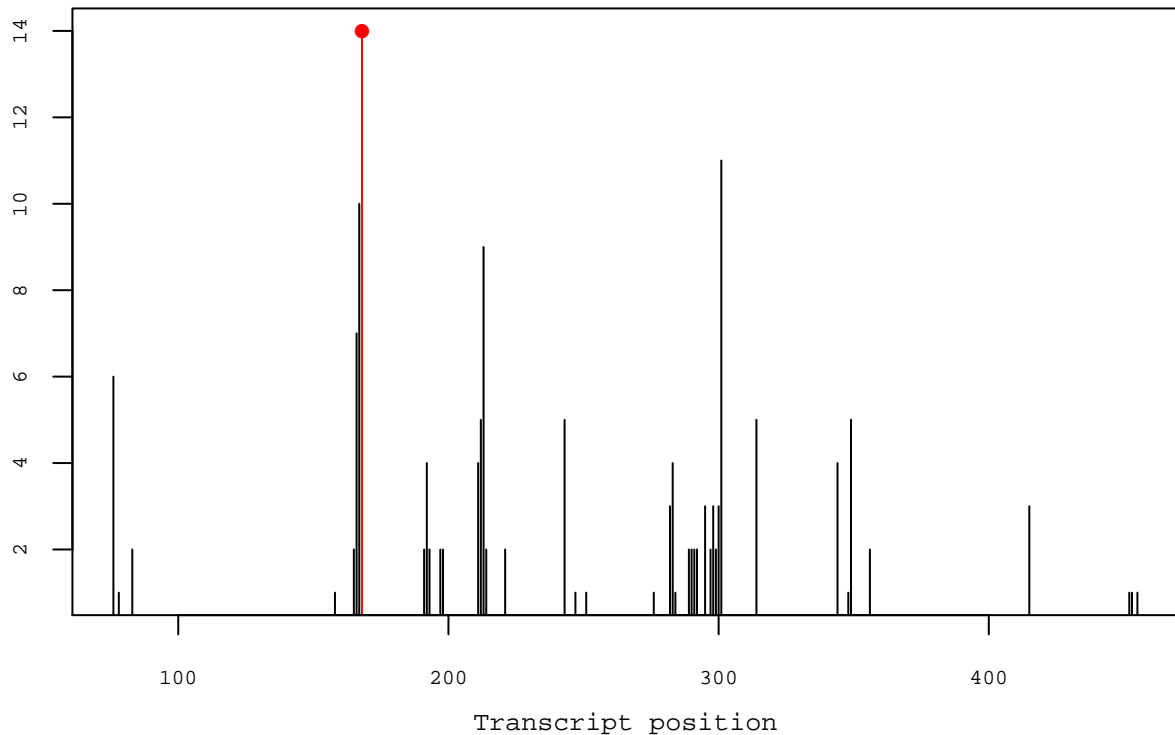


5' AAATTTTCTAAGTCATTCCAAAATTATCTTT '3
 ||o||||||||||||||||||
 3' TAGAAGATTCAGTAAGGTTTTAAA '5

Fragment Abundance



Cleavage site: 168

sRNA abundance: 29

Tag abundance: 14

Alignment score: 0.5

Weighted abundance: 0.406

MFE ratio: 0.92

Category: 0

p-value: 0.038