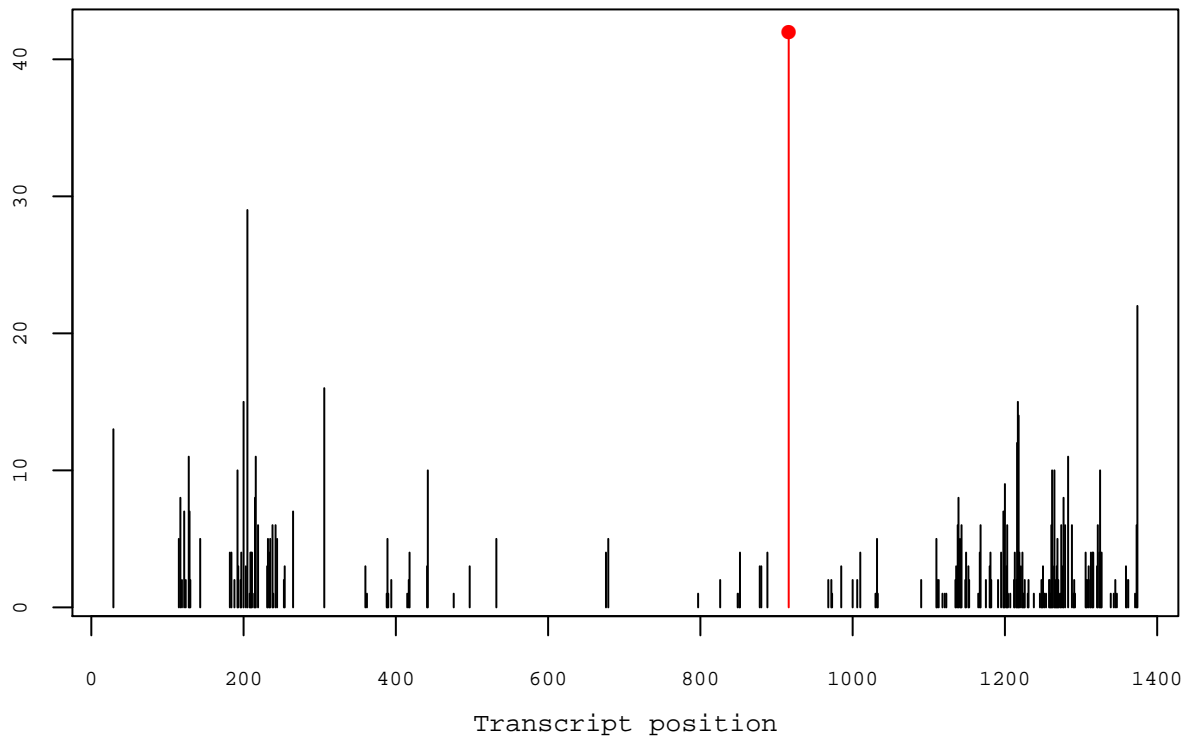


PggD1526\_c1\_g1\_i2

5' CCCAACTCACGTGCCCTGCTTCTCCAATGGTA '3  
|||||  
3' ACGTGCACGGGACGAAGAGGT '5

Fragment Abundance



Cleavage site: 916 Tag abundance: 42 Weighted abundance: 42 Category: 0  
sRNA abundance: 235 Alignment score: 2 MFE ratio: 0.898 p-value: 0.004