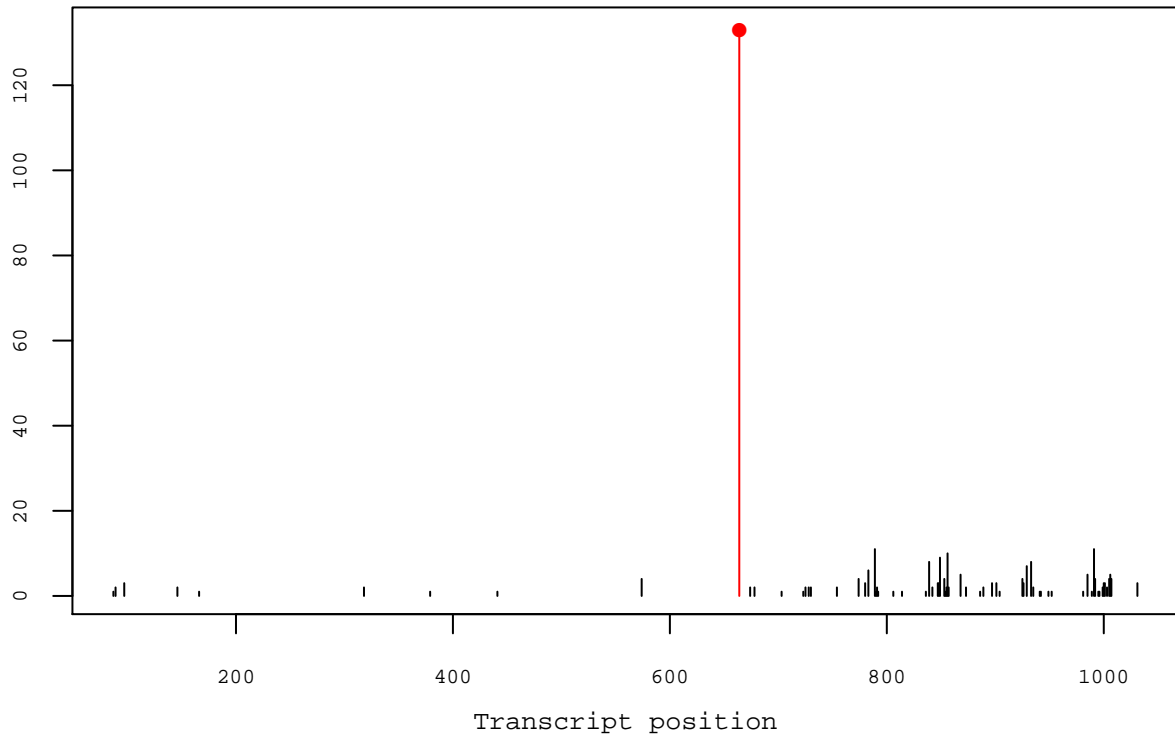


PggD1457_c16_g2_i2

5' GCTTGAGCAGGTGCCCTGCTTCTCCAGCTTTT '3
| o | | | | | | | | | | | | | | | |
3' TTGTGCACGGGACGAAGAGGT '5

Fragment Abundance



Cleavage site: 664	Tag abundance: 133	Weighted abundance: 66.5	Category: 0
sRNA abundance: 13	Alignment score: 1.5	MFE ratio: 0.926	p-value: 0.006