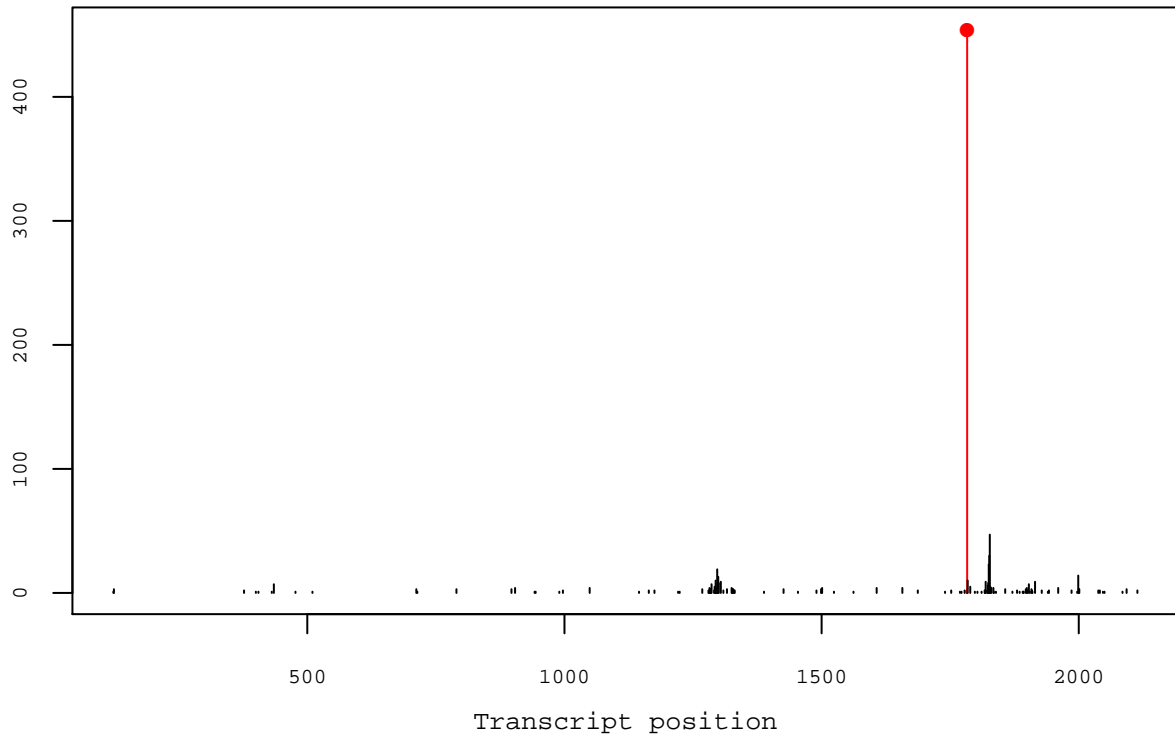


PggD620\_c8\_g1\_i1

5' CGCTGGA-GCAGCATCATCAGGATTCTTATC '3  
| | | | | | | | | | | | | | | |  
3' TCCGTCGTAGTAGTTCTAAGA '5

Fragment Abundance



Cleavage site: 1783    Tag abundance: 454    Weighted abundance: 454    Category: 0  
sRNA abundance: 139    Alignment score: 2    MFE ratio: 0.879    p-value: 0