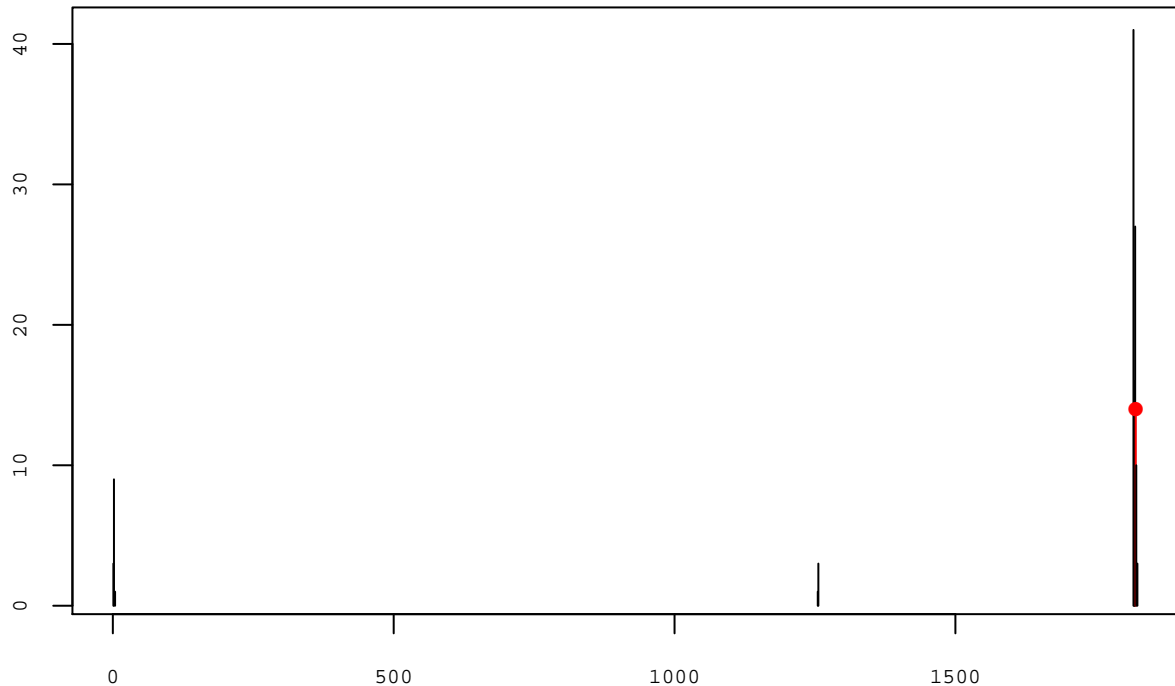


PggD12227\_c59\_g1\_i1

5' ATACCATCCAAATCCTATGTATTTACATATCT '3  
|||o||||||||||||||o|||  
3' TGGTGGGTTTAGGATACATAGATG '5

Fragment Abundance



Transcript position

Cleavage site: 1821 Tag abundance: 14 Weighted abundance: 0.206 Category: 3  
sRNA abundance: 8 Alignment score: 1.5 MFE ratio: 0.875 p-value: 0.004