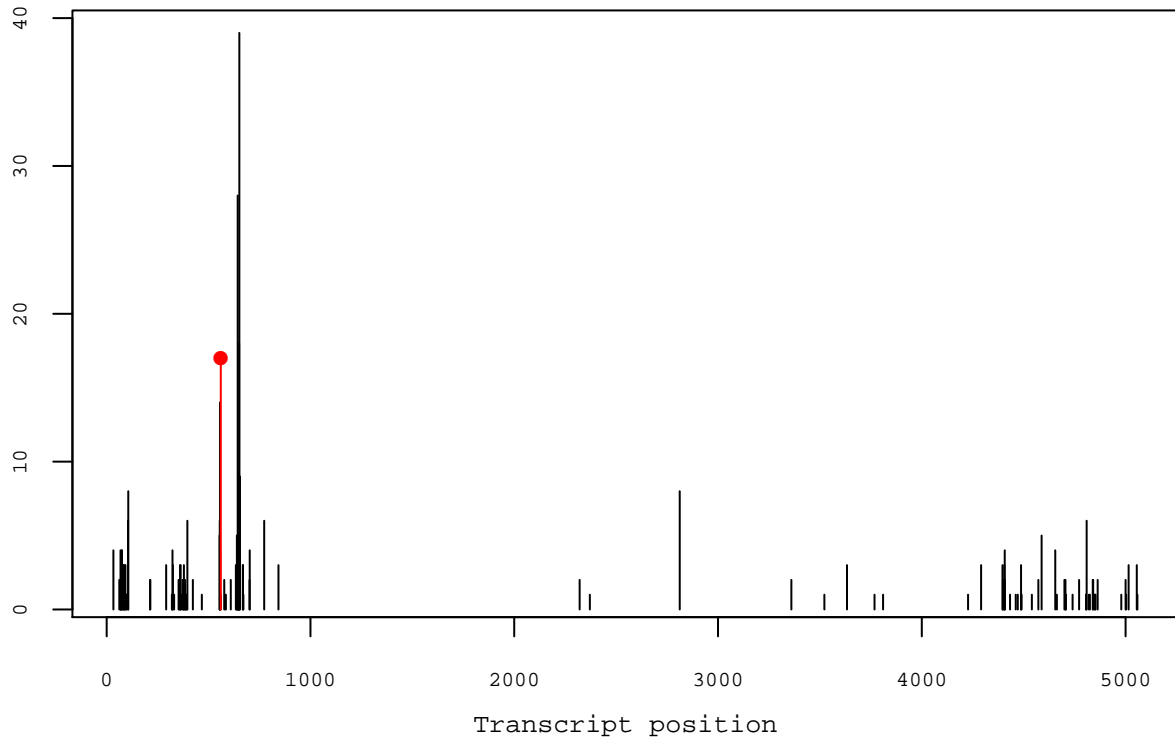


5' AATGCA~TCGGTGTCCGATCCTGACCGTAGATC '3
|| |||||
3' GTACACCACAGGCTAGGACTGGC '5

Fragment Abundance



Cleavage site: 560

Tag abundance: 17

Weighted abundance: 5.667

Category: 2

sRNA abundance: 8

Alignment score: 3

MFE ratio: 0.828

p-value: 0.023