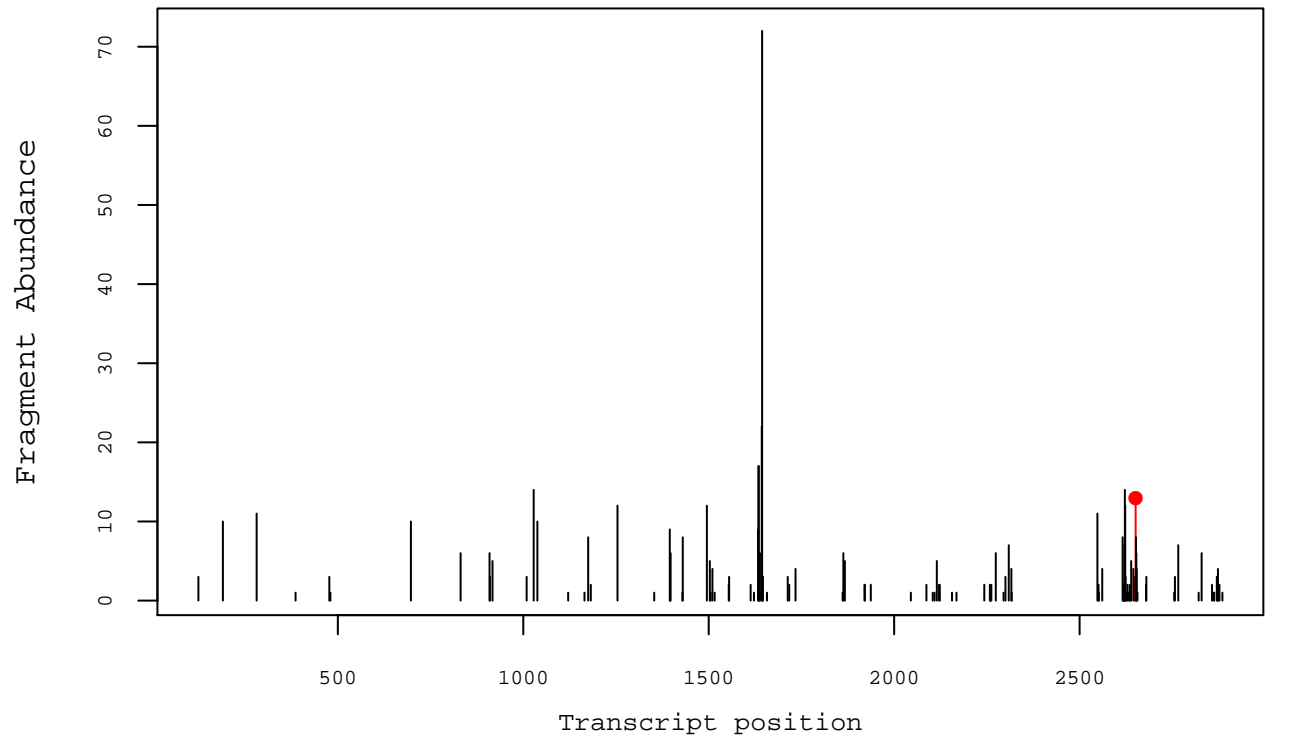


PggD14546_c38_g1_i6

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



Cleavage site: 2651 Tag abundance: 13 Weighted abundance: 3.25 Category: 2
sRNA abundance: 18 Alignment score: 4 MFE ratio: 0.802 p-value: 0.047