



SUPPLEMENTARY FIGURE 11 | Enriched and depleted resident microbial groups in inoculated plants. Enrichment and depletion of families in DKB177 (A), SX7341 (B), and P3707VYH (C) shown for WW- (blue) and DS-treated (red) plants. Only significant ($P \leq 0.05$) shifts in inoculation were considered, with OTUs collapsed at the taxonomic level of the family. Families comprised of a single OTU were omitted. "Unknown" taxonomic levels were collapsed to the deepest common levels, comma-separated, and "unknown" taxa represented as "-". FC, fold change; Un., unknown.