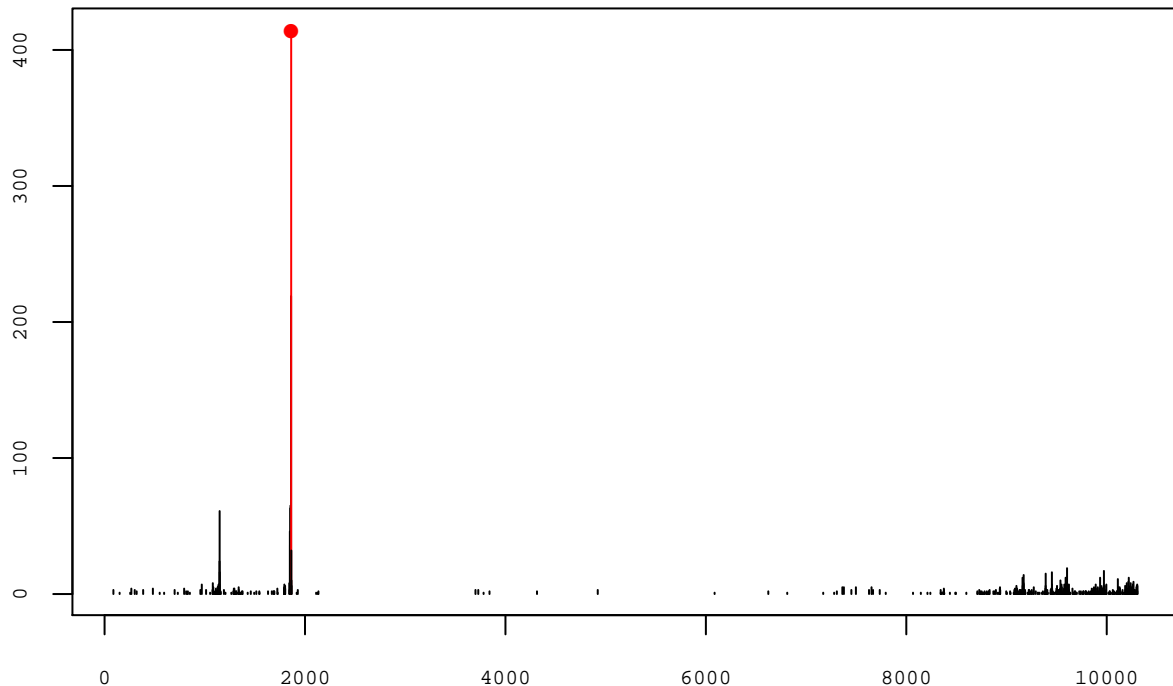


PggD51\_c2\_g1\_i7

5' AAATTT-AGGTGTTAGATATGTAAATACATAAG '3  
||| ||||| |o| ||||| |||  
3' AAAGTCCACAATTATACATTAT '5

Fragment Abundance



Transcript position

Cleavage site: 1862	Tag abundance: 414	Weighted abundance: 0.653	Category: 0
sRNA abundance: 5	Alignment score: 2	MFE ratio: 0.82	p-value: 0.03