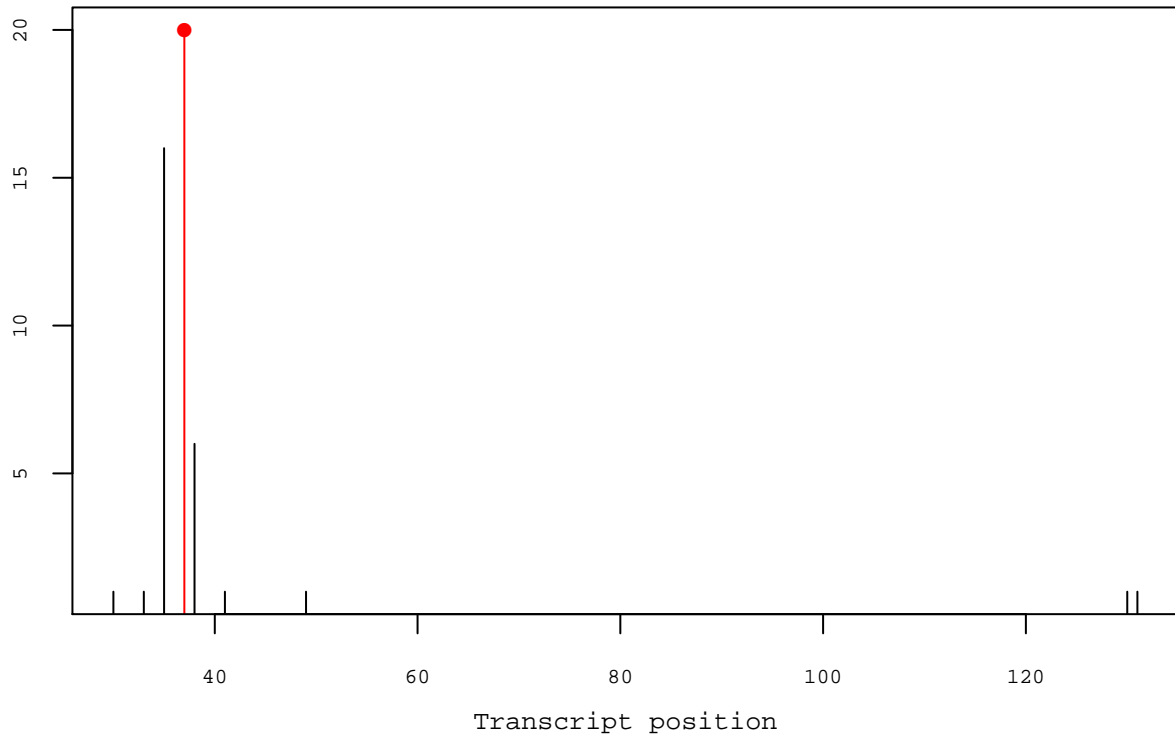


PggD3203_c174_g1_i1

5' CATGTAGAACTACGTCAGGAGTCCGGACAGAA '3
||||| |||||
3' ACATCTCGATGCAGTCCTCAGGCG '5

Fragment Abundance



Cleavage site: 37 Tag abundance: 20 Weighted abundance: 19.333 Category: 0
sRNA abundance: 27 Alignment score: 2 MFE ratio: 0.824 p-value: 0.006