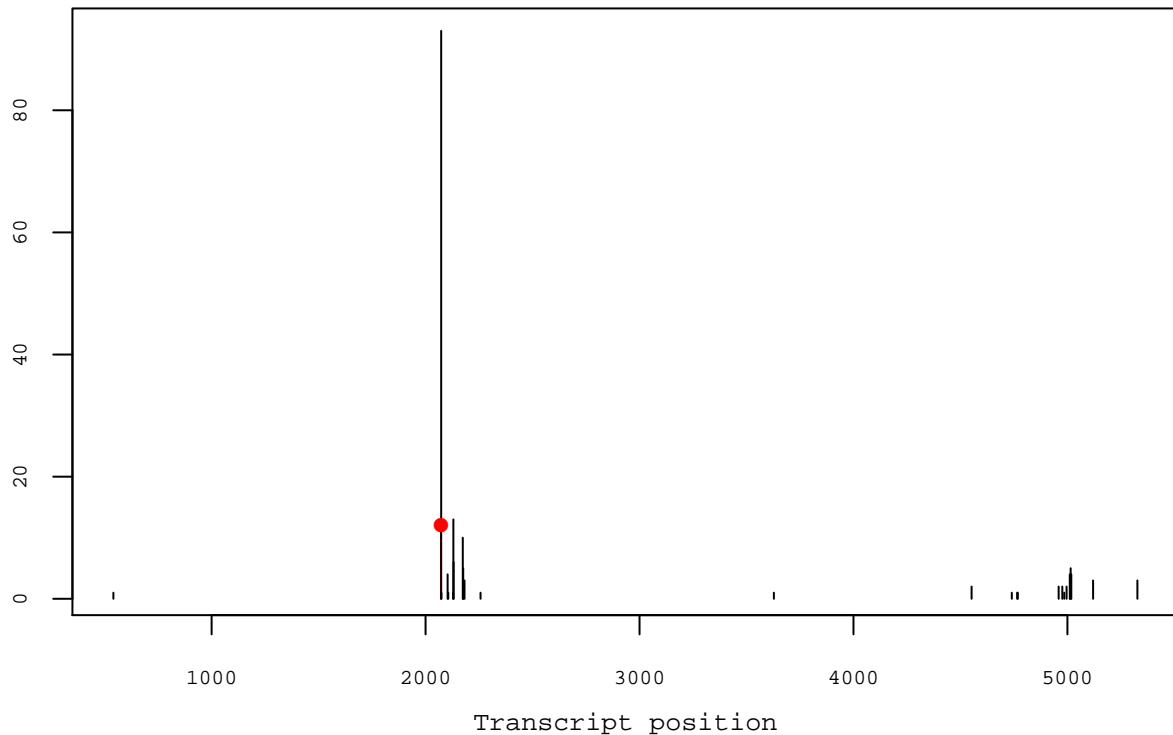


5' ACGA-ACAAAAAGTCCTACATTGGCTAGGTGTA '3  
| | | | | | | | | | | | | | | |  
3' TCTGTTTATCAGGGTGTAGCCGAT '5

Fragment Abundance



Cleavage site: 2072

Tag abundance: 12

Weighted abundance: 0.238

Category: 2

sRNA abundance: 619

Alignment score: 4

MFE ratio: 0.709

p-value: 0.026