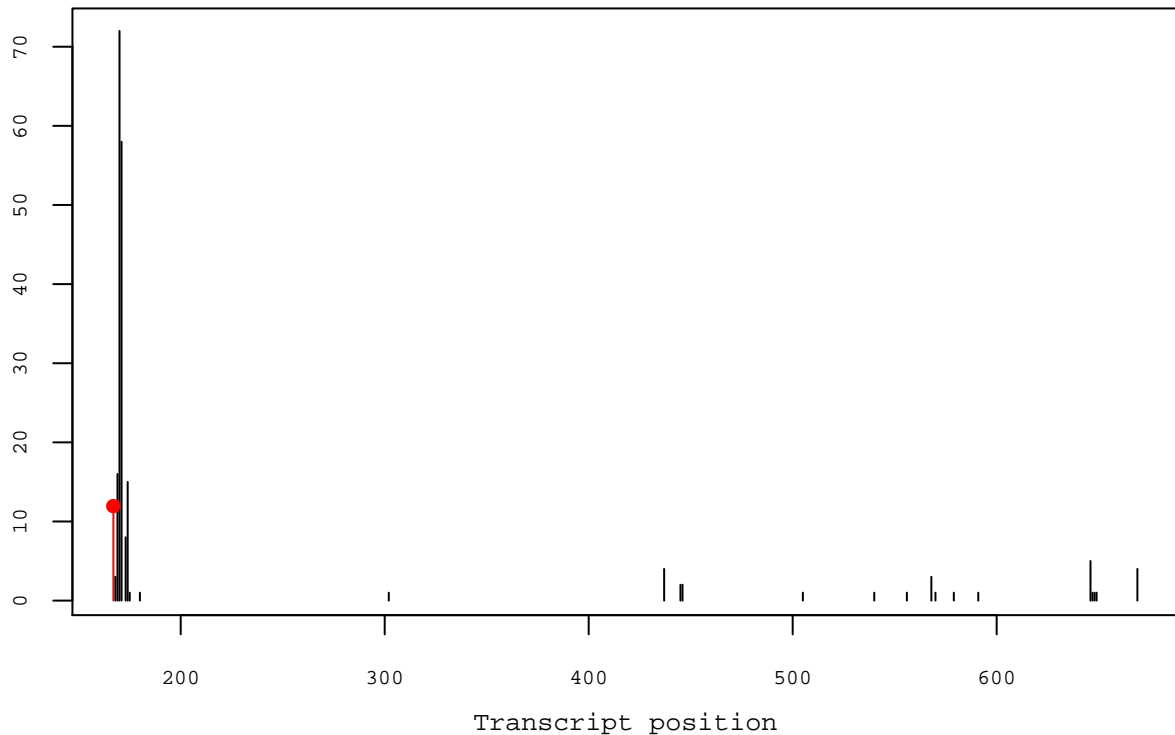


5' ACCACTGTCATCCAGCCGCGATGTGCGGCTGT '3
 ||||| |||○ |||||
 3' GTGACAGTATGTTGGCGCTACACG '5

Fragment Abundance



Cleavage site: 167

sRNA abundance: 45

Tag abundance: 12

Alignment score: 2

Weighted abundance: 3.917

MFE ratio: 0.897

Category: 3

p-value: 0.029