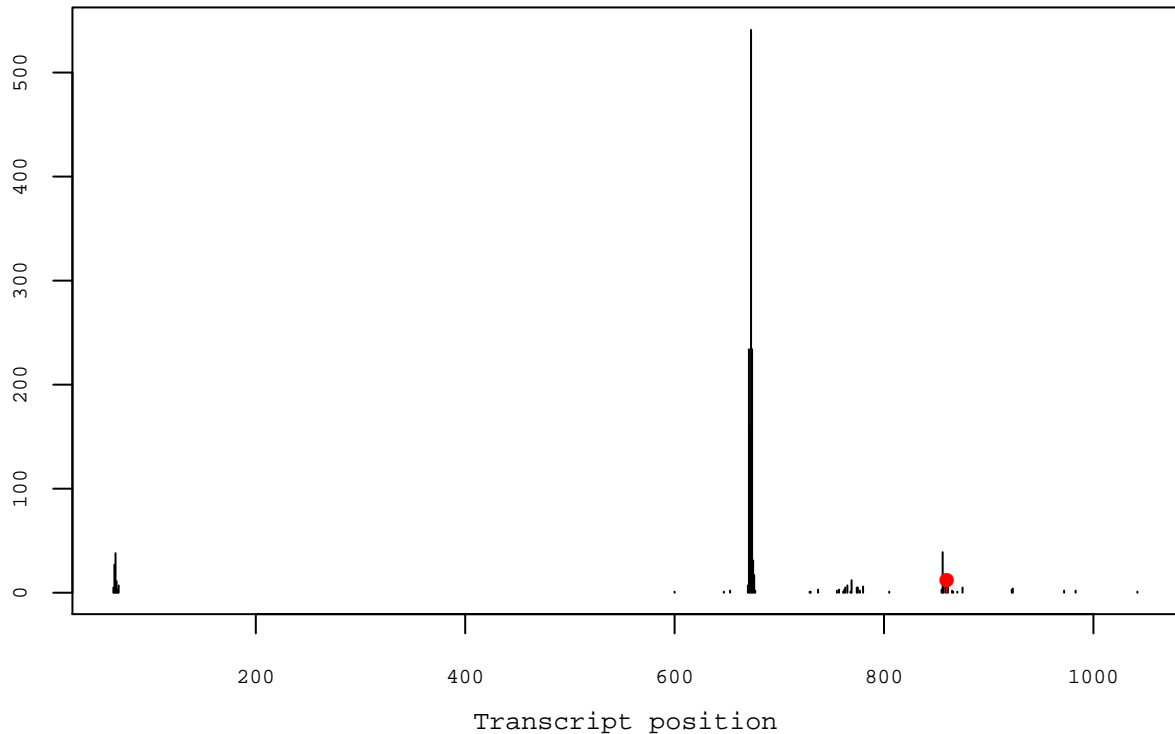


5' CTTTCTTAACTTAAGTC-GAATCTAAAACATCT '3
 |||||
 3' GAATTGAATTCAGATTAGGTT '5

Fragment Abundance



Cleavage site: 860

Tag abundance: 12

Weighted abundance: 0.039

Category: 2

sRNA abundance: 6

Alignment score: 4

MFE ratio: 0.748

p-value: 0.011