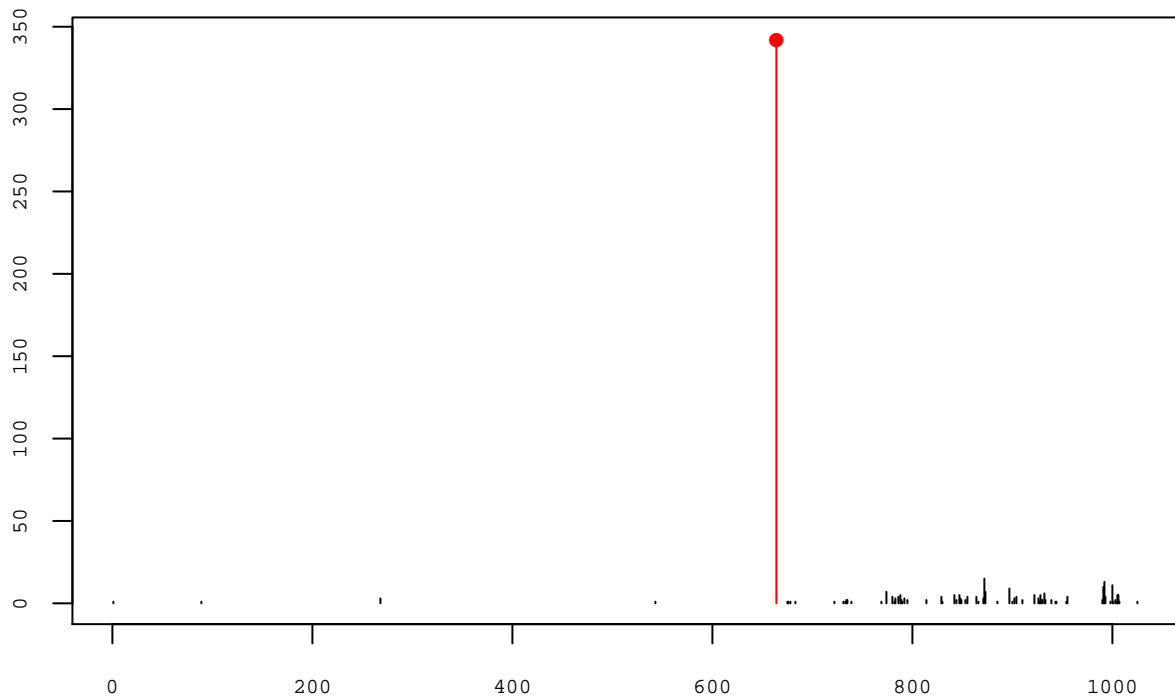


PggD1457_c16_g2_i2

5' GCTTGAGCAGGTGCCCTGCTTCTCCAGCTTTT 3'
|| |||||
3' GTGCACGGGACGAAGAGGT 5'

Fragment Abundance



Transcript position

Cleavage site: 664 Tag abundance: 342 Weighted abundance: 171 Category: 0
sRNA abundance: 17 Alignment score: 1 MFE ratio: 0.919 p-value: 0.007