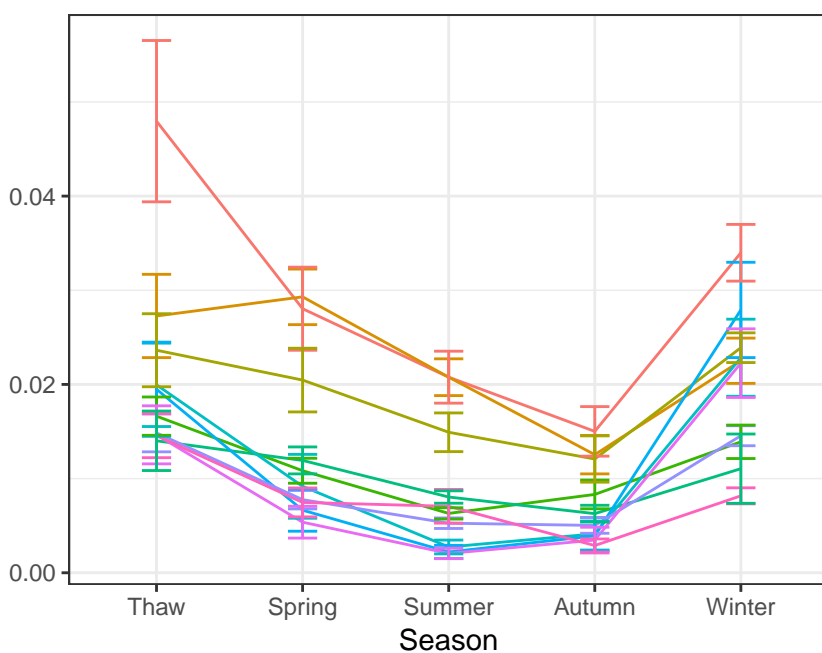


A

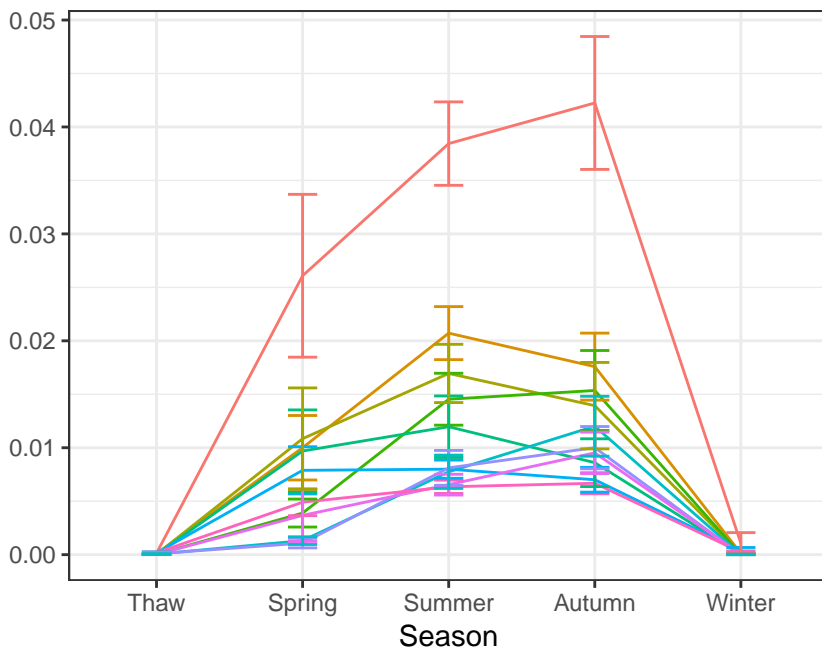
Mean Relative Abundance



- k__Bacteria; p__Acidobacteria; c__DA052; o__Ellin6513; f__; g__; s__
- k__Bacteria; p__Acidobacteria; c__DA052; o__Ellin6513; f__; g__; s__
- k__Bacteria; p__Proteobacteria; c__Alphaproteobacteria; o__Rhodospirillales; f__Acetobacteraceae; g__; s__
- k__Bacteria; p__Acidobacteria; c__Acidobacteriia; o__Acidobacteriales; f__Acidobacteriaceae; g__; s__
- k__Bacteria; p__Actinobacteria; c__Actinobacteria; o__Actinomycetales; f__; g__; s__
- k__Bacteria; p__Acidobacteria; c__Acidobacteriia; o__Acidobacteriales; f__Koribacteraceae; g__; s__
- k__Bacteria; p__Acidobacteria; c__Acidobacteriia; o__Acidobacteriales; f__Acidobacteriaceae; g__; s__
- k__Bacteria; p__Acidobacteria; c__Acidobacteriia; o__Acidobacteriales; f__Koribacteraceae; g__; s__
- k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Xanthomonadales; f__Sinobacteraceae; g__; s__
- k__Bacteria; p__Acidobacteria; c__Solibacteres; o__Solibacterales; f__; g__; s__

B

Mean Relative Abundance



- k__Bacteria; p__Proteobacteria; c__Alphaproteobacteria; o__Rhizobiales
- k__Bacteria; p__Proteobacteria; c__Alphaproteobacteria; o__Rhodospirillales; f__Rhodospirillaceae; g__; s__
- k__Bacteria; p__Actinobacteria; c__Acidimicrobiia; o__Acidimicrobiales; f__EB1017; g__; s__
- k__Bacteria; p__AD3; c__ABS-6; o__; f__; g__; s__
- k__Bacteria; p__Acidobacteria; c__[Chloracidobacteria]; o__RB41; f__; g__; s__
- k__Bacteria; p__Gemmatimonadetes; c__Gemmatimonadetes; o__N1423WL; f__; g__; s__
- k__Bacteria; p__Bacteroidetes; c__[Saprospirae]; o__[Saprospirales]; f__Chitinophagaceae; g__; s__
- k__Bacteria
- k__Bacteria; p__WPS-2; c__; o__; f__; g__; s__
- k__Bacteria; p__Verrucomicrobia; c__[Spartobacteria]; o__[Chthoniobacteriales]; f__[Chthoniobacteraceae]; g__DA101; s__