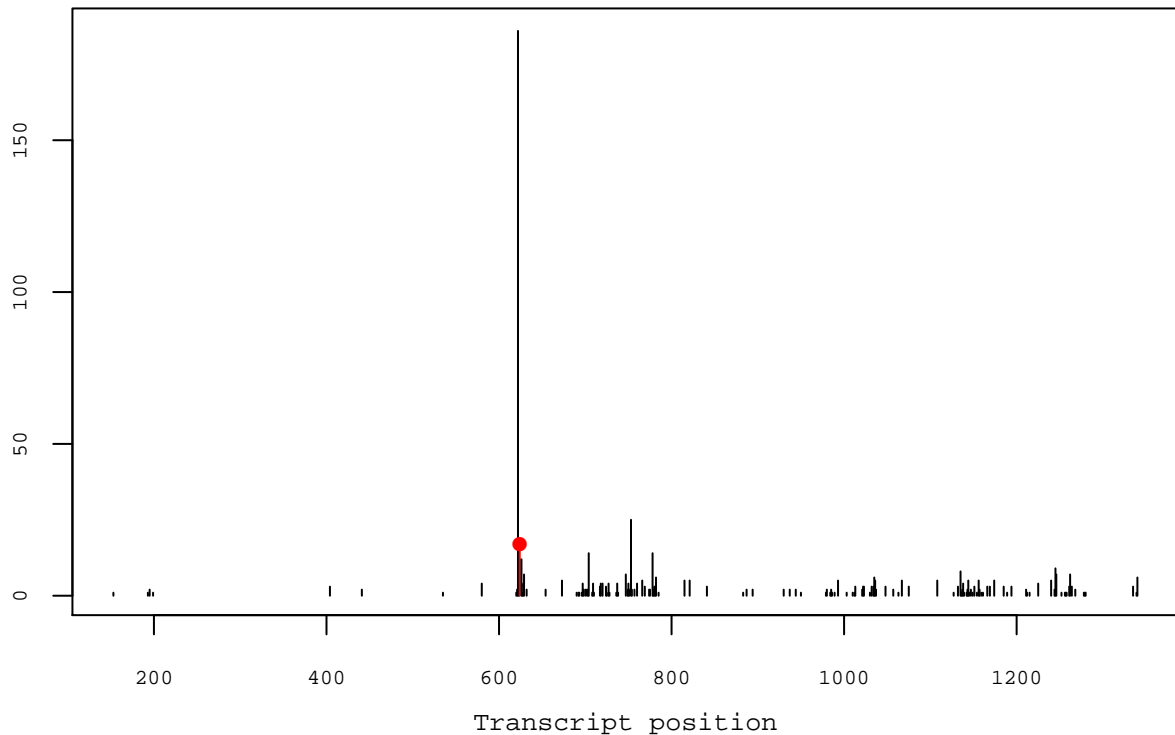


PggD13991_c0_g1_i1

5' GCTGATACATTGTGTGATATCTTAATTGGATG '3
|||||||oo|||||||
3' ATGTAAATGCTATAGAATTAA '5

Fragment Abundance



Cleavage site: 624 Tag abundance: 17 Weighted abundance: 8.5 Category: 2
sRNA abundance: 5 Alignment score: 1.5 MFE ratio: 0.806 p-value: 0.016