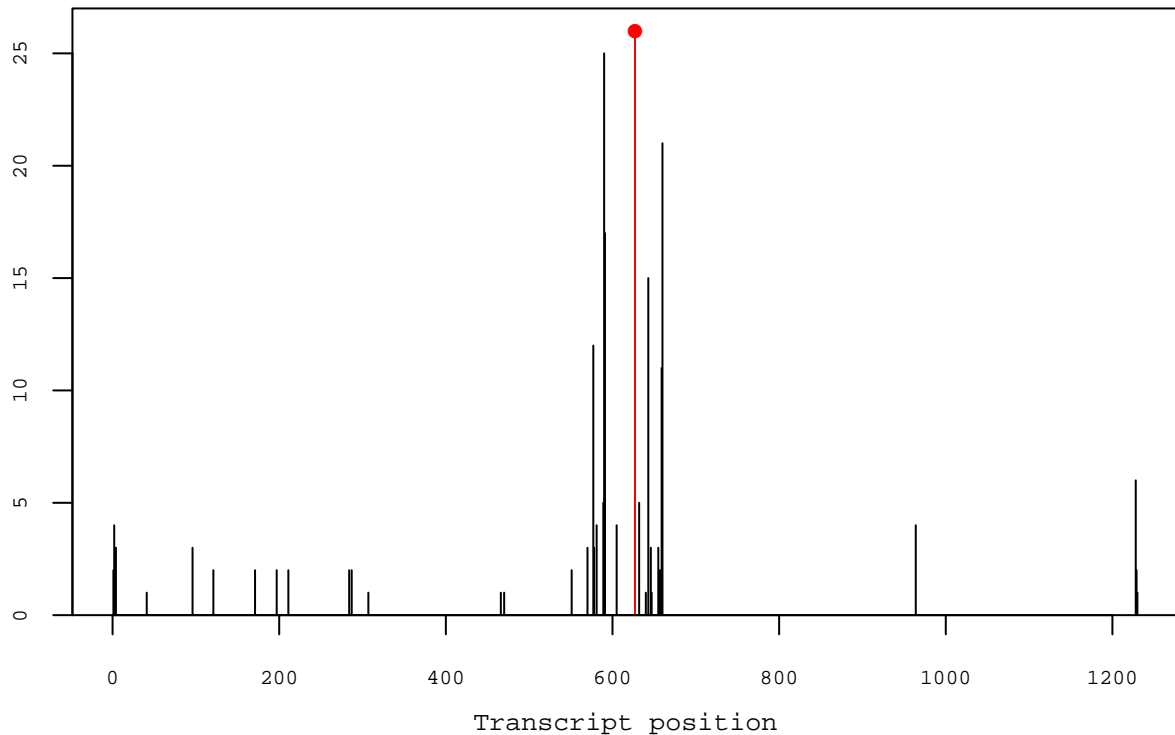


PggD13189_c49_g1_i2

5' TCTGTAAC TTGTTTGCAGGTTACAGACGTACG '3
||| ||||| ||||| ||||| |||||
3' TTGTACAAACGTCCAATGTCT '5

Fragment Abundance



Cleavage site: 627
sRNA abundance: 10

Tag abundance: 26
Alignment score: 1

Weighted abundance: 1.083
MFE ratio: 0.881

Category: 0
p-value: 0.009