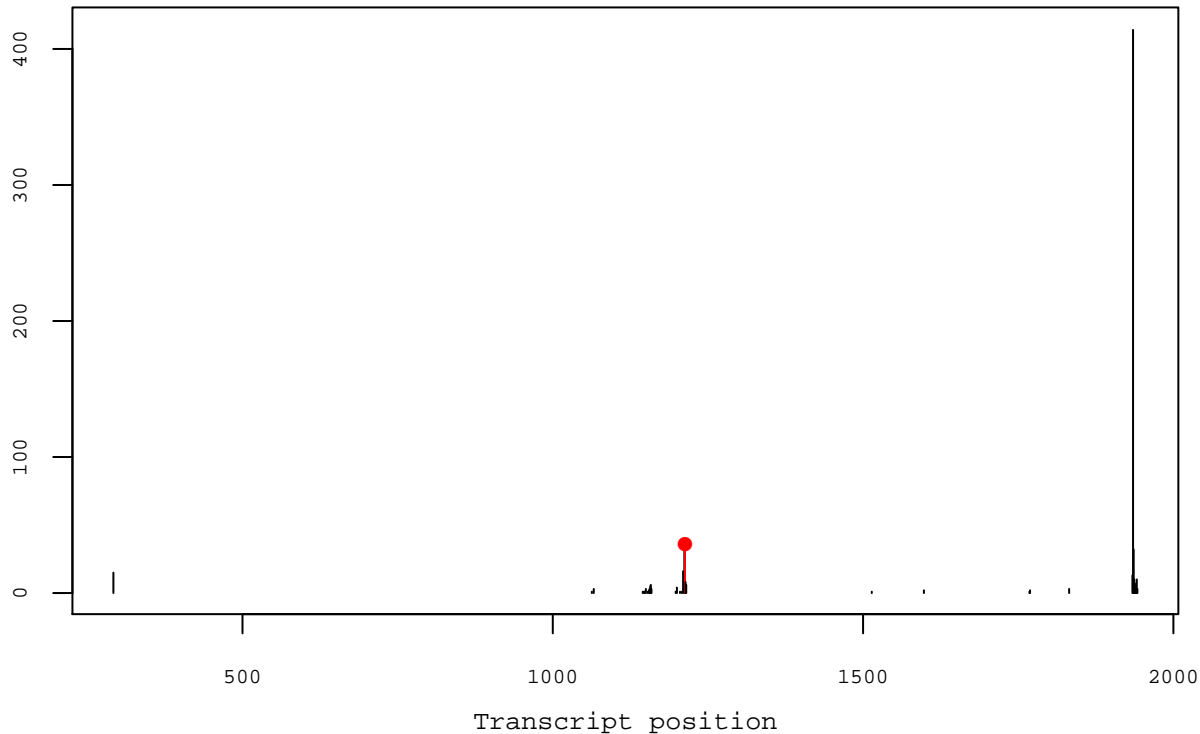


PggD15095_c245_g1_i1

5' ACGTAGAGCTACGTCAGGAGTCCG-GACAGAAA '3
|||||
3' CATCTCGATGCAGTCCTCAGACGT '5

Fragment Abundance



Cleavage site: 1213 Tag abundance: 36 Weighted abundance: 18 Category: 2
sRNA abundance: 13 Alignment score: 3.5 MFE ratio: 0.828 p-value: 0.001