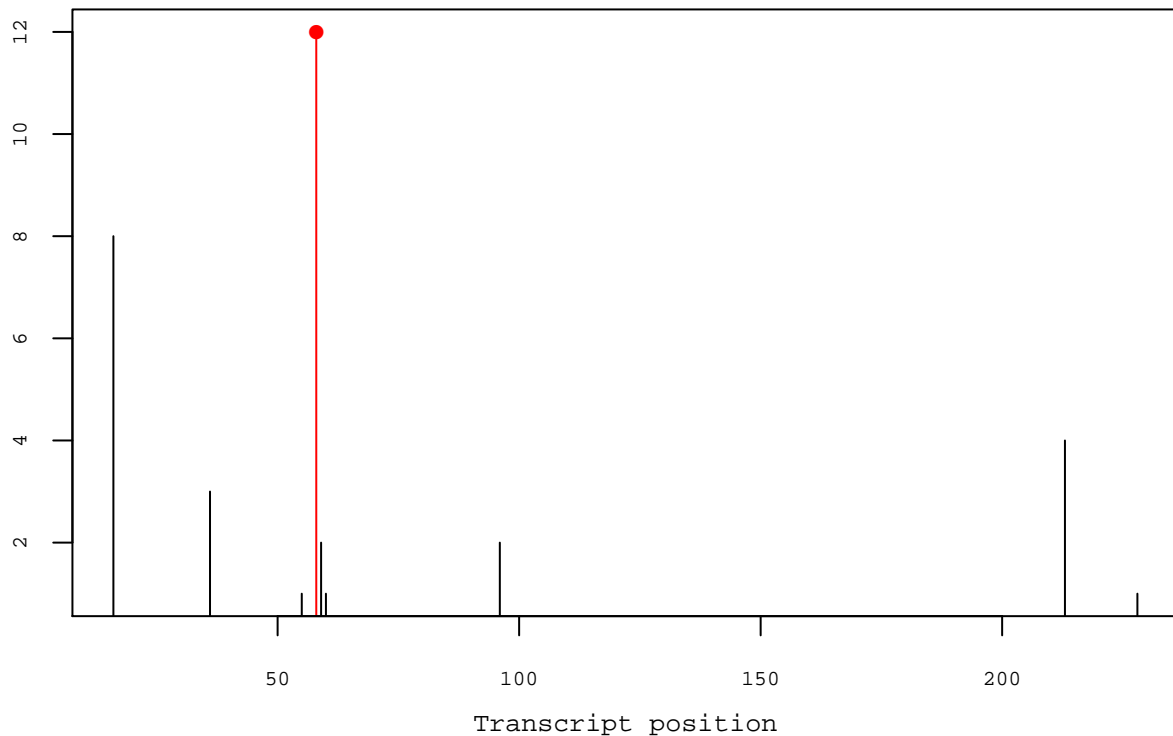


PggD285\_c1074\_g1\_i1

5' GGTTCGACAAAGATGCCTACGTCCTCGGGAGC '3  
|||||o|||o|||  
3' AAGCTGTTTCTATGGATGTAGGAG '5

Fragment Abundance



Cleavage site: 58 Tag abundance: 12 Weighted abundance: 0.229 Category: 0  
sRNA abundance: 7 Alignment score: 2 MFE ratio: 0.99 p-value: 0.012