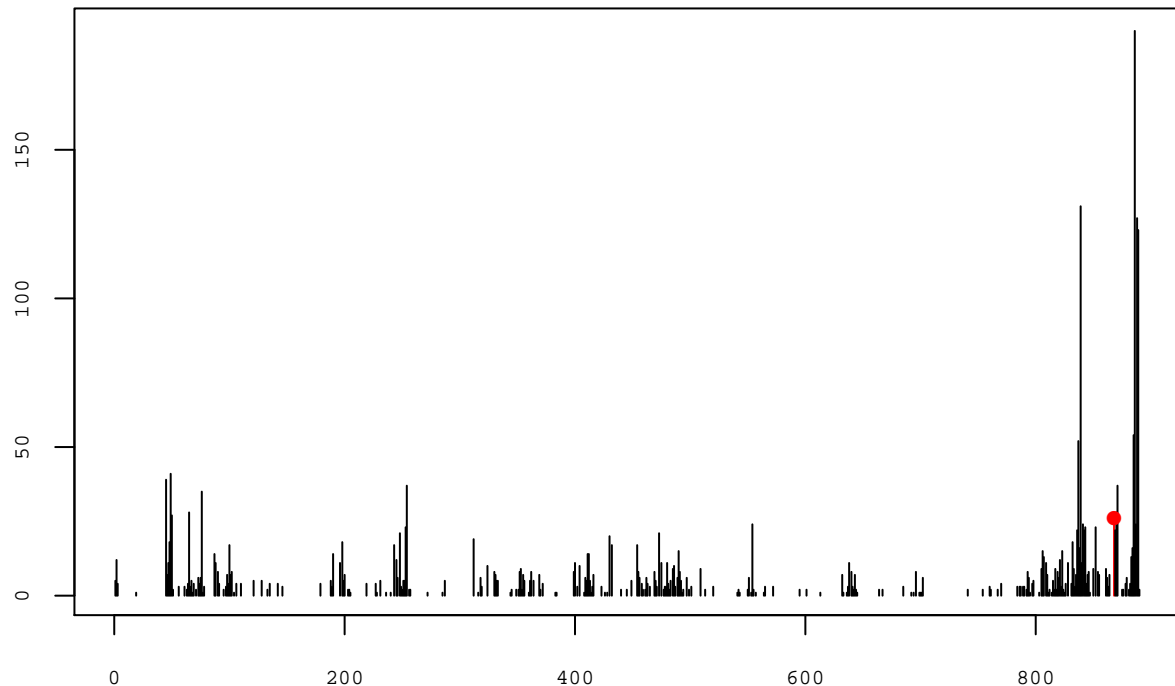


PggD15204_c33_g1_i1

5' CCTAGAAGGTCAATAACACCTGAATTCTGAAA '3
||| | o |||||
3' ATCTTGT-GTTATTGTGGACTTAT '5

Fragment Abundance



Transcript position

Cleavage site: 868 Tag abundance: 26 Weighted abundance: 26 Category: 2
sRNA abundance: 7 Alignment score: 3.5 MFE ratio: 0.762 p-value: 0.047