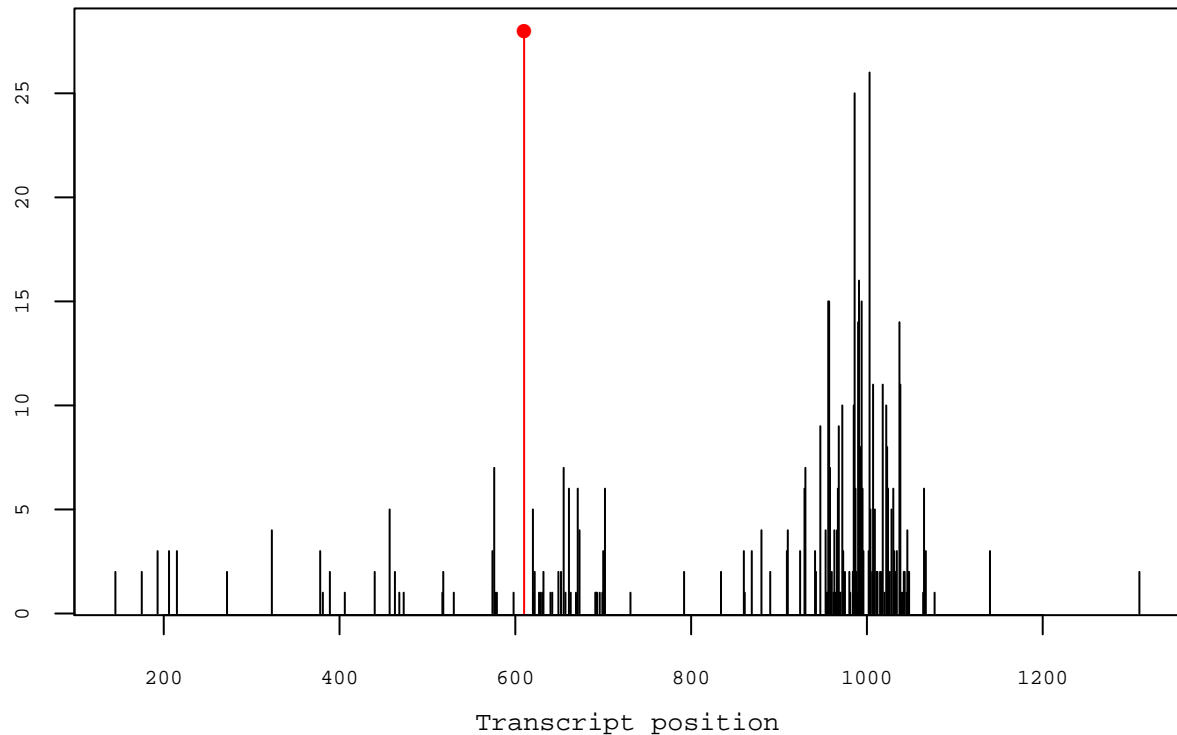


PggD13945_c35_g1_i1

5' AAGGAACTGCGCGGTCTTGAGCAAAACATTGA '3
|||||
3' CAACGCGCCAGAACTCGTTTT '5

Fragment Abundance



Cleavage site: 610 Tag abundance: 28 Weighted abundance: 7 Category: 0
sRNA abundance: 8 Alignment score: 2 MFE ratio: 0.932 p-value: 0.007