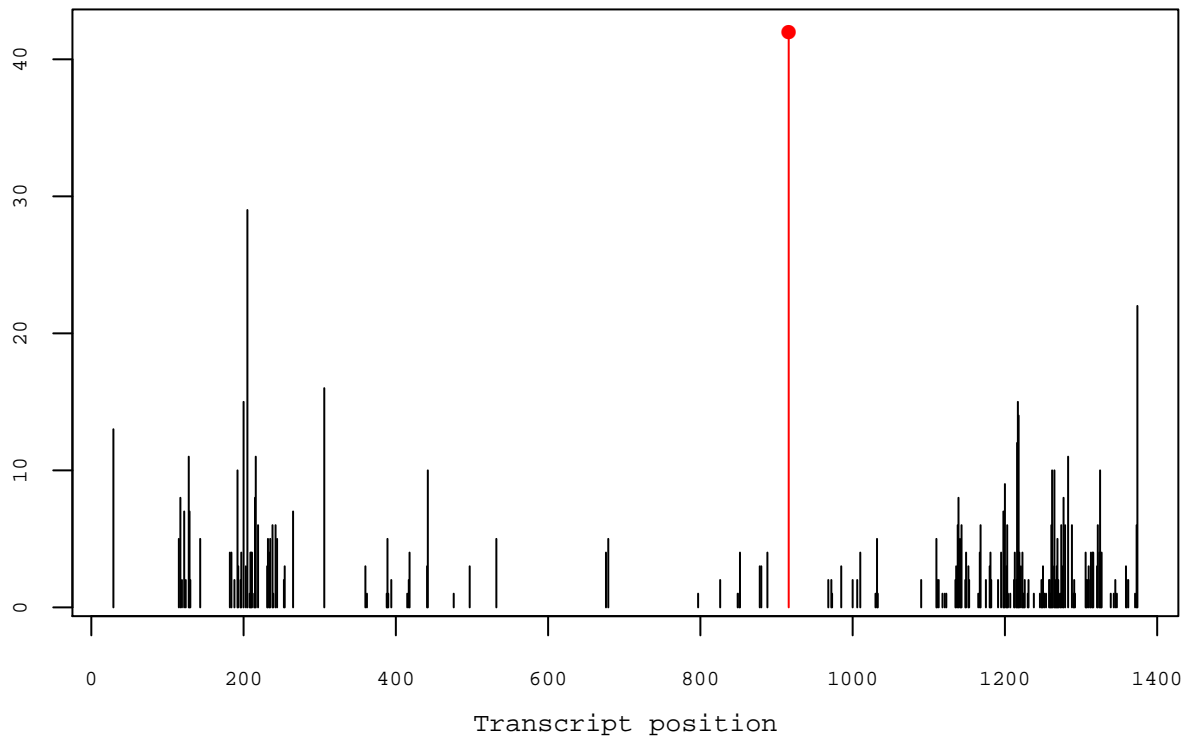


PggD1526_c1_g1_i2

5' CCCAACTCACGTGCCCTGCTTCTCCAATGGTA '3
|||||
3' CGTGCACGGGACGAAGAGGT '5

Fragment Abundance



Cleavage site: 916 Tag abundance: 42 Weighted abundance: 42 Category: 0
sRNA abundance: 45 Alignment score: 1 MFE ratio: 0.932 p-value: 0.004