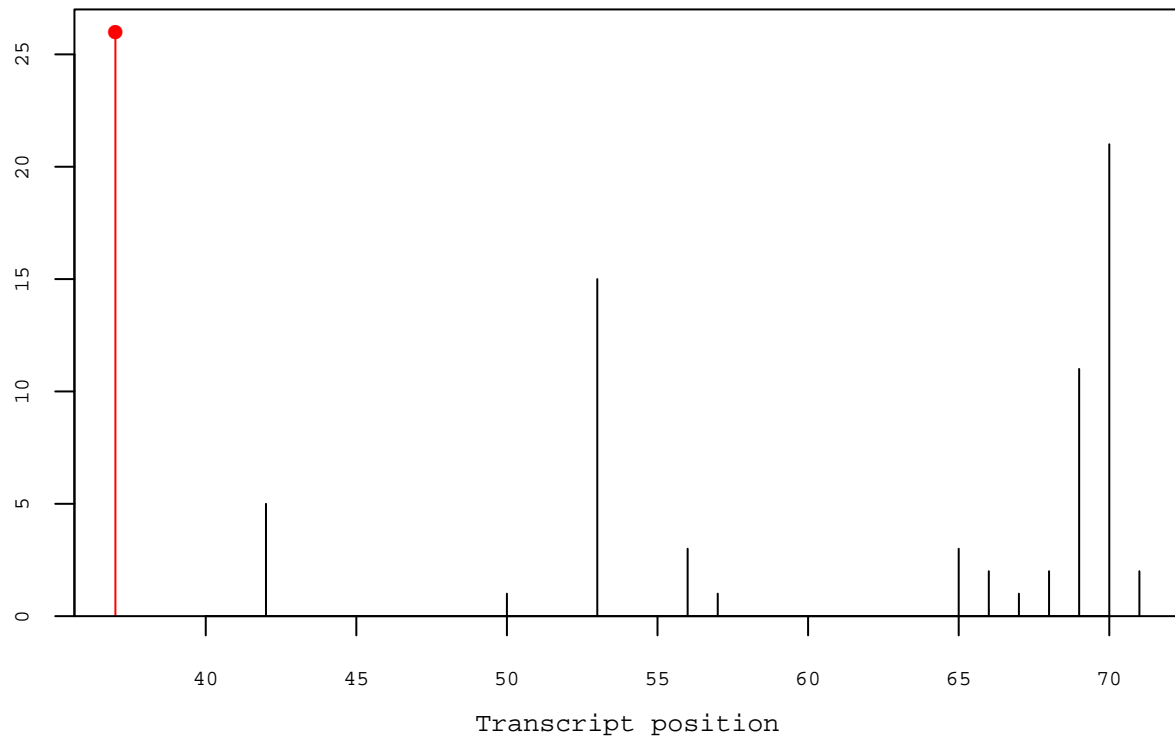


PggD2278_c153_g1_i1

5' CCTATAACCTGTTTGCAGGTTACAGACGTACG '3
||| ||||| ||||| ||||| |||||
3' TTGTACAAACGTCCAATGTCT '5

Fragment Abundance



Cleavage site: 37 Tag abundance: 26 Weighted abundance: 1.083 Category: 0
sRNA abundance: 10 Alignment score: 1 MFE ratio: 0.869 p-value: 0.011