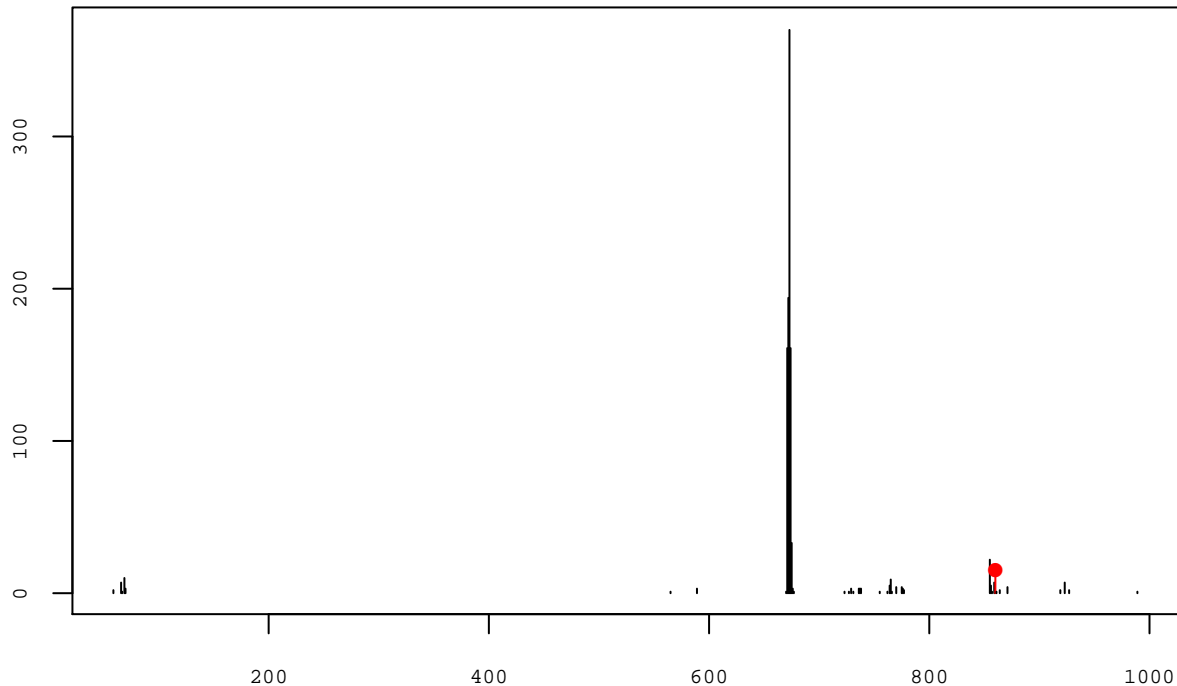


5' CTTTCTTAACTTAAGTC-GAATCTAAAACATCT '3
 ||||| o|||o||
3' GAATTGAATTCAGATTAGGTT '5

Fragment Abundance



Transcript position

Cleavage site: 860 Tag abundance: 15 Weighted abundance: 0.05 Category: 2
sRNA abundance: 26 Alignment score: 4 MFE ratio: 0.748 p-value: 0.045