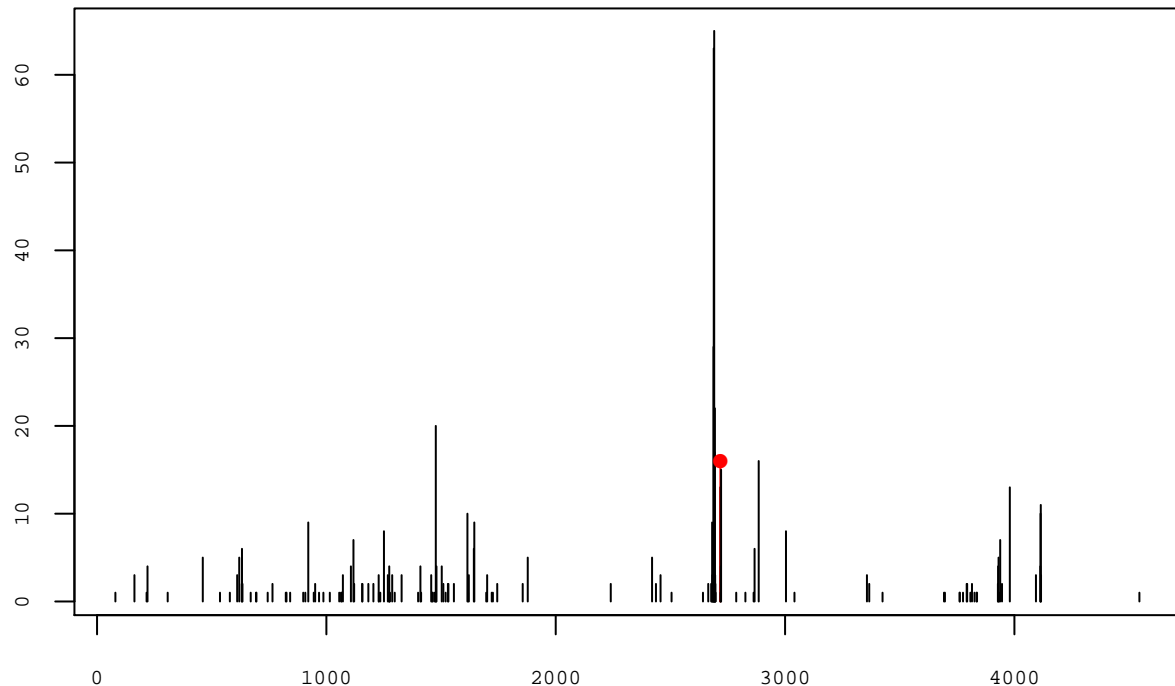


5' TAAATACATAGGAAGTGAGTGGTATGTATGTA '3
||||| | | | | | | | | | | | | | | |
3' TTATGTACCCTTTACTCACCATAC '5

Fragment Abundance



Transcript position

Cleavage site: 2718 Tag abundance: 16 Weighted abundance: 0.096 Category: 2
sRNA abundance: 28 Alignment score: 2 MFE ratio: 0.831 p-value: 0.022