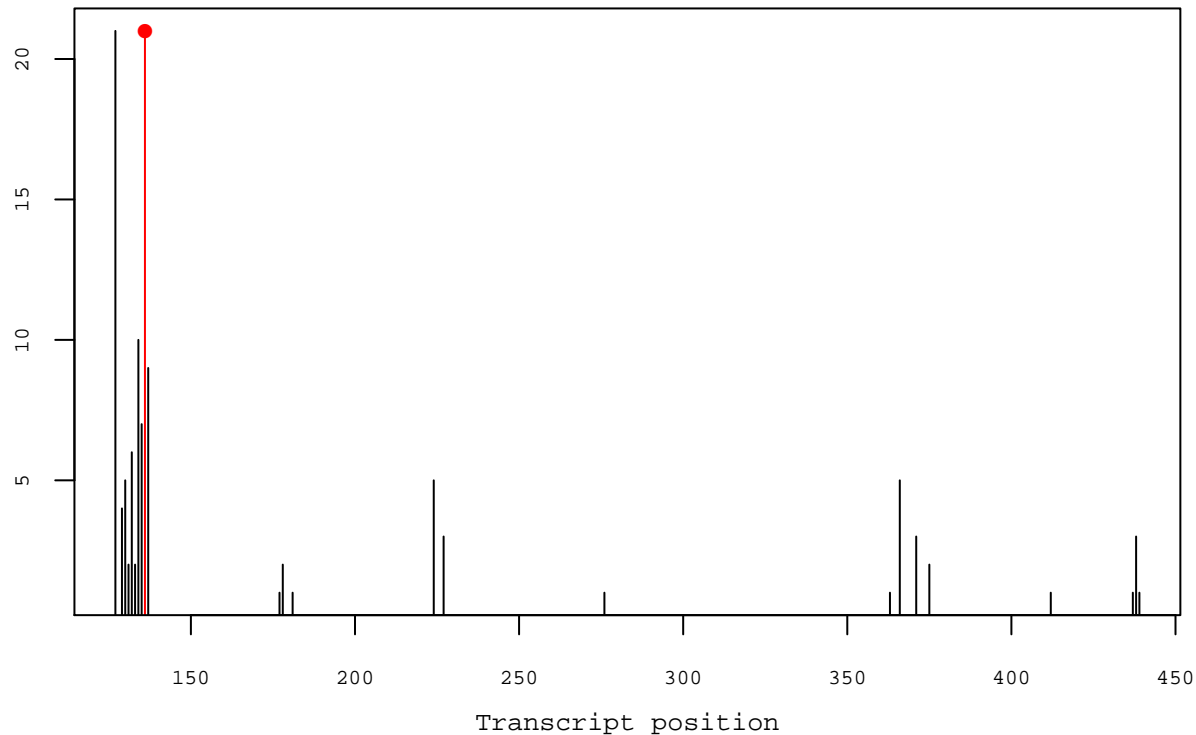


5' CGAACAC-AGCTGTTACGAACAAAGAGTTGTT '3  
||||| |||||  
3' TGTGCTCGACAAGTGCTTGTATTAT '5



|                    |                    |                          |                |
|--------------------|--------------------|--------------------------|----------------|
| Cleavage site: 136 | Tag abundance: 21  | Weighted abundance: 5.25 | Category: 1    |
| sRNA abundance: 23 | Alignment score: 3 | MFE ratio: 0.81          | p-value: 0.029 |