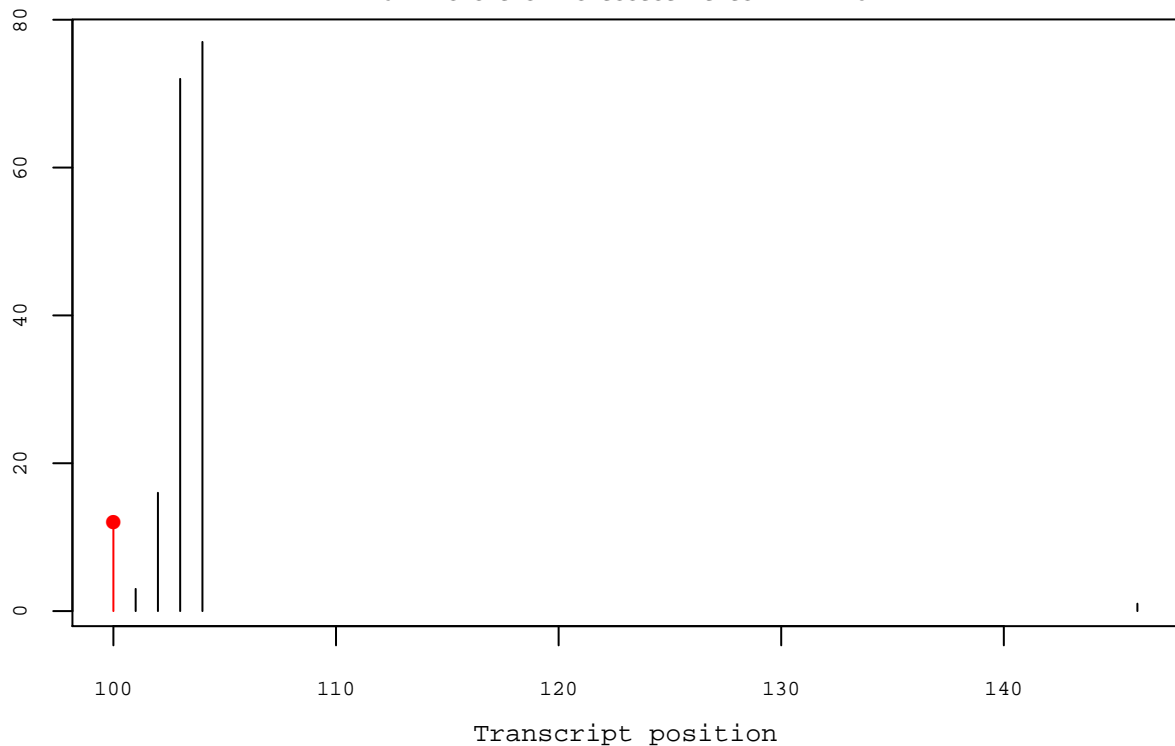


5' GCTGCTGTCATACAGTCGCGATGTGCGGCTGT '3  
oo|||||||o|||||||  
3' GTGACAGTATGTCGGCGCTACACG '5

Fragment Abundance



Cleavage site: 100 Tag abundance: 12 Weighted abundance: 3.917 Category: 3  
sRNA abundance: 190 Alignment score: 2 MFE ratio: 0.913 p-value: 0.024