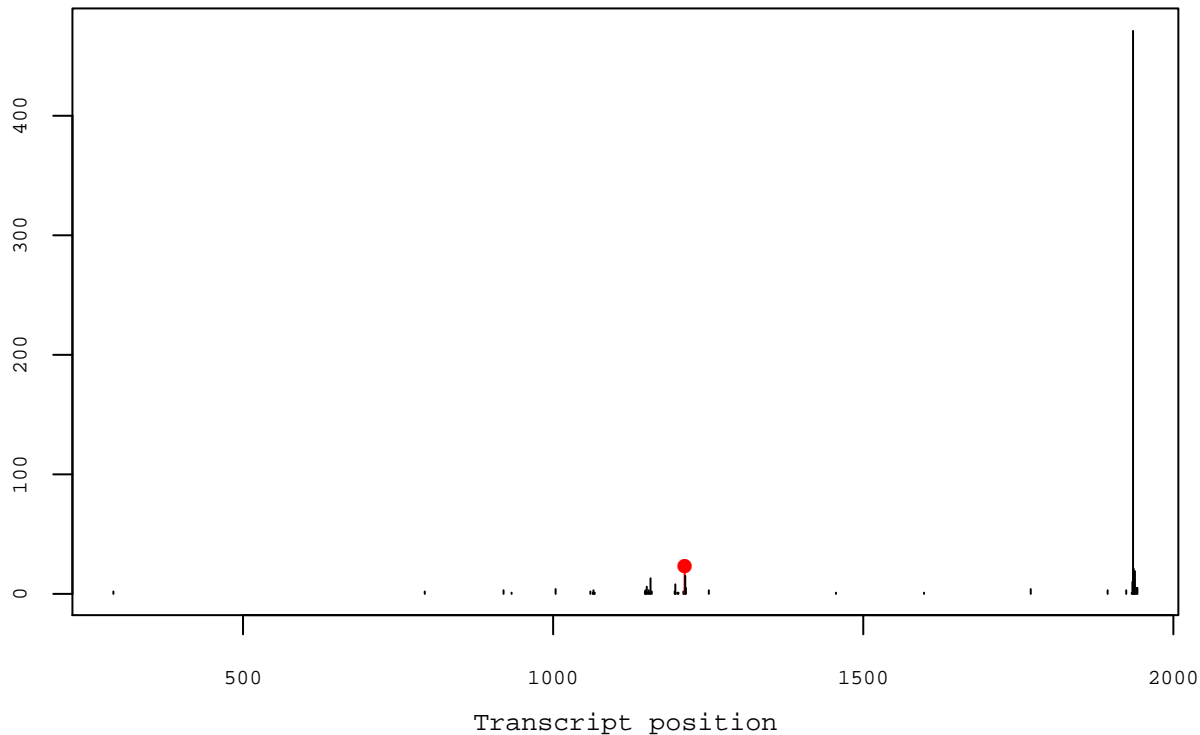


PggD15095_c245_g1_i1

5' CACGTAGAGCTACGTCAGGAGTCCGGACAGAA 3'
|||||
3' GCATCTCGATGCAGTCCCTCAGGCC 5'



Cleavage site: 1212 Tag abundance: 23 Weighted abundance: 11.5 Category: 3
sRNA abundance: 6 Alignment score: 0 MFE ratio: 1 p-value: 0.013