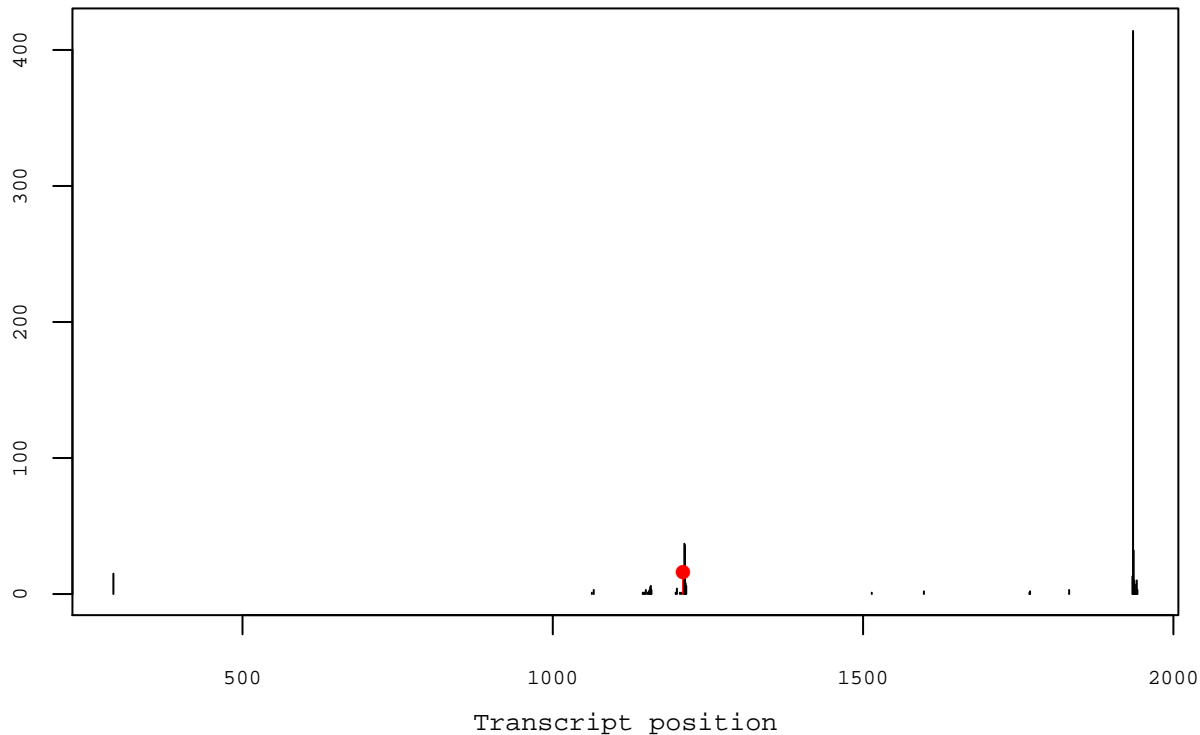


PggD15095_c245_g1_i1

5' TGCACGTAGAGCTACGTCAGGAGTCCGGACAG '3
|||||
3' GTGCATCTCGATGCAGTCCTAAGG '5

Fragment Abundance



Cleavage site: 1210 Tag abundance: 16 Weighted abundance: 3.619 Category: 3
sRNA abundance: 6 Alignment score: 2 MFE ratio: 0.916 p-value: 0.022