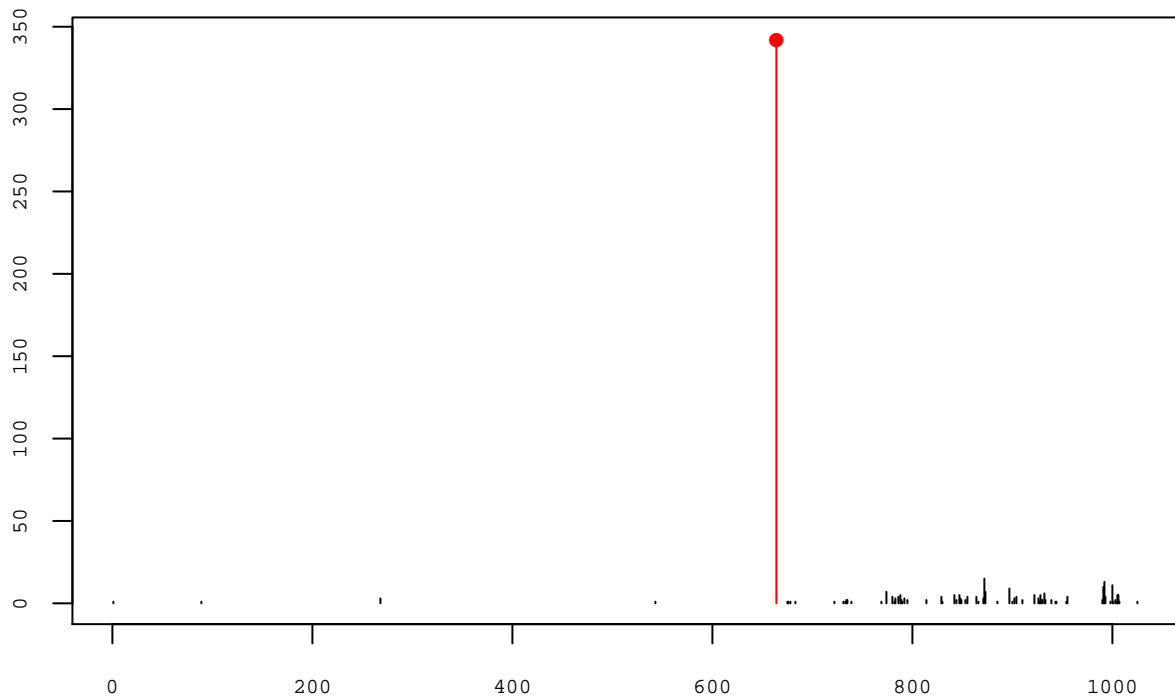


PggD1457_c16_g2_i2

5' GCTTGAGCAGGTGCCCTGCTTCTCCAGCTTTT '3
||| ||||| ||||| ||||| |||||
3' ACGTGCACGGGACGAAGAGGT '5

Fragment Abundance



Transcript position

Cleavage site: 664 Tag abundance: 342 Weighted abundance: 171 Category: 0
sRNA abundance: 412 Alignment score: 2 MFE ratio: 0.926 p-value: 0.005