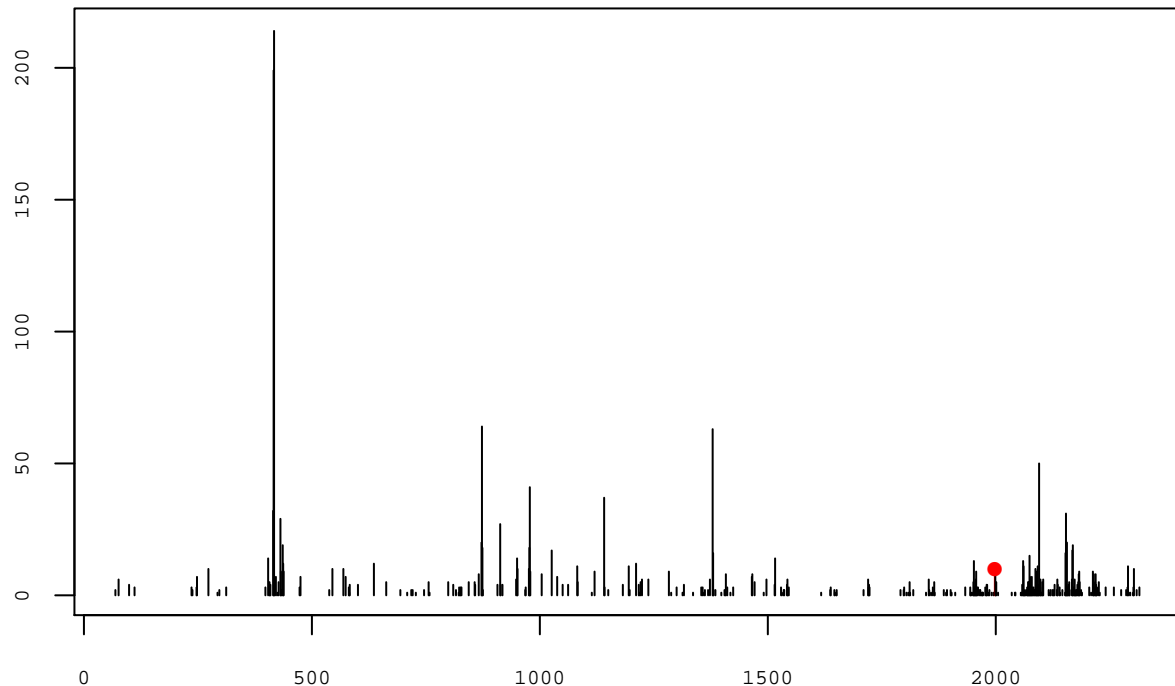


PggD14850\_c0\_g1\_i1

5' CCTACG-CCGCACATAGAGCTATGTCAGGAGTC '3  
| | | | | | | | | | | | | | | |  
3' TCAGGCGTGCATCTCGATGCAGT '5

Fragment Abundance



Transcript position

Cleavage site: 1998 Tag abundance: 10 Weighted abundance: 10 Category: 2  
sRNA abundance: 9 Alignment score: 4 MFE ratio: 0.721 p-value: 0.044