Supplementary Figure 1. Rhizosphere bacterial community composition across studies of wild, landrace and modern plants. Principal Coordinate Analysis (PCoA) of Bray-Curtis dissimilarities of 16S rRNA data. a) PCoA with samples colored by plant species. b) PCoA with samples colored by study. The source of the data (study) was the main explaining variable as assessed by PERMANOVA (29.1%; \( P<0.005 \)).
Supplementary Figure 2. α-diversity of 16S sequence data of wild, landrace and modern plant species for rhizosphere and roots. (a) Chao1 and (b) Phylogenetic diversity for all the plants included in the meta-analysis. Bacterial diversity on/in the roots is less than in the rhizosphere.