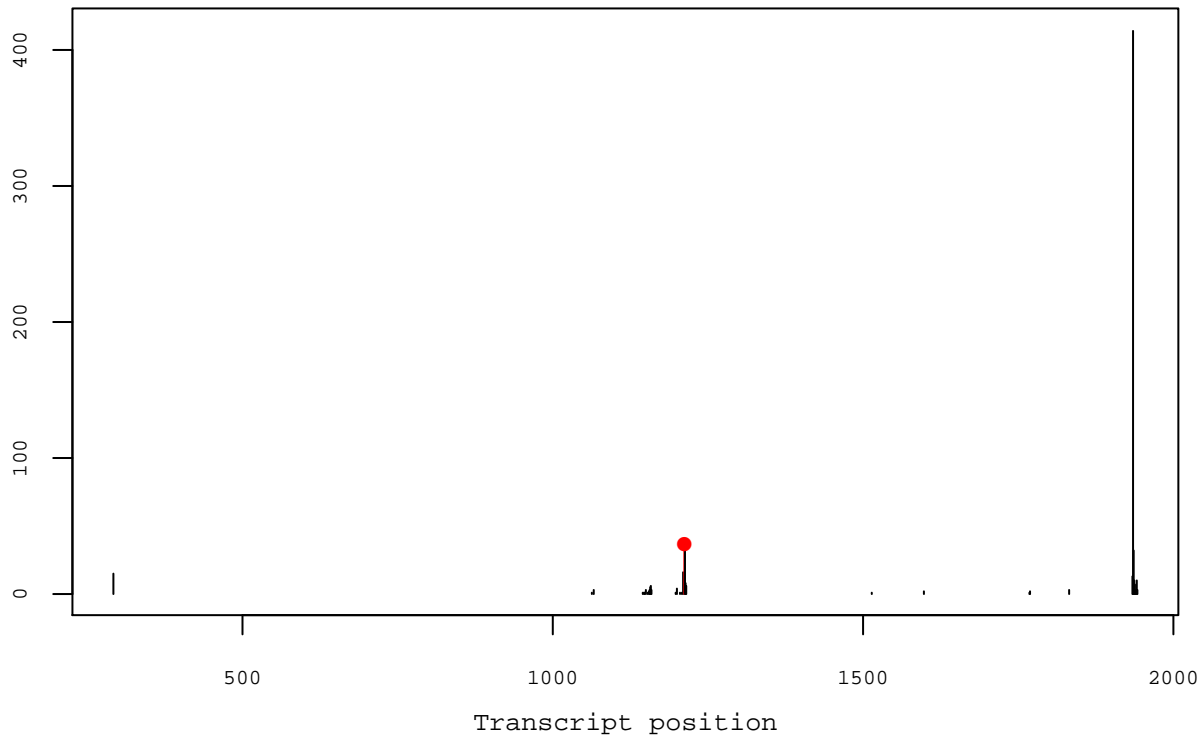


5' CACGTAGAGCTACGTCAGGAGTCCGGACAGAA 3'
|||||
3' GCATCTCGATGCAGTCCCTCAGGCC 5'



Cleavage site: 1212 Tag abundance: 37 Weighted abundance: 18.333 Category: 2
sRNA abundance: 64 Alignment score: 0 MFE ratio: 1 p-value: 0.001