



Figure S3. Bar plot showing relative mean proportion of fungal taxa in the endophytic fungal community in all sample groups ($n = 4$ per group). The top 10% of filtered, fungal reads are included and aggregated at genus level (if not available, the lowest possible taxonomic level). Fungal taxa prefixes follow UNITE notation (“c_”, class; “f_”, family; “g_”, genus; “o_”, order; “p_”, phylum). “Other” means a pool of ASVs from all other taxa. Sample codes: year_sample, where sample type is either “soil”, “carr” (soil or carrot both from site 1), 1037 or 326 or 112 (site 2 horseradish samples from three soil types) or A-W (accession codes from site 1).