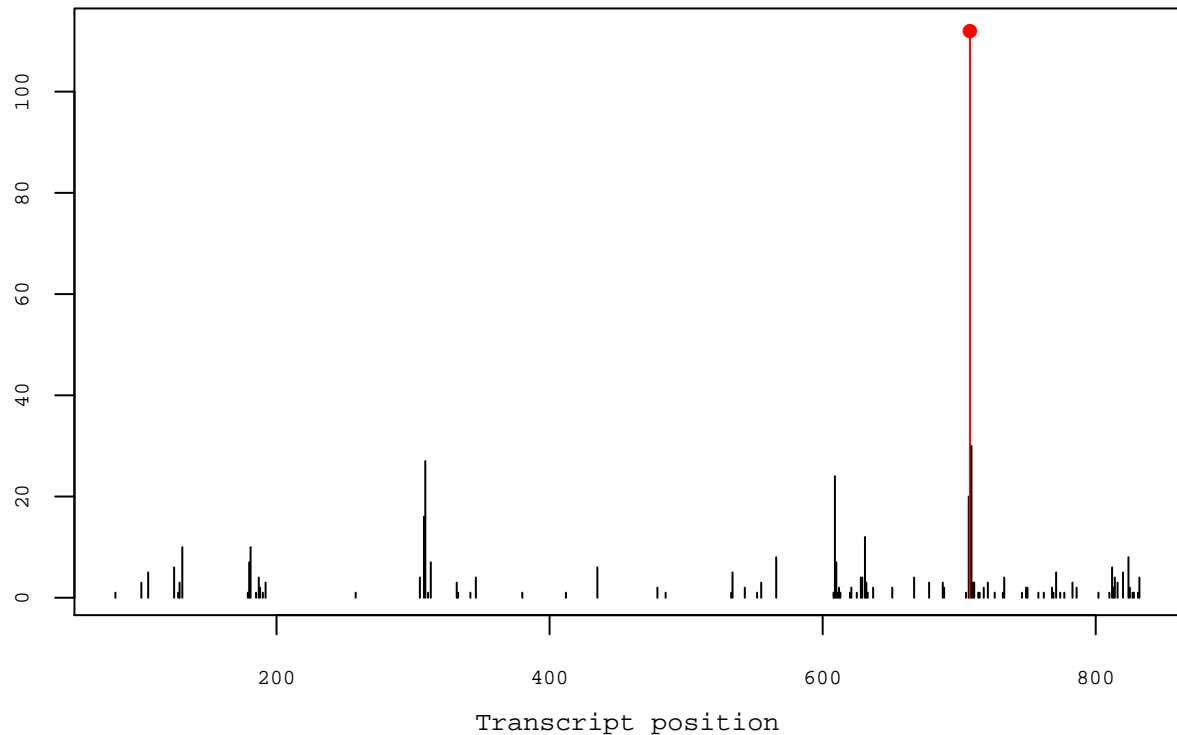


PggD3526_c13_g1_i5

5' AAAGTGCCTTTGGGTTTTTAACTAAACGTAG '3
||o| |||||o|||||||
3' TGGCACAAACTCAAAAATGTGATT '5

Fragment Abundance



Cleavage site: 708 Tag abundance: 112 Weighted abundance: 28 Category: 0
sRNA abundance: 6 Alignment score: 2 MFE ratio: 0.804 p-value: 0.013