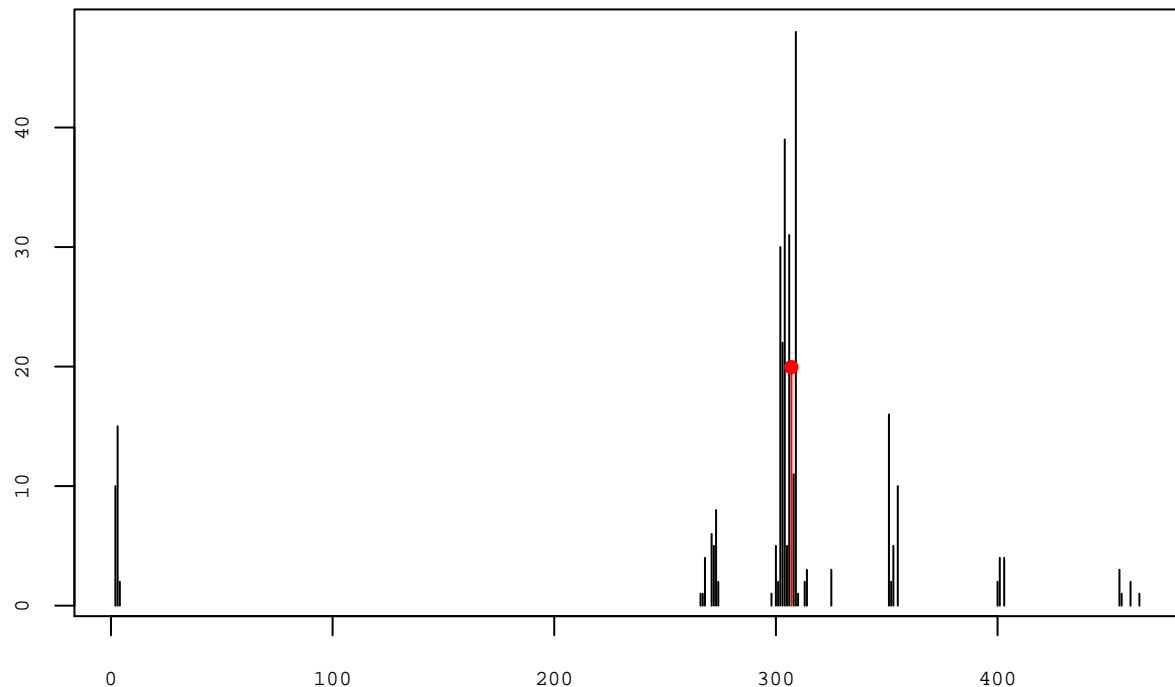


PggD11874_c54_g1_i1

5' CAATTCCTATCTATTTACATATCTAACACCTA '3
 ||||| ||||| |||
3' TAAGGATAGTTAAATGTATAGGT-G '5

Fragment Abundance



Transcript position

Cleavage site: 307 Tag abundance: 20 Weighted abundance: 0.059 Category: 2
sRNA abundance: 11 Alignment score: 3 MFE ratio: 0.76 p-value: 0.036