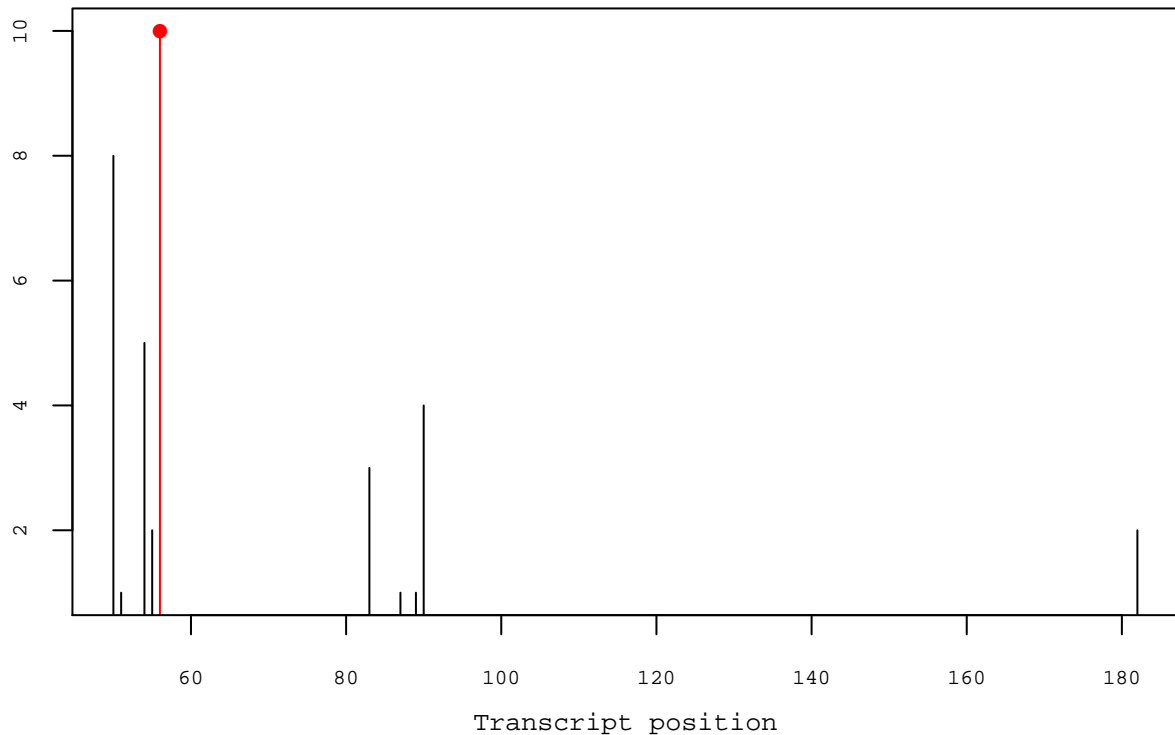


5' ATCATGATCAACAGTCAGGATCGGACACCGCA '3
|| ||||| || |||||
3' AG-ACTAGATGCCAGTCCTAGCCTG '5

Fragment Abundance



Cleavage site: 56
srRNA abundance: 109

Tag abundance: 10
Alignment score: 3

Weighted abundance: 2.5
MFE ratio: 0.703

Category: 0
p-value: 0.013