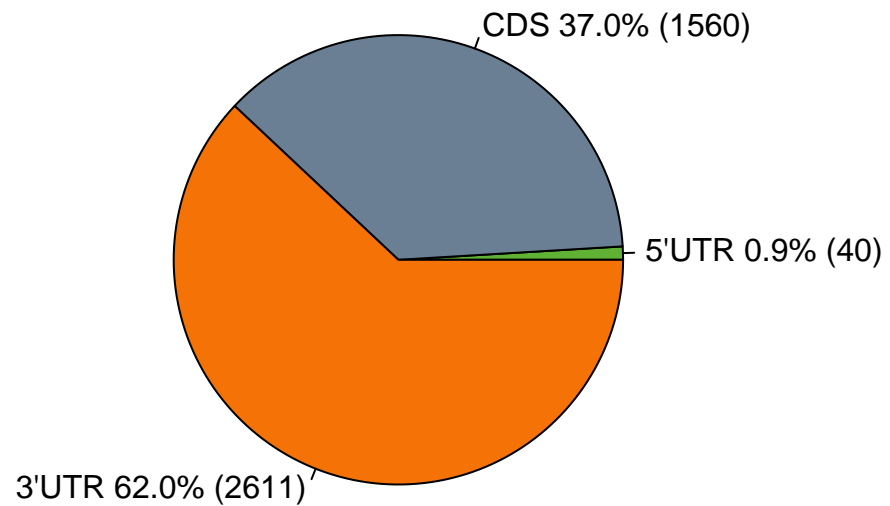


Distribution of 4211 m6A sites across transcript sections



Distribution of 861643 null_m6A sites across transcript sections

