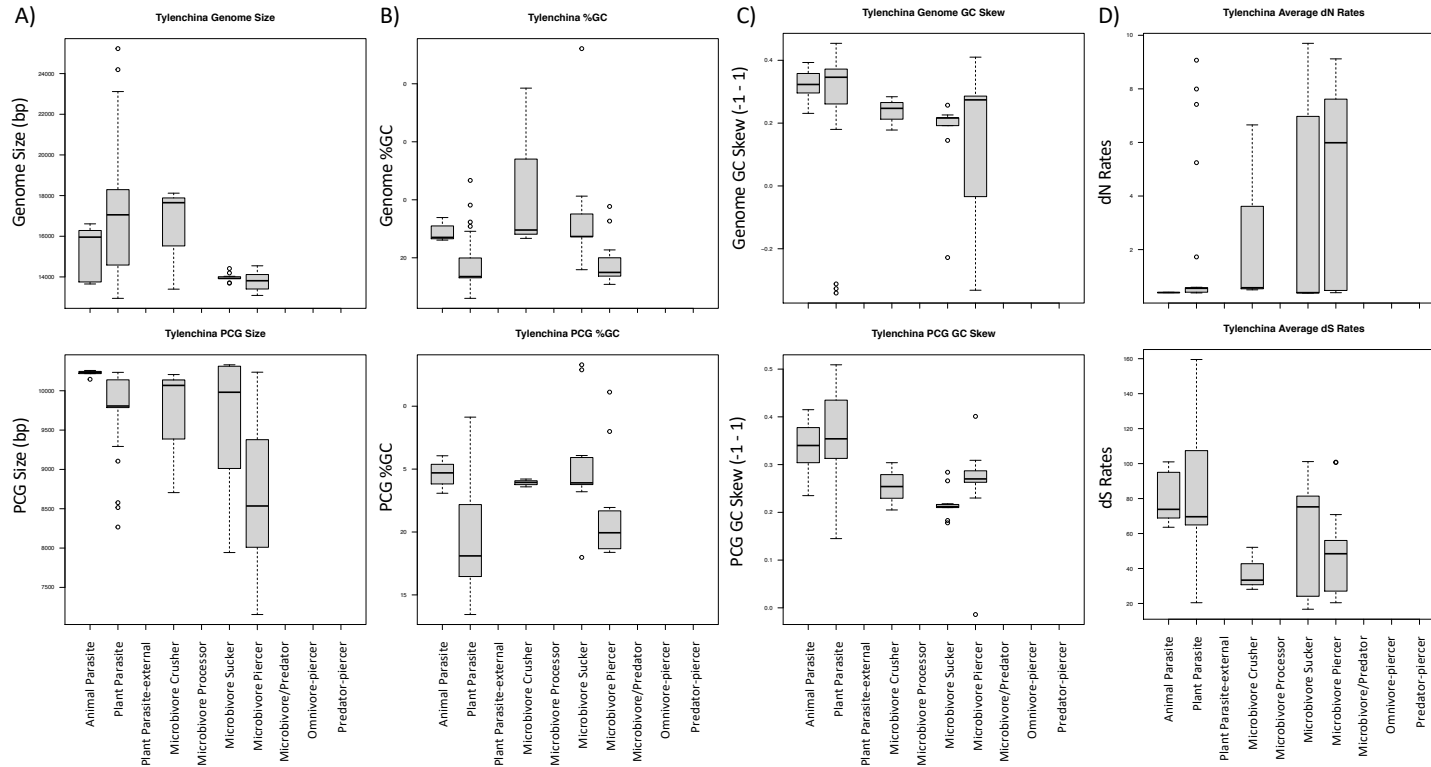


Supplemental Figure 7: Tylenchina Mitogenome Characteristics and Substitution Rates by Habit



SI Figure 7: Tylenchina 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 Tylenchina suborder. Medians and quantiles were calculated for each characteristic based on the life trait classification for feeding Habit. Tylenchina feeding habit was significant for genome %GC, Genome GC skews, and PCG GC skews.