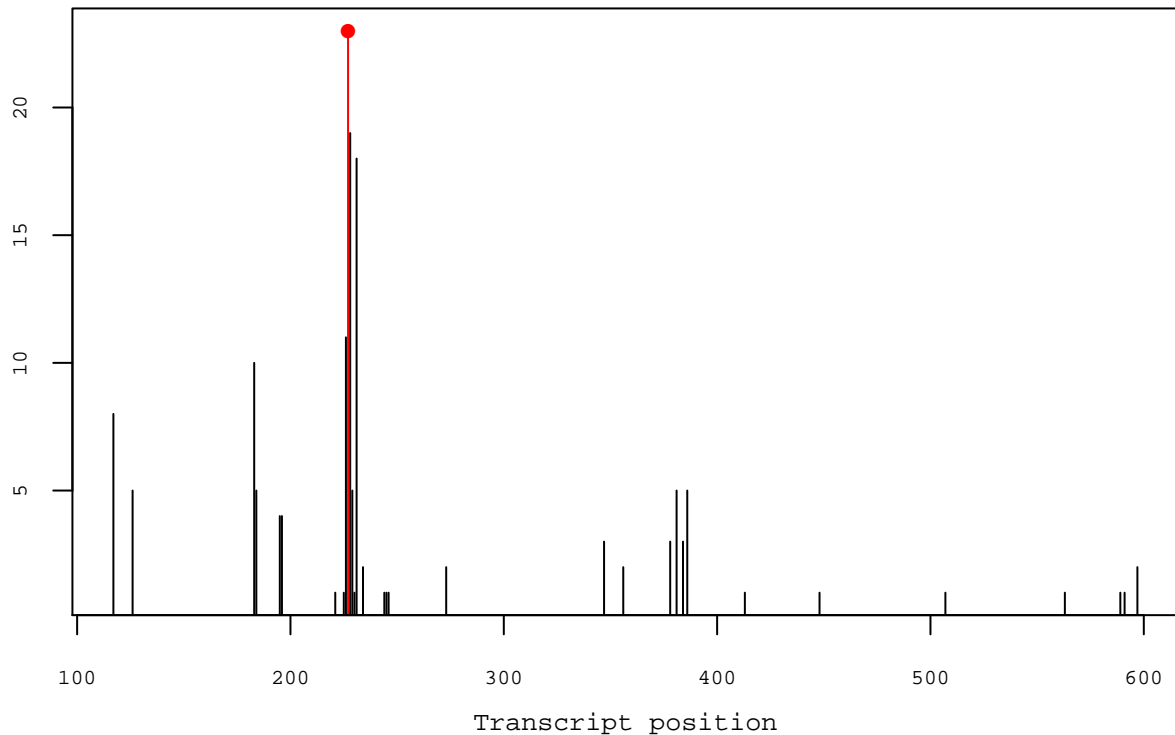


5' TGTCTGACATTTTGGTTTGTGCGTTTGGAAAA '3
||||| |||||
3' AGACTGACAAACCAACACGCAAA '5

Fragment Abundance



Cleavage site: 227 Tag abundance: 23 Weighted abundance: 10.167 Category: 0
sRNA abundance: 6 Alignment score: 2 MFE ratio: 0.794 p-value: 0.015