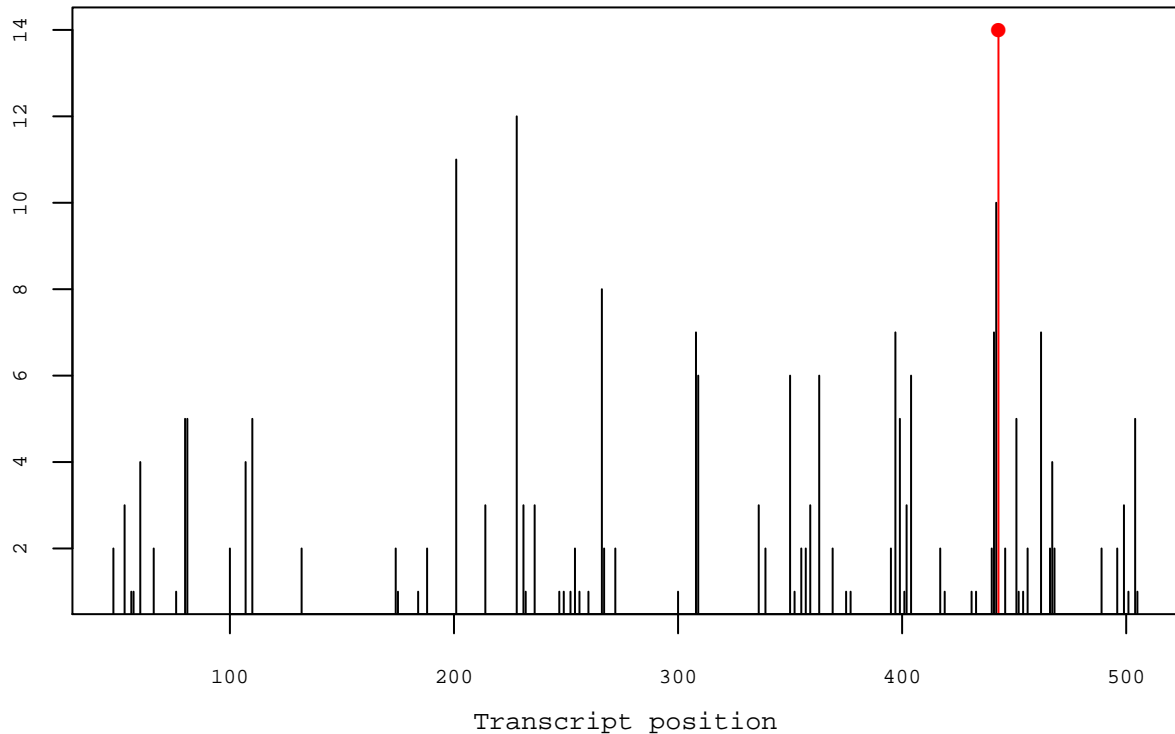


PggD5464\_c34\_g1\_i1

5' AAGTCTTCTAAGTCATTCCAAAATTATCTTT '3  
o |||||||||||||||||||||  
3' TAGAAGATTTCAGTAAGGTTTTAAA '5

Fragment Abundance



Cleavage site: 443 Tag abundance: 14 Weighted abundance: 0.406 Category: 0  
sRNA abundance: 29 Alignment score: 0.5 MFE ratio: 1 p-value: 0.041