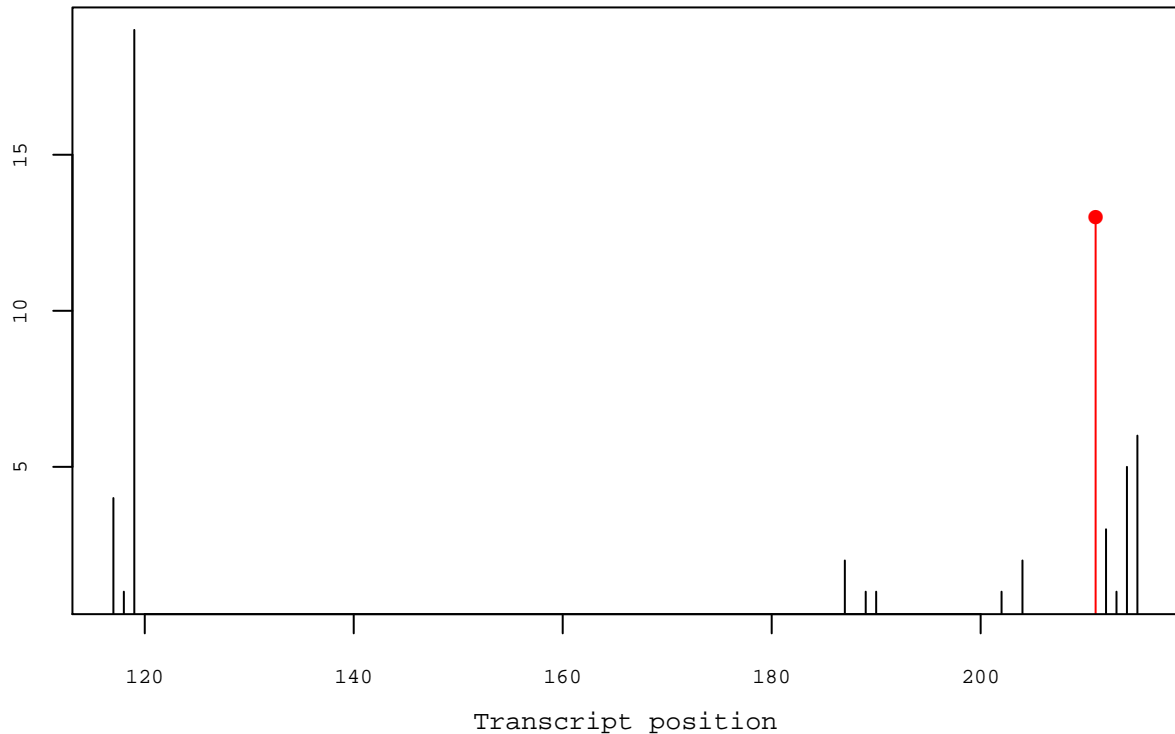


PggD9015_c120_g1_i1

5' CGGACTCTTGACGTAGCTCTACGTGCGGACCA '3
 |||||o||||||| |||||
 3' CTGAGGACTGCATCAAGATGCACG '5



Cleavage site: 211	Tag abundance: 13	Weighted abundance: 6.5	Category: 2
sRNA abundance: 10	Alignment score: 2.5	MFE ratio: 0.849	p-value: 0.014