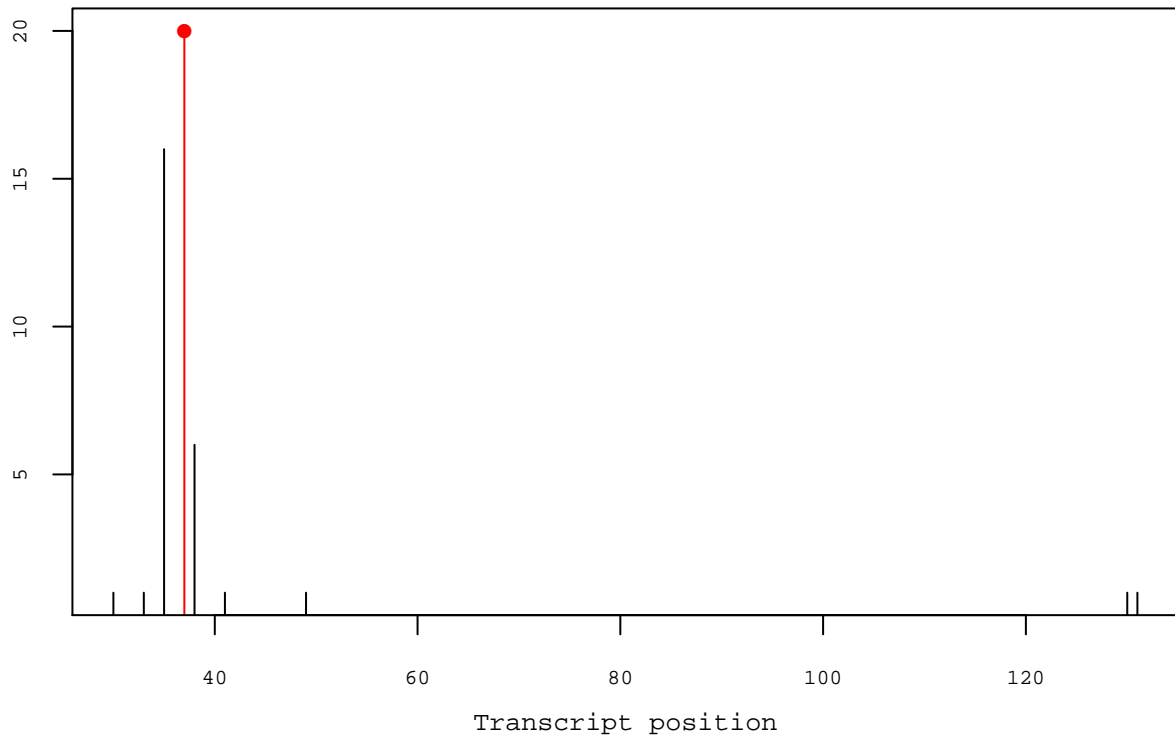


PggD3203_c174_g1_i1

5' CATGTAGAACTACGTCAGGAGTCCGGACAGAA '3

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3' GCATCTCGATGCAGTCCTCAGGCC 5'



Cleavage site: 37 Tag abundance: 20 Weighted abundance: 19.333 Category: 0
sRNA abundance: 64 Alignment score: 1.5 MFE ratio: 0.848 p-value: 0.006