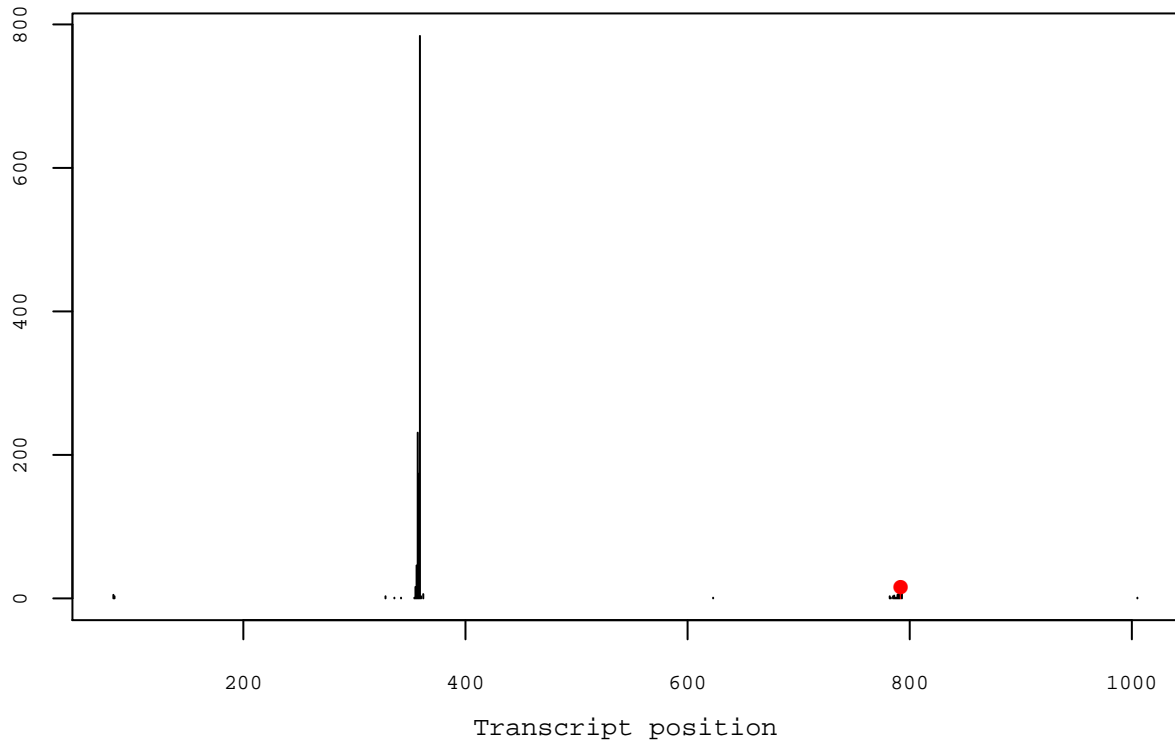


PggD72_c23_g1_i1

5' TTGTTTCGCGAACAACCTCGTGTTCGGCTTGTTA '3
|o||| | |||||
3' CGAGCACAAAGTTGAGCACAAAGCCG '5



Cleavage site: 792 Tag abundance: 16 Weighted abundance: 6.333 Category: 2
sRNA abundance: 20 Alignment score: 3.5 MFE ratio: 0.717 p-value: 0.045