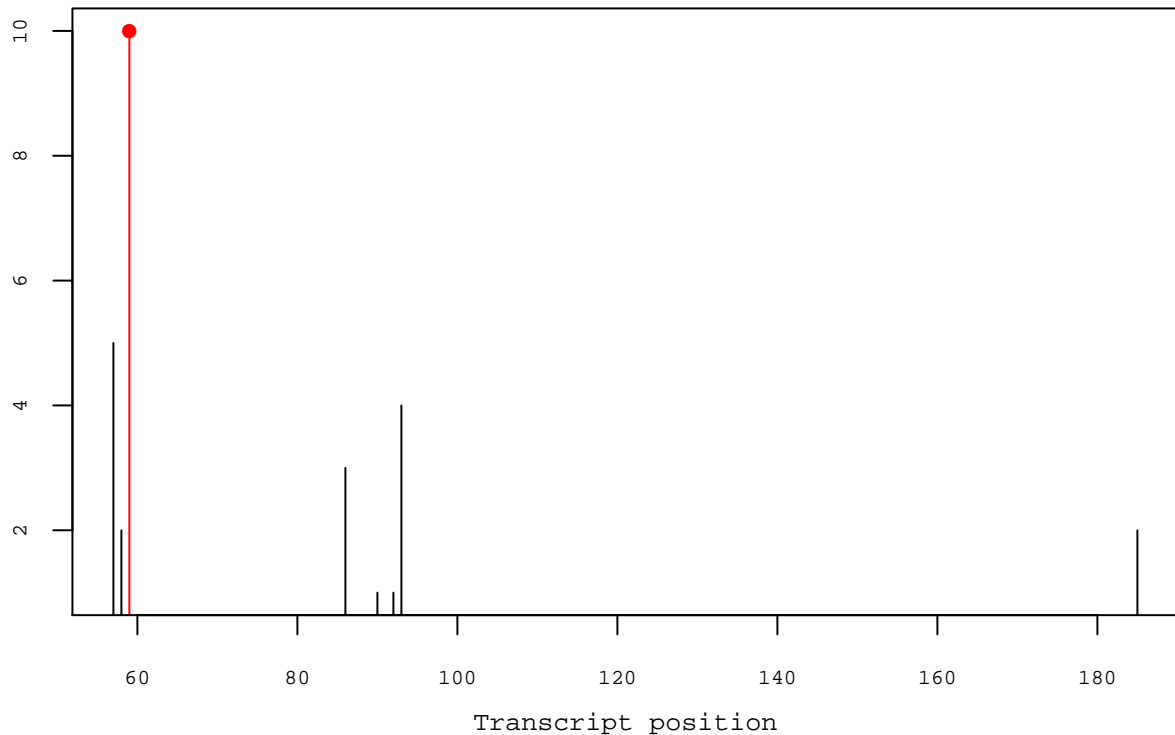


5' TTCGTGATCTATGGTCAGGATCGGACACCGCA '3
||| ||| | oo ||| ||| ||| |||
3' GCACTAGATGTCAGTCCTAGCCTG '5

Fragment Abundance



Cleavage site: 59
sRNA abundance: 1891

Tag abundance: 10
Alignment score: 1

Weighted abundance: 2.5
MFE ratio: 0.931

Category: 0
p-value: 0.003