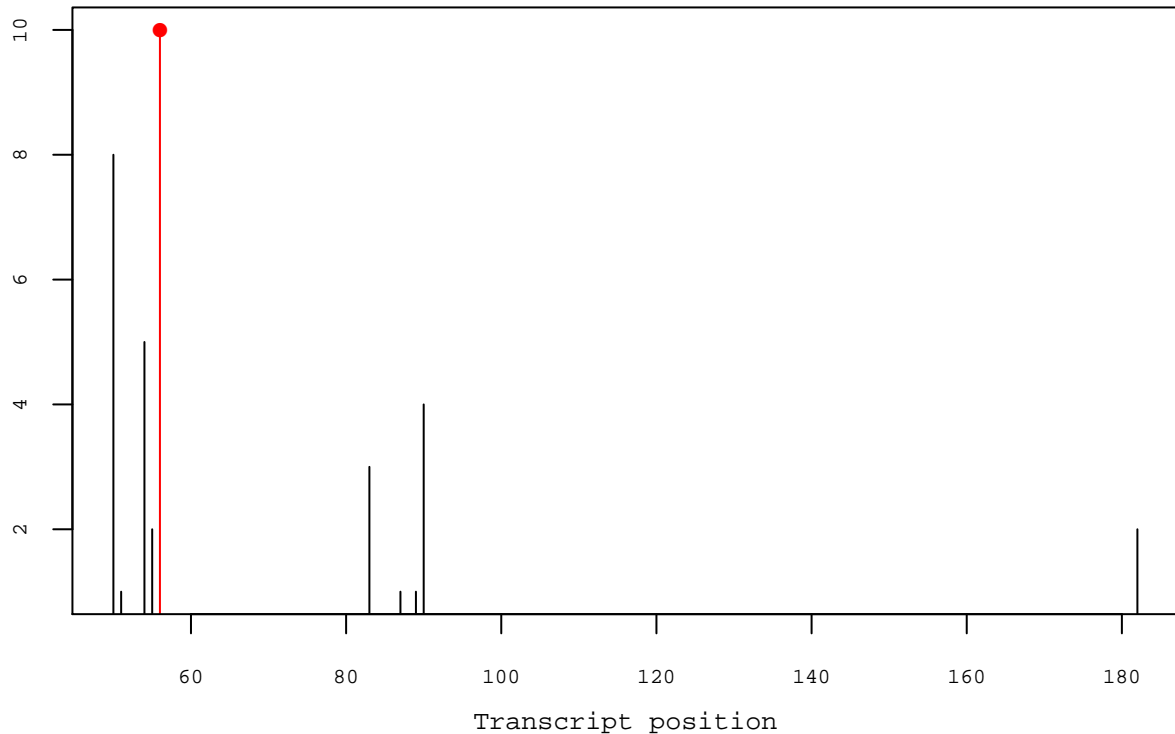


5' ATCATGATCAACAGTCAGGATCGGACACCGCA '3  
| | | | | | | | | | | | | | | | | | | | | |  
3' GCACTAGATGTCAGTCCTAGCCTG '5

Fragment Abundance



Cleavage site: 56  
sRNA abundance: 1891

Tag abundance: 10  
Alignment score: 2

Weighted abundance: 2.5  
MFE ratio: 0.805

Category: 0  
p-value: 0.007