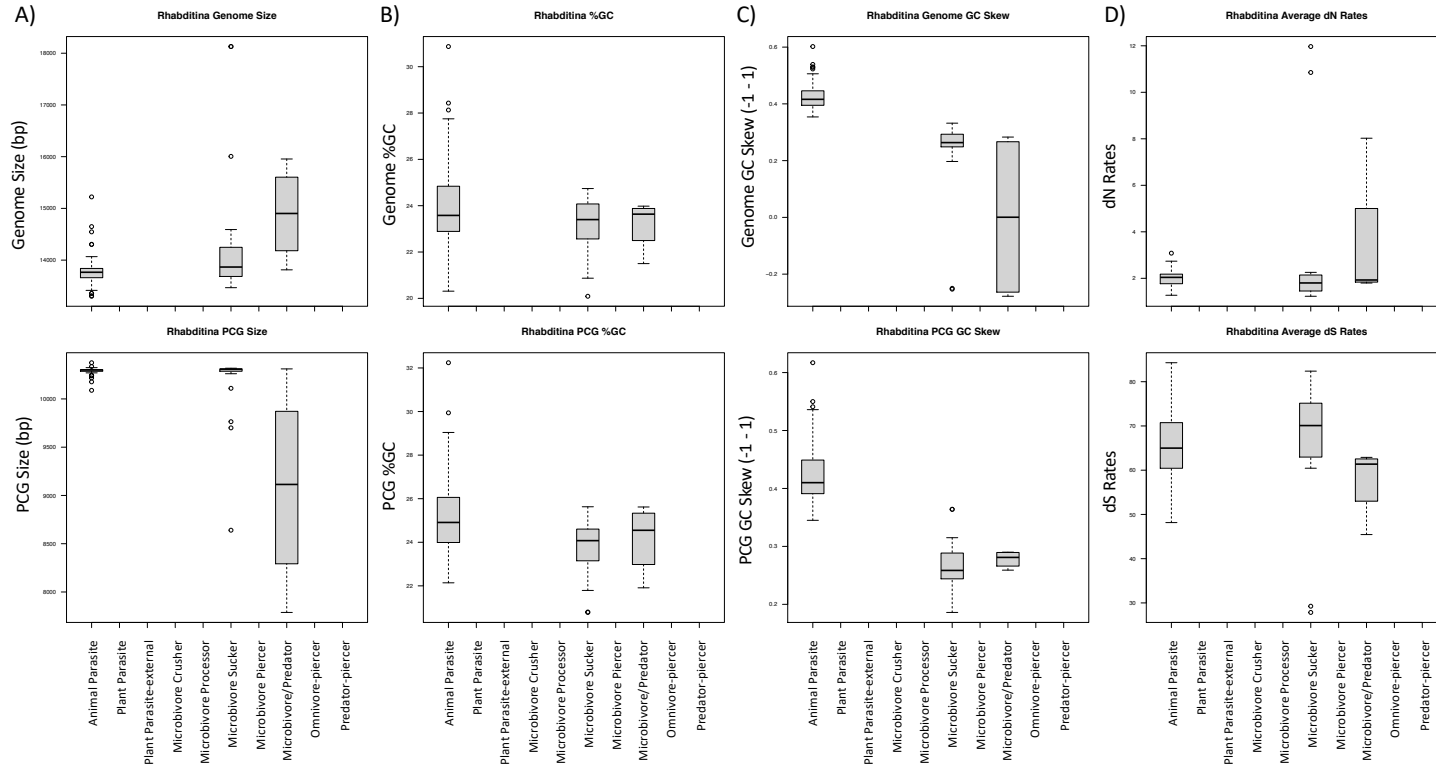


Supplemental Figure 10: Rhabditina Mitogenome Characteristics and Substitution Rates by Habit



SI Figure 10: Rhabditina Mitogenome Characteristics by Feeding Habit

Box and whisker plots for total genome and PCG characteristics for A) size, B) %GC content, C) GC compositional skew, and D) substitution rates for PCG sequences for the Rhabditina suborder. Medians and quantiles were calculated for each characteristic based on the life trait classification for feeding Habit. Rhabditina feeding habits were significant for PCG proportion of the genome, genome GC skews, and PCG GC skews.