

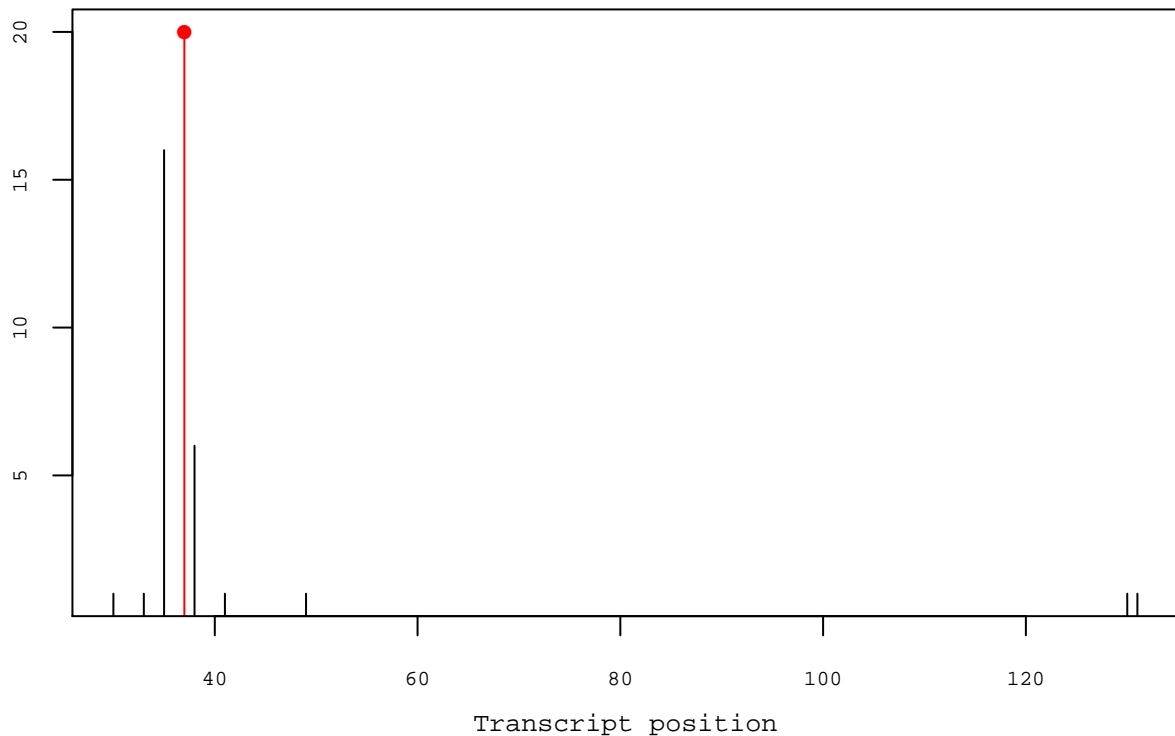
PggD3203\_c174\_g1\_i1

5' CATGTAGAACTACGTCAGGAGTCCGGACAGAA '3

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3' GCATCTCGATGCAGTTCTCAGGCG '5

Fragment Abundance



Cleavage site: 37 Tag abundance: 20 Weighted abundance: 19.333 Category: 0  
sRNA abundance: 13 Alignment score: 3.5 MFE ratio: 0.792 p-value: 0.006