

## Supplementary Material

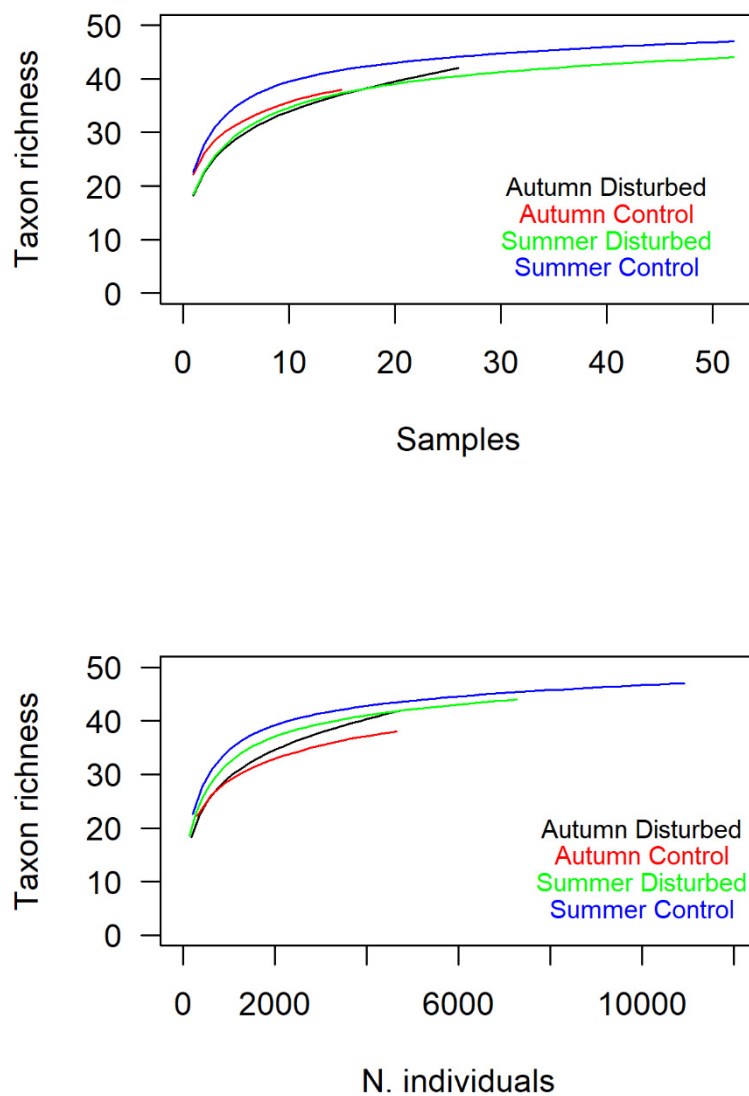


Fig. S1. Taxa Accumulation Curves obtained using the “rarefaction” method in the vegan R package, version 2.5-6 (Oksanen et al. 2015). The y-axis represents the expected taxon richness, while the x-axis represents the number of samples (top) and the number of individuals (bottom) for each site and each season.