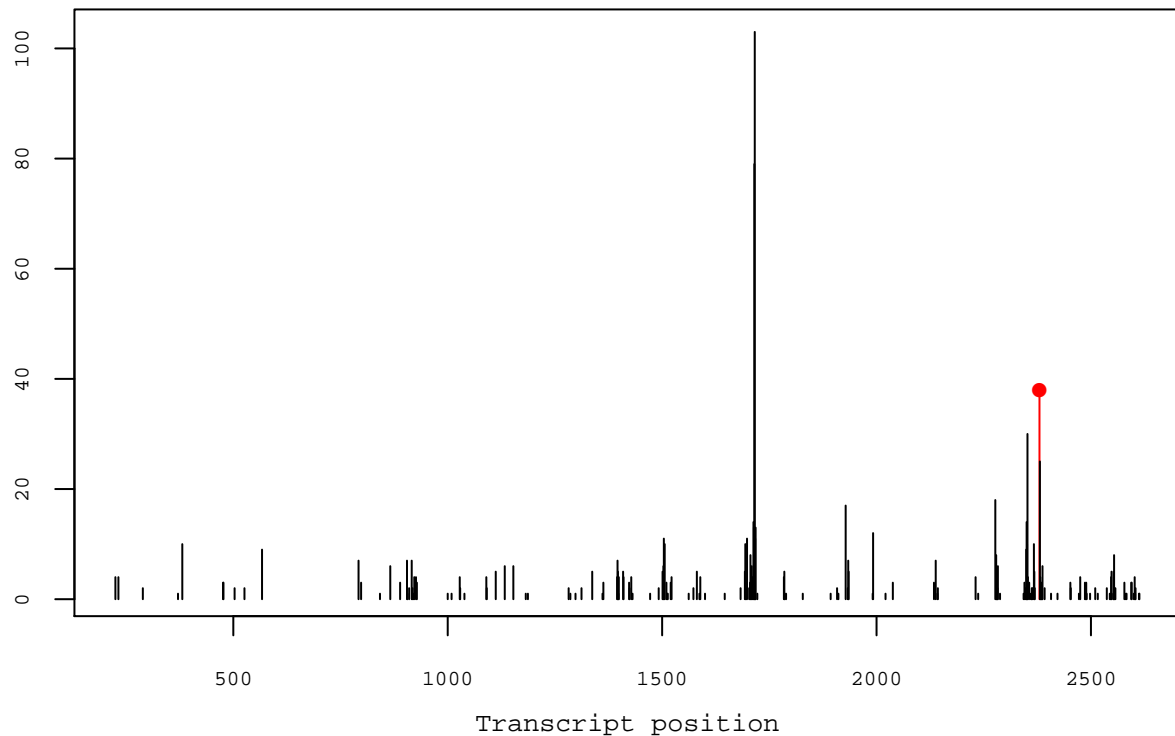


PggD14546\_c38\_g1\_i1

5' CAGTAGAGATCGGACGACGCATGTATCTCTGA '3  
|| ||||| ||||| o |||| o ||  
3' CACCTCTAGCCTGCTGTGTACGTA '5

Fragment Abundance



Cleavage site: 2380 Tag abundance: 38 Weighted abundance: 9.5 Category: 2  
sRNA abundance: 6 Alignment score: 3 MFE ratio: 0.808 p-value: 0.035