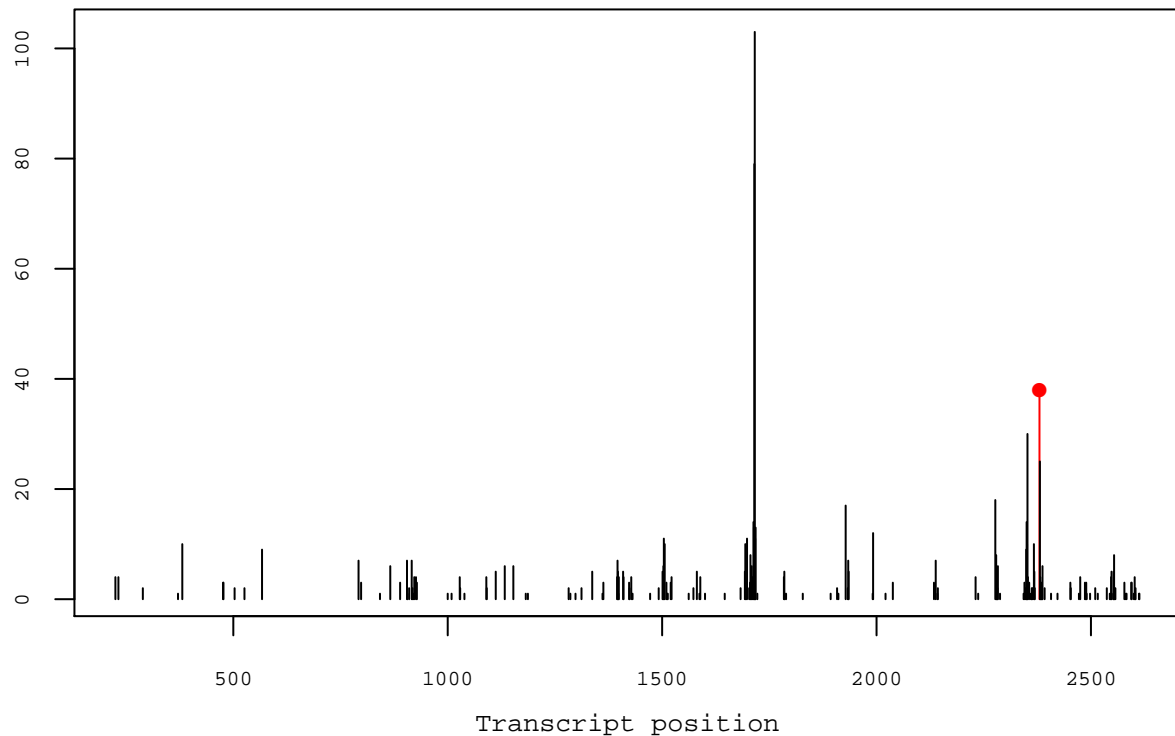


PggD14546_c38_g1_i1

5' CAGTAGAGATCGGACGACGCATGTATCTCTGA '3
|||||o |||||o ||
3' CATCTCTAGCCTGTGGCGTACGTA '5

Fragment Abundance



Cleavage site: 2380 Tag abundance: 38 Weighted abundance: 9.5 Category: 2
sRNA abundance: 62 Alignment score: 4 MFE ratio: 0.802 p-value: 0.043