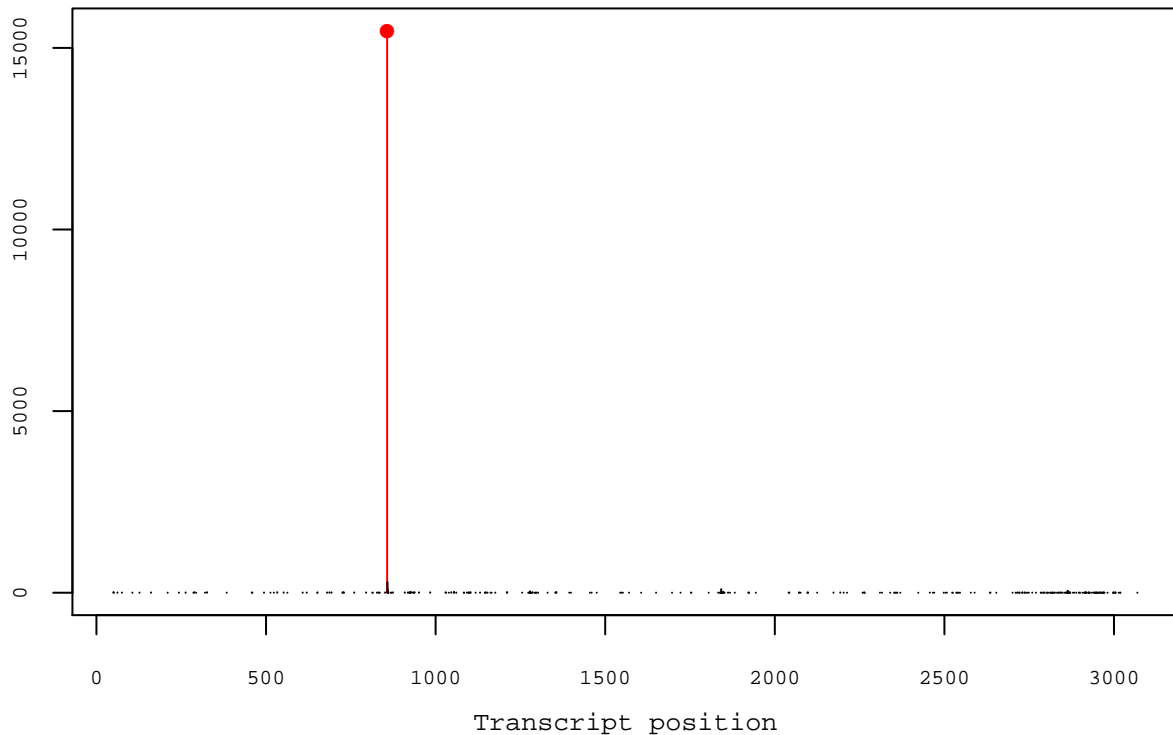


PggD11091_c20_g1_i1

5' GATGGTCGGGATGAAGCCTGGTCCGGATTCTA '3
|||o|||||||||||o
3' CCCCTTACTTCGGACCAGGCT '5

Fragment Abundance



Cleavage site: 857 Tag abundance: 15468 Weighted abundance: 7734 Category: 0
srRNA abundance: 19192 Alignment score: 3 MFE ratio: 0.855 p-value: 0.001