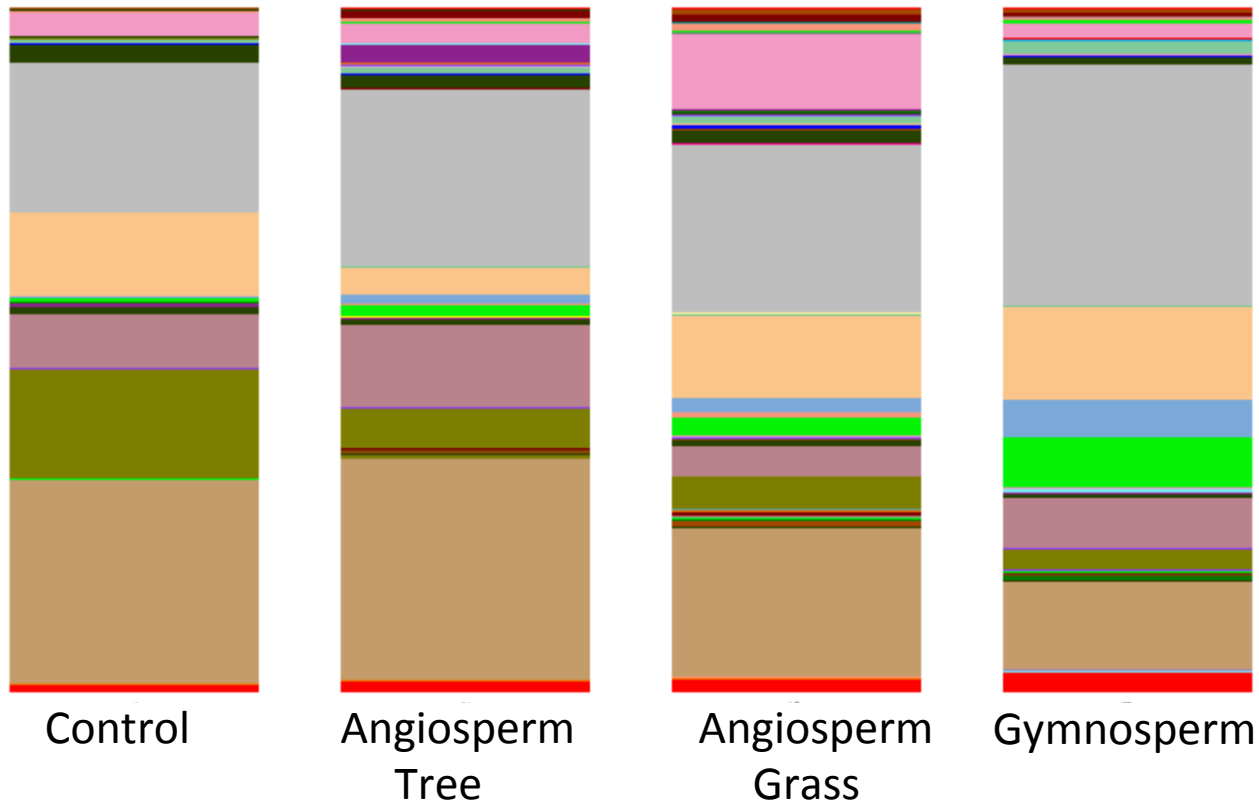


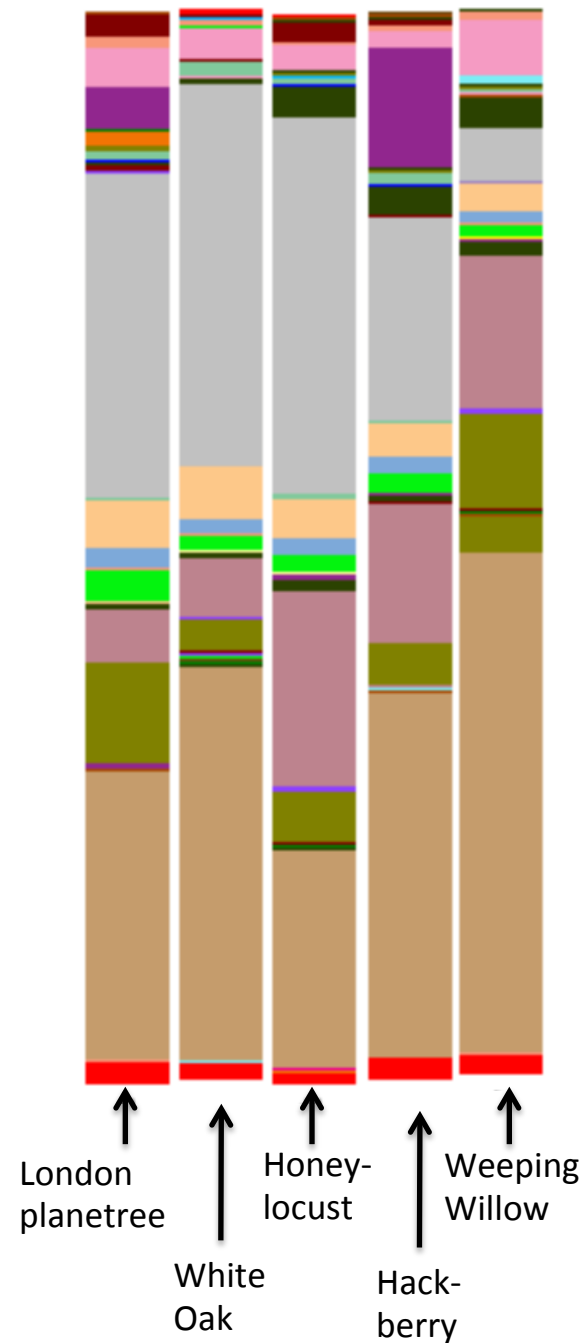
Composition by Broad Taxonomic Group of Plant Host

Percentage
Relative
Abundance



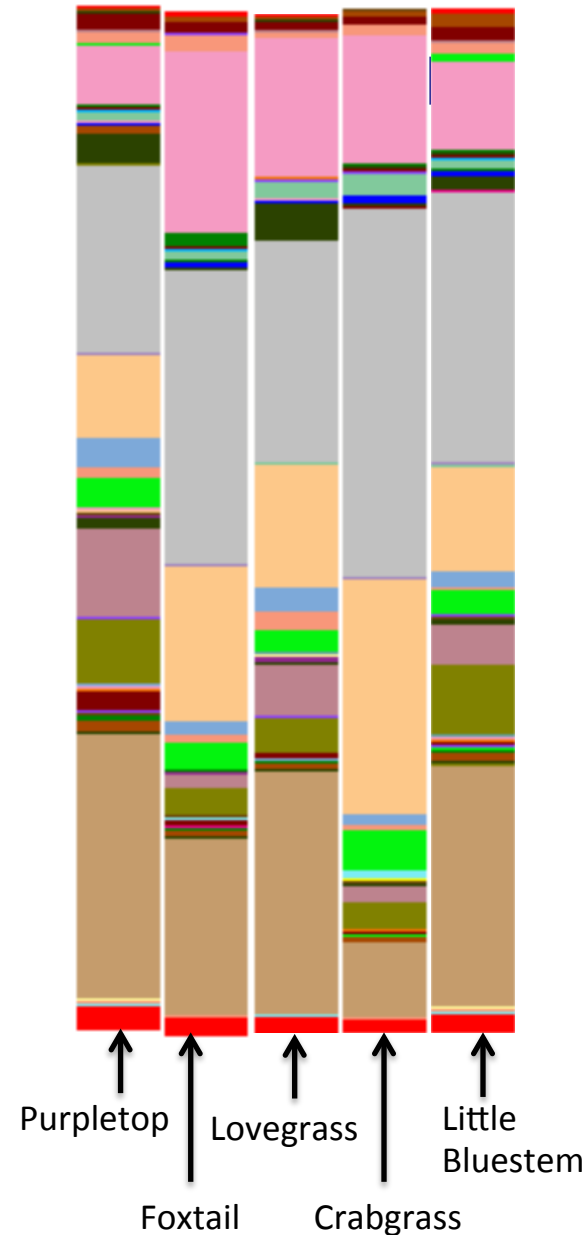
	Unassigned;Other;Other;Other
	k_Bacteria;p_Actinobacteria;c_Actinobacteria;o_Actinomycetales
	k_Bacteria;p_Armatimonadetes;c_[Fimbriimonadia];o_[Fimbriimonadales]
	k_Bacteria;p_Firmicutes;c_Bacilli;o_Bacillales
	k_Bacteria;p_Firmicutes;c_Bacilli;o_Lactobacillales
	k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales
	k_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Rhizobiales
	k_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Rhodospirillales
	k_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Sphingomonadales
	k_Bacteria;p_Proteobacteria;c_Betaproteobacteria;o_Burkholderiales
	k_Bacteria;p_Proteobacteria;c_Betaproteobacteria;o_Neisseriales
	k_Bacteria;p_Proteobacteria;c_Deltaproteobacteria;o_Myxococcales
	k_Bacteria;p_Proteobacteria;c_Gammaproteobacteria;o_Oceanospirillales
	k_Bacteria;p_Proteobacteria;c_Gammaproteobacteria;o_Pseudomonadales
	k_Bacteria;p_TM7;c_TM7-3;o_

Composition of the Angiosperm Tree Samples (% Relative Abundance)



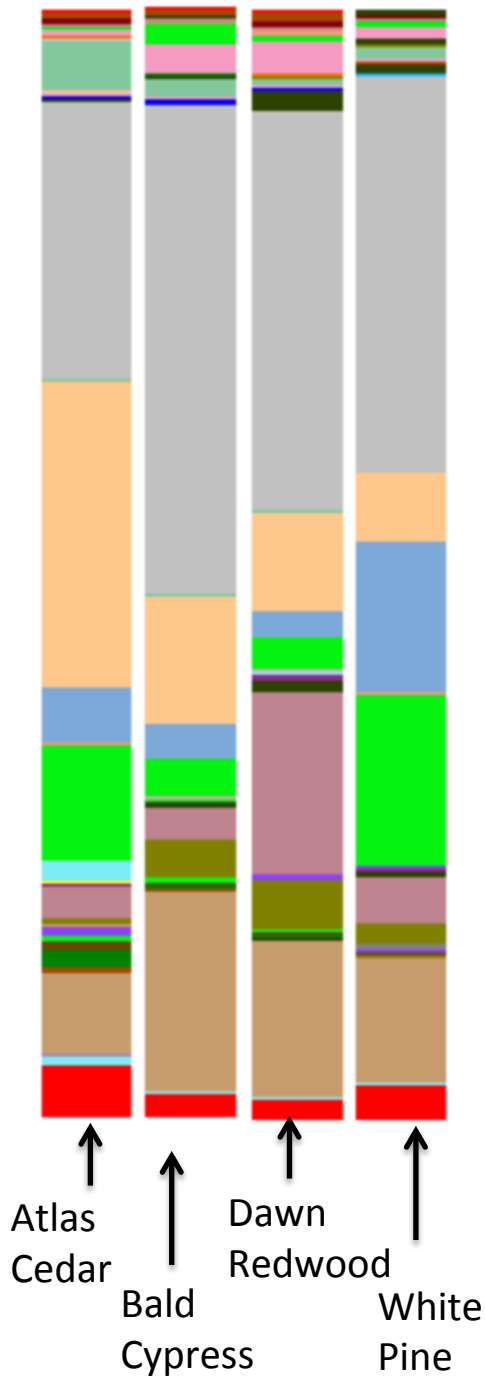
	Unassigned;Other;Other;Other
	k_Bacteria;p_Actinobacteria;c_Actinobacteria;o_Actinomycetales
	k_Bacteria;p_Firmicutes;c_Bacilli;o_Bacillales
	k_Bacteria;p_Firmicutes;c_Bacilli;o_Lactobacillales
	k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales
	k_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Rhizobiales
	k_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Rhodospirillales
	k_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Sphingomonadales
	k_Bacteria;p_Proteobacteria;c_Betaproteobacteria;o_Burkholderiales
	k_Bacteria;p_Proteobacteria;c_Betaproteobacteria;o_Neisseriales
	k_Bacteria;p_Proteobacteria;c_Deltaproteobacteria;o_Myxococcales
	k_Bacteria;p_Proteobacteria;c_Gammaproteobacteria;o_Legionellales
	k_Bacteria;p_Proteobacteria;c_Gammaproteobacteria;o_Oceanospirillales
	k_Bacteria;p_Proteobacteria;c_Gammaproteobacteria;o_Pseudomonadales
	k_Bacteria;p_TM7;c_TM7-3;o_

Composition of the Angiosperm Grass Samples (% Relative Abundance)



	Unassigned;Other;Other;Other
	k_Bacteria;p_Actinobacteria;c_Actinobacteria;o_Actinomycetales
	k_Bacteria;p_Actinobacteria;c_Actinobacteria;o_Bifidobacteriales
	k_Bacteria;p_Actinobacteria;c_Thermoleophila;o_Solirubrobacterales
	k_Bacteria;p_Chloroflexi;c_Chloroflexi;o_Roseiflexales
	k_Bacteria;p_Firmicutes;c_Bacilli;o_Bacillales
	k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales
	k_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Rhizobiales
	k_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Rhodobacterales
	k_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Rhodospirillales
	k_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Sphingomonadales
	k_Bacteria;p_Proteobacteria;c_Betaproteobacteria;o_Burkholderiales
	k_Bacteria;p_Proteobacteria;c_Betaproteobacteria;o_Neisseriales
	k_Bacteria;p_Proteobacteria;c_Deltaproteobacteria;o_Myxococcales
	k_Bacteria;p_Proteobacteria;c_Gammaproteobacteria;o_Methylococcales
	k_Bacteria;p_Proteobacteria;c_Gammaproteobacteria;o_Oceanospirillales
	k_Bacteria;p_Proteobacteria;c_Gammaproteobacteria;o_Pseudomonadales
	k_Bacteria;p_Proteobacteria;c_Gammaproteobacteria;o_Xanthomonadales
	k_Bacteria;p_TM7;c_TM7-3;o_

Composition of the Gymnosperm Samples (% Relative Abundance)



	Unassigned;Other;Other; Other
	k_Bacteria;p_Actinobacteria;c_Actinobacteria;o_Actinomycetales
	k_Bacteria;p_Armatimonadetes;c_[Fimbriimonadia];o_[Fimbriimonadales]
	k_Bacteria;p_Firmicutes;c_Bacilli;o_Bacillales
	k_Bacteria;p_Firmicutes;c_Bacilli;o_Lactobacillales
	k_Bacteria;p_Firmicutes;c_Clostridia;o_Clostridiales
	k_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_
	k_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Rhizobiales
	k_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Rhodospirillales
	k_Bacteria;p_Proteobacteria;c_Alphaproteobacteria;o_Sphingomonadales
	k_Bacteria;p_Proteobacteria;c_Betaproteobacteria;o_Burkholderiales
	k_Bacteria;p_Proteobacteria;c_Betaproteobacteria;o_Neisseriales
	k_Bacteria;p_Proteobacteria;c_Deltaproteobacteria;o_Myxococcales
	k_Bacteria;p_Proteobacteria;c_Gammaproteobacteria;o_Pseudomonadales
	k_Bacteria;p_Proteobacteria;c_Gammaproteobacteria;o_Vibrionales