

Supporting Material

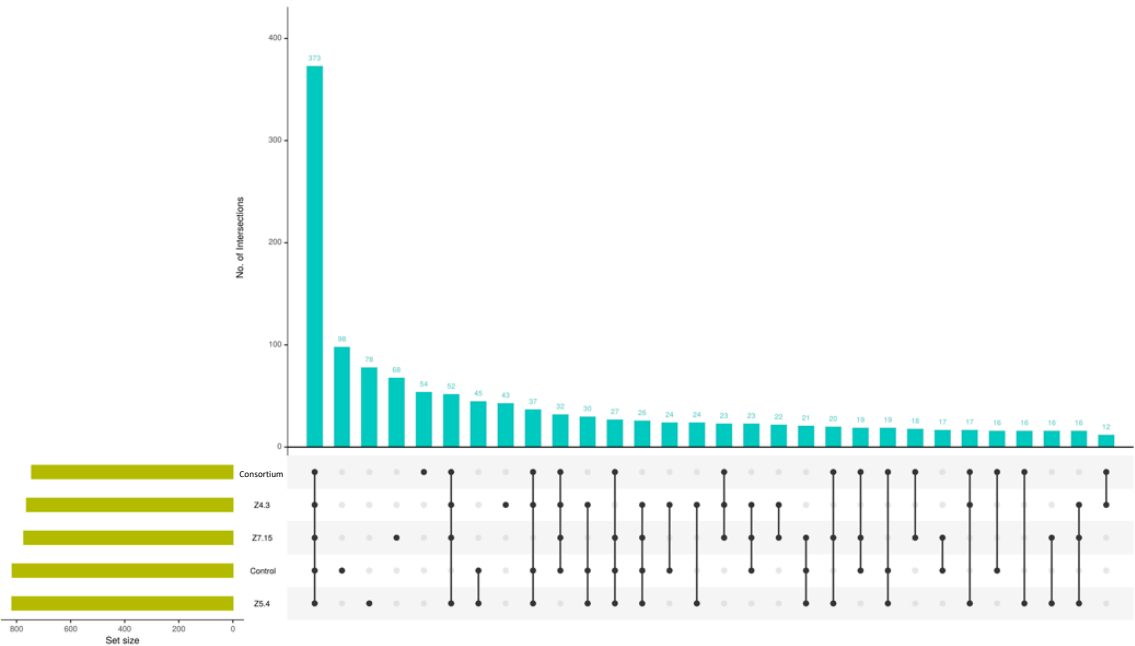


Figure S1. Number of unique bacterial taxa (species) in the different treatments in Autumn sampling (plants in nursery, before transplanting).

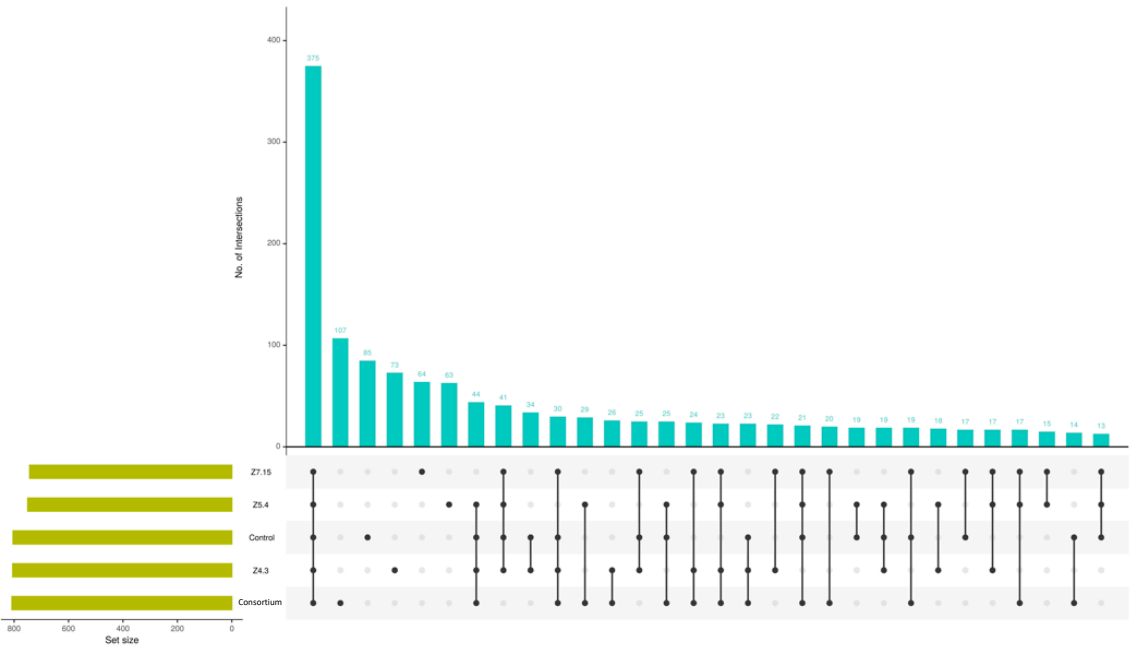


Figure S2. Number of unique bacterial taxon (species) in the different treatments in Spring sampling (after trasplanting).

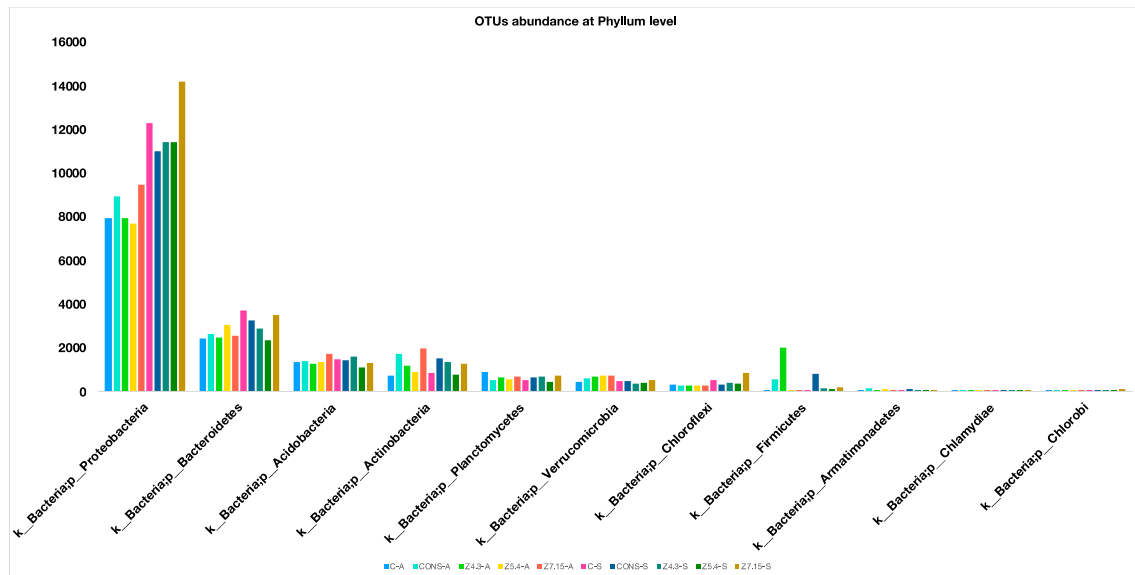


Figure S3. OTUS abundance at phylum level. Data obtained from analysis done by EasyMap.



Figure S4. OTUS abundance at class level: a) phylum Proteobacteria; b) phylum Bacteroidetes; c) phylum Acidobacteria and d) phylum Actinobacteria. Data obtained from analysis done by EasyMap.

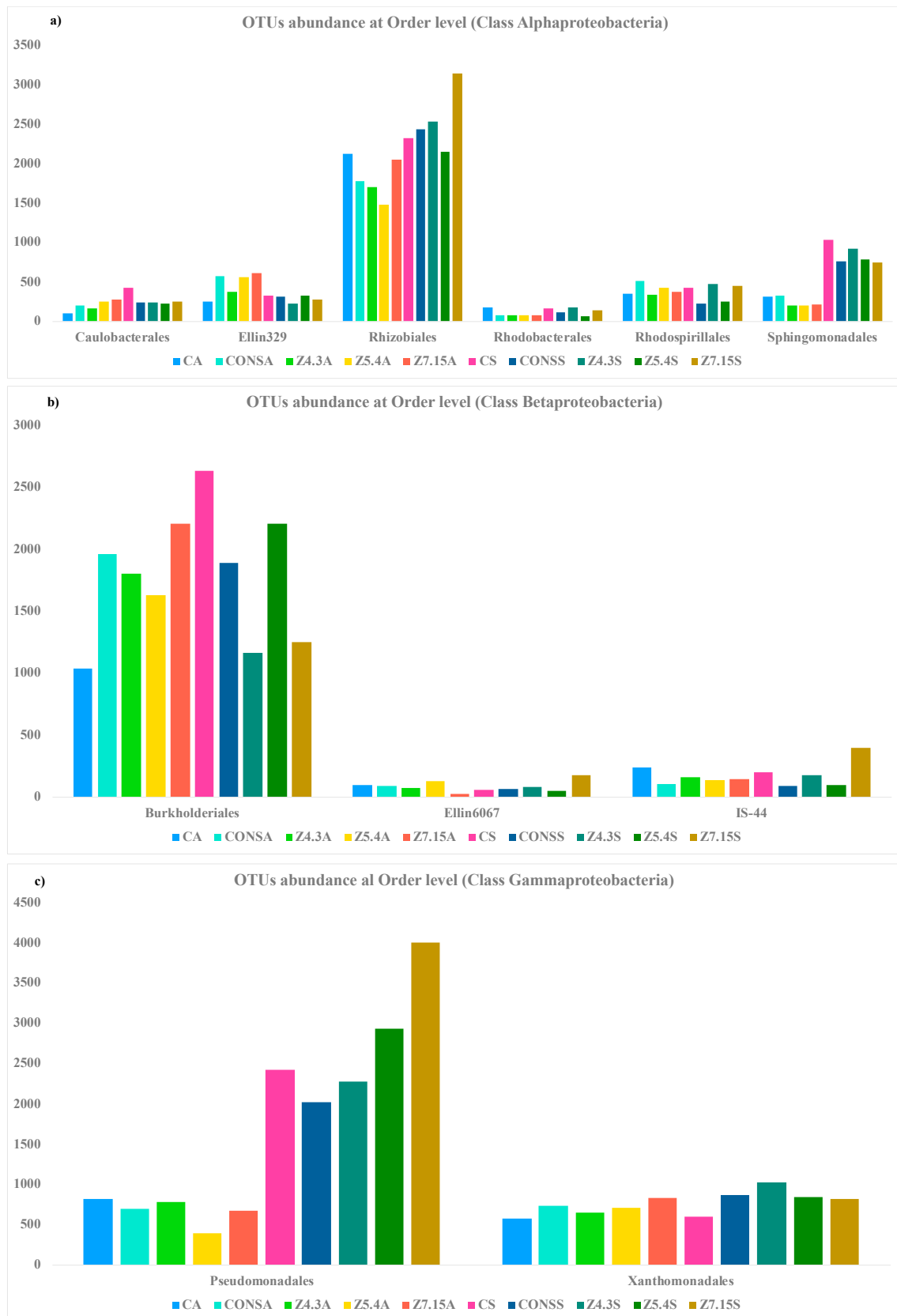


Figure S5. OTUs abundance at order level: a) class Alphaproteobacteria; b) class Betaproteobacteria and c) class Gammaproteobacteria. Data obtained from analysis done by EasyMap.



Figure S6. OTUS abundance at order level: a) phylum Bacteroidetes; b) phylum Acidobacteria and c) phylum Actinobacteria. Data obtained from analysis done by EasyMap.