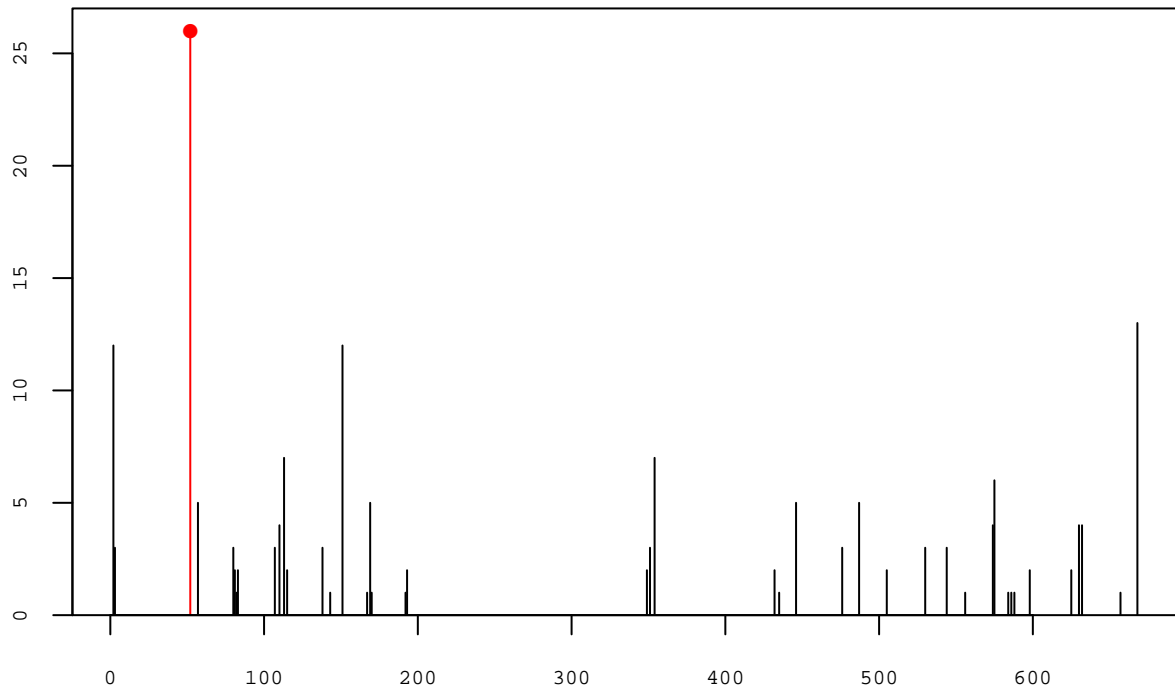


PggD7341\_c83\_g1\_i2

5' CCTATAATCTGTTTGCAGGTTACAGACGTACG '3  
||o |||||  
3' TTGTACAAACGTCCAATGTCT '5

Fragment Abundance



Transcript position

Cleavage site: 52 Tag abundance: 26 Weighted abundance: 1.083 Category: 0  
sRNA abundance: 10 Alignment score: 1.5 MFE ratio: 0.836 p-value: 0.018