

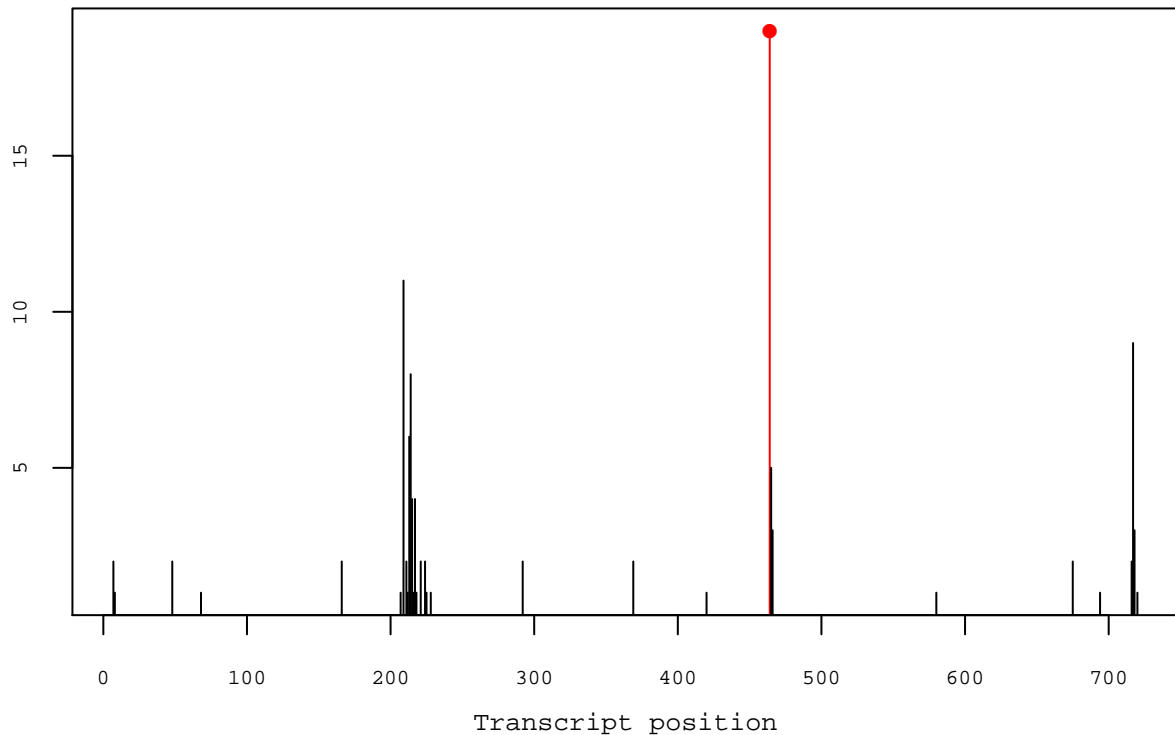
PggD10889_c79_g1_i1

5' CATCCTCCTCCCAAAGTCTATAA-AAGTGCCAA '3

|||||

3' TCAGGGTTTCAGATATTGTT '5

Fragment Abundance



Cleavage site: 464 Tag abundance: 19 Weighted abundance: 1.9 Category: 0
sRNA abundance: 6 Alignment score: 4 MFE ratio: 0.731 p-value: 0.011