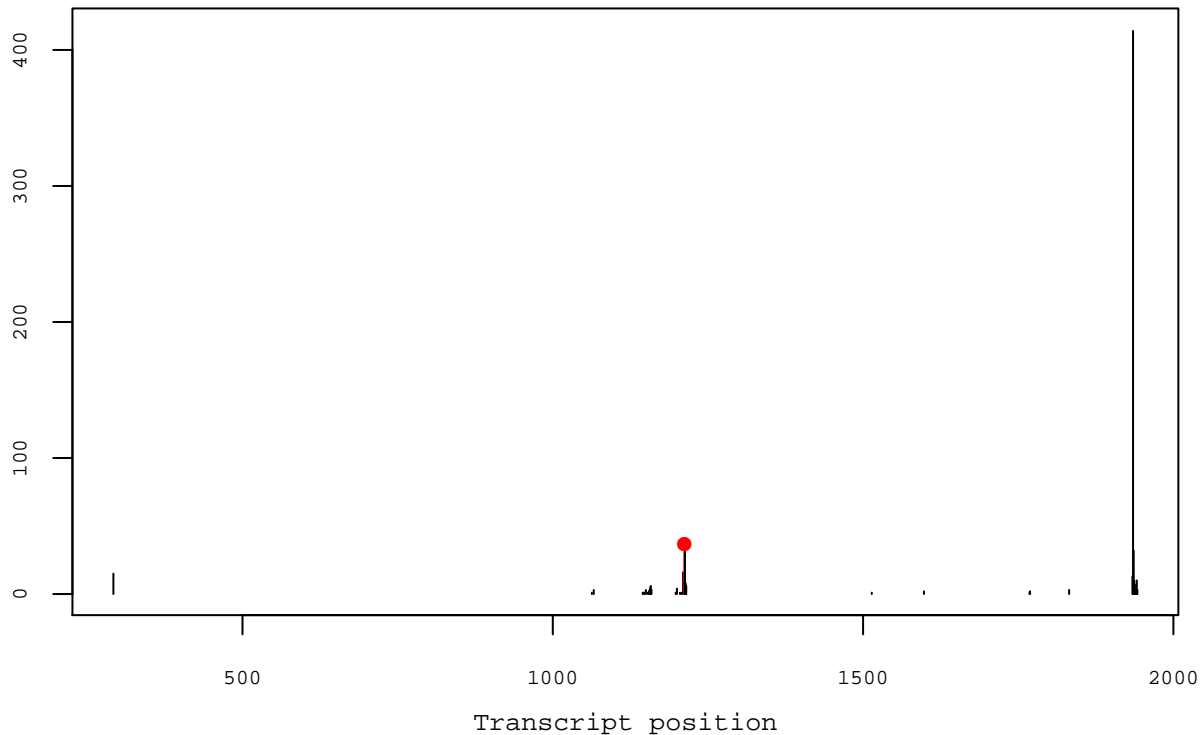


PggD15095\_c245\_g1\_i1

5' CACGTAGAGCTACGTCAGGAGTCCGGACAGAA 3'  
 |||||  
 3' GCATCTCGATGCAGTTCTCAGGCG 5'



Cleavage site: 1212    Tag abundance: 37    Weighted abundance: 18.333    Category: 2  
sRNA abundance: 13    Alignment score: 2    MFE ratio: 0.952    p-value: 0.001