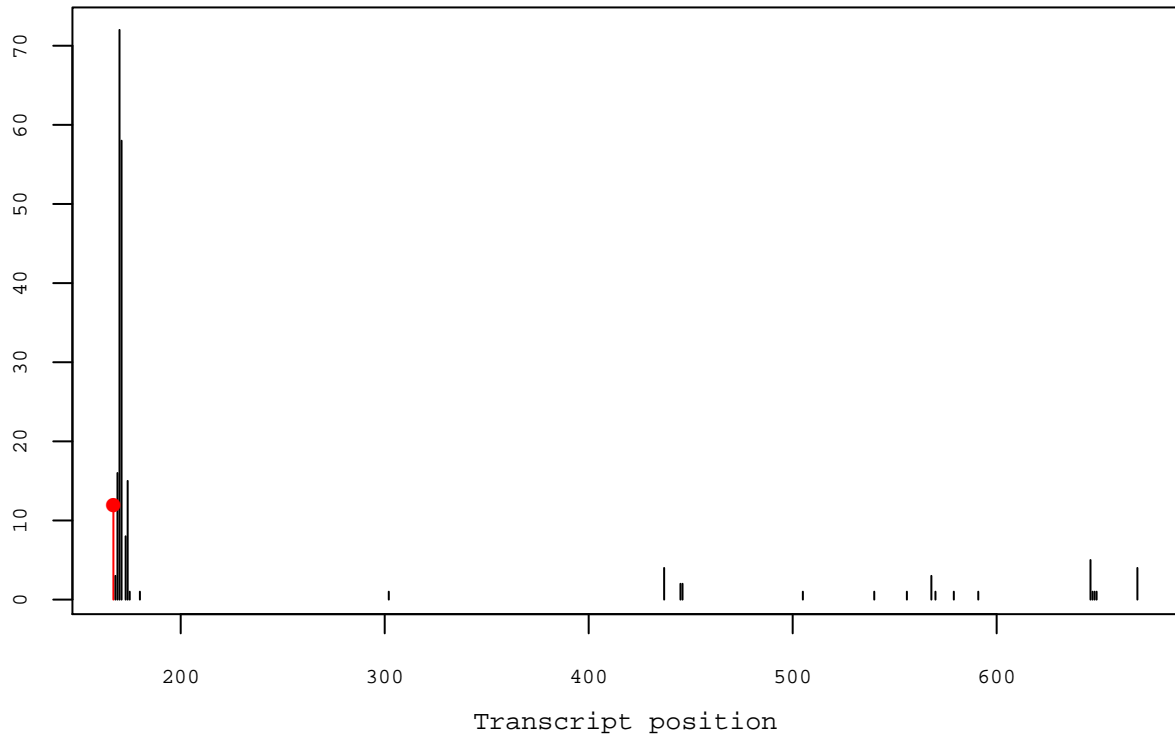


5' ACCACTGTCATCCAGCCGCGATGTGCGGCTGT '3
||| ||| |
3' GTGACAGTATGTTGGCGCTACACA '5

Fragment Abundance



Cleavage site: 167
sRNA abundance: 31

Tag abundance: 12
Alignment score: 3

Weighted abundance: 3.917
MFE ratio: 0.88

Category: 3
p-value: 0.029