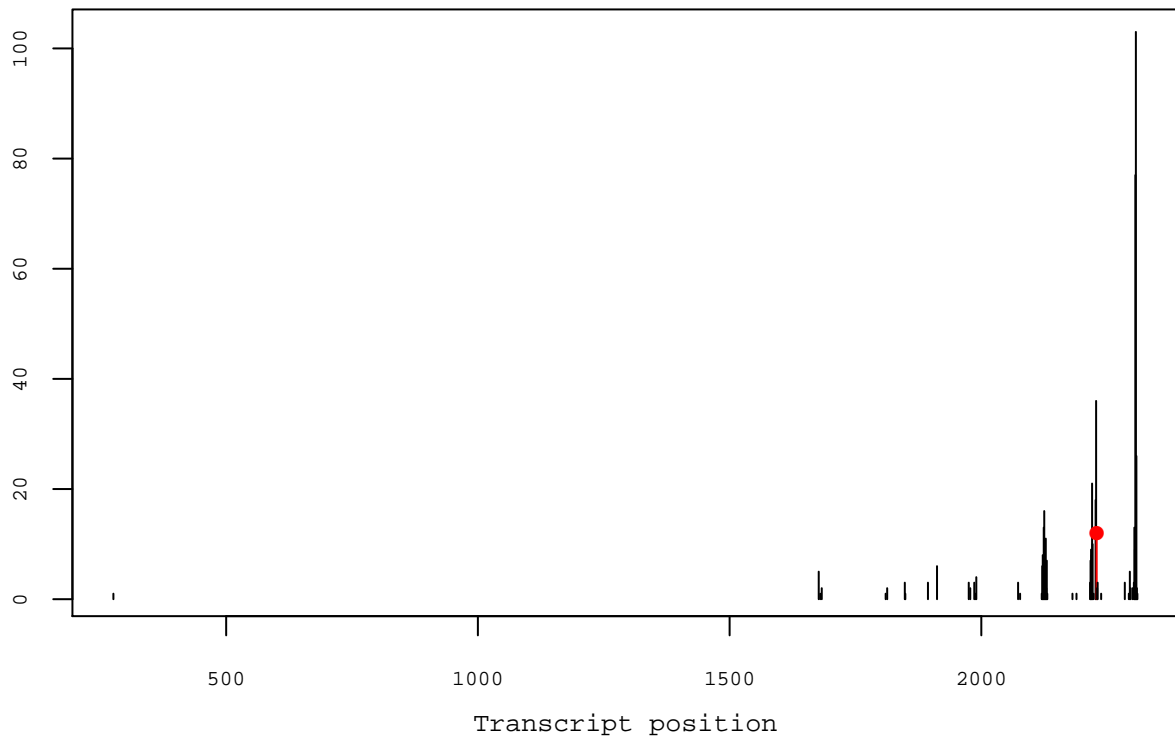


PggD12774_c125_g1_i2

5' CAACTCGTGTTTCGACTCGTTAAG-AAGCTCGTG '3
|||||
3' CGAGCACAAGCTGAGCAATTCGTT '5

Fragment Abundance



Cleavage site: 2229 Tag abundance: 12 Weighted abundance: 0.983 Category: 2
sRNA abundance: 22 Alignment score: 3 MFE ratio: 0.834 p-value: 0.027