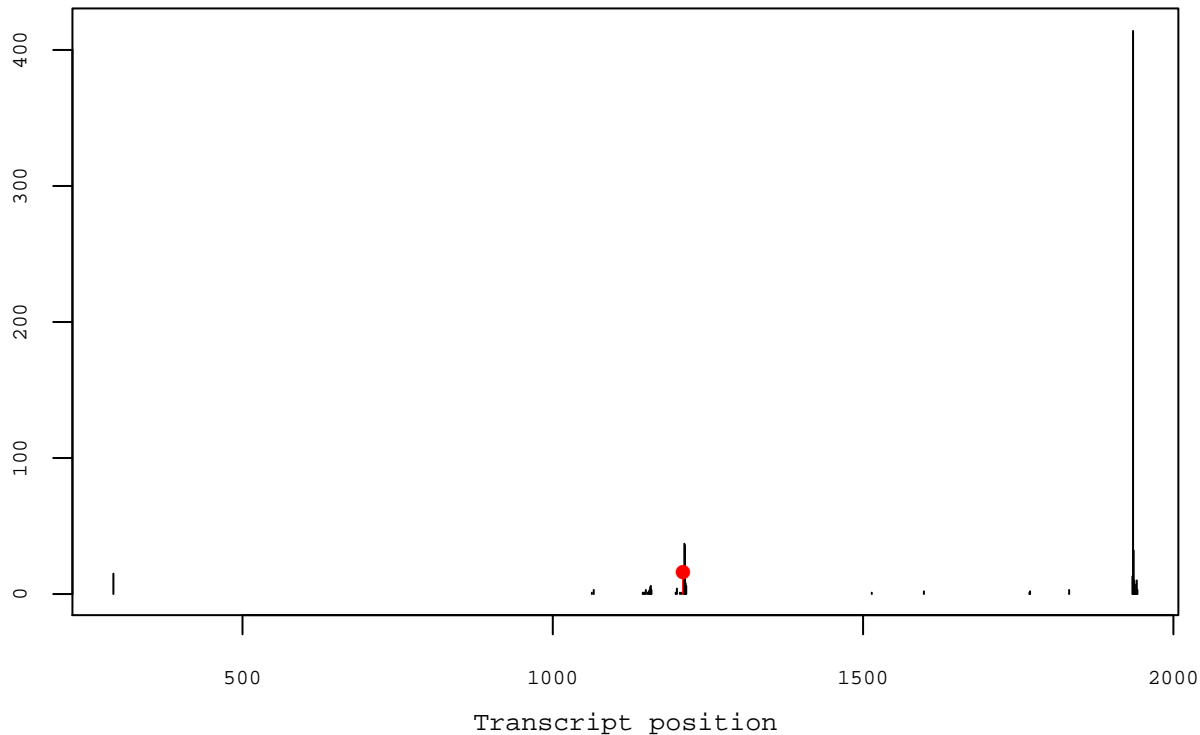


PggD15095\_c245\_g1\_i1

5' TGCACGTAGAGCTACGTCAGGAGTCCGGACAG '3  
|||||  
3' GTGCATCTCGATGCAGTCCTTAGG '5

Fragment Abundance



Cleavage site: 1210 Tag abundance: 16 Weighted abundance: 3.619 Category: 3  
sRNA abundance: 9 Alignment score: 1 MFE ratio: 1 p-value: 0.022