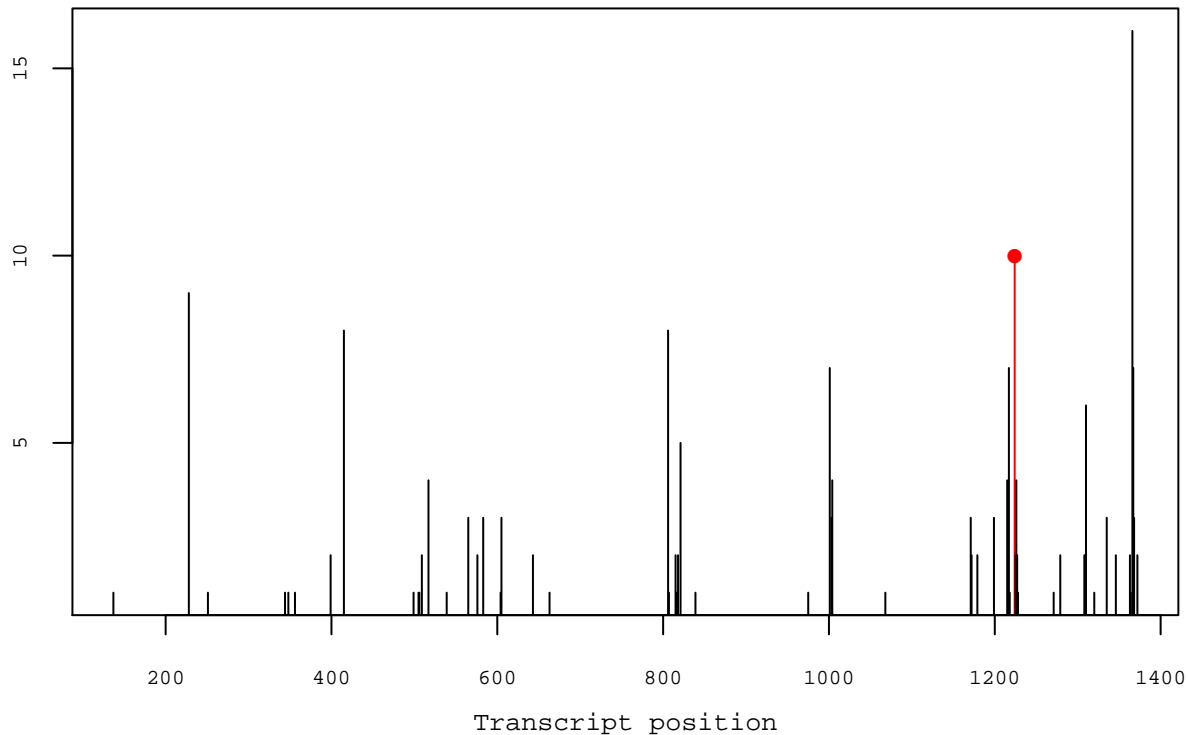


PggD3714\_c75\_g1\_i6

5' CAACAGCTGCCCCAAACAGACCCTTAGTTGAA '3  
||| |o| ||||| ||||| |||||  
3' CGTCGATGGGGTTTGTCTGGAAT '5

Fragment Abundance



Cleavage site: 1224 Tag abundance: 10 Weighted abundance: 10 Category: 2  
sRNA abundance: 5 Alignment score: 1.5 MFE ratio: 0.952 p-value: 0.02