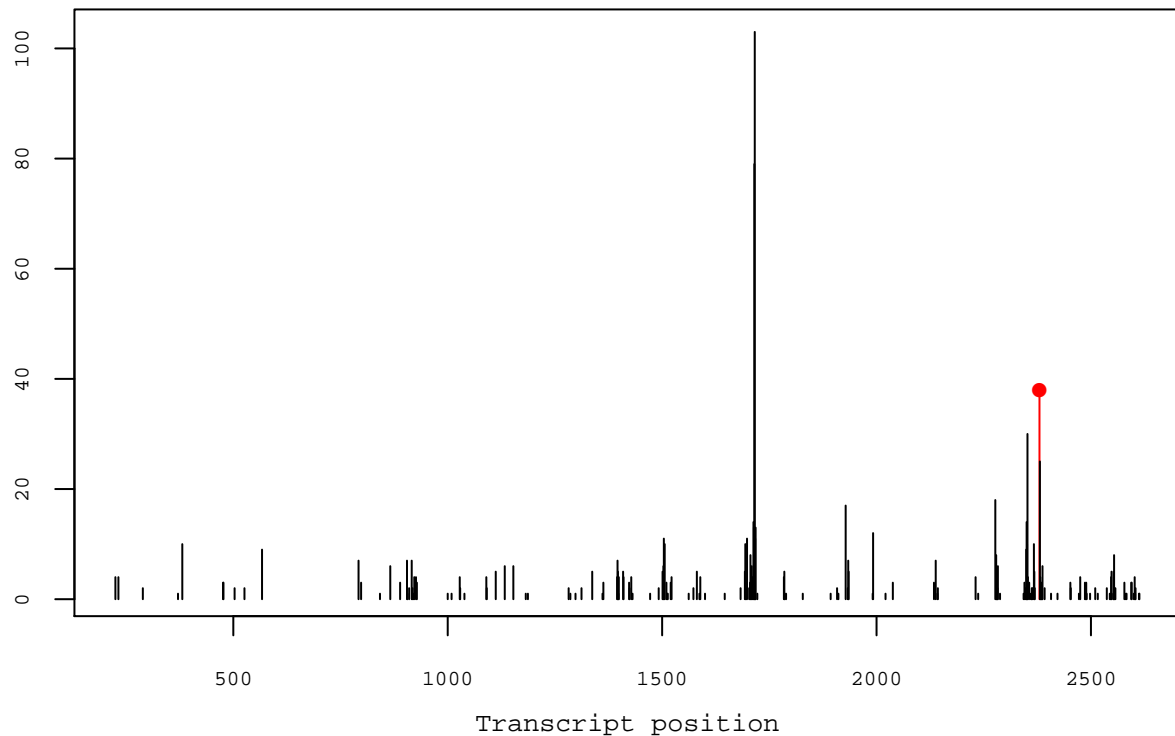


PggD14546\_c38\_g1\_i1

5' CAGTAGAGATCGGACGACGCATGTATCTCTGA '3  
|||||  
3' CATCTCTAGCCTGCTGCGTACGTA '5

Fragment Abundance



Cleavage site: 2380 Tag abundance: 38 Weighted abundance: 9.5 Category: 2  
sRNA abundance: 7 Alignment score: 1 MFE ratio: 0.957 p-value: 0.035