

Supplemental Figure 7

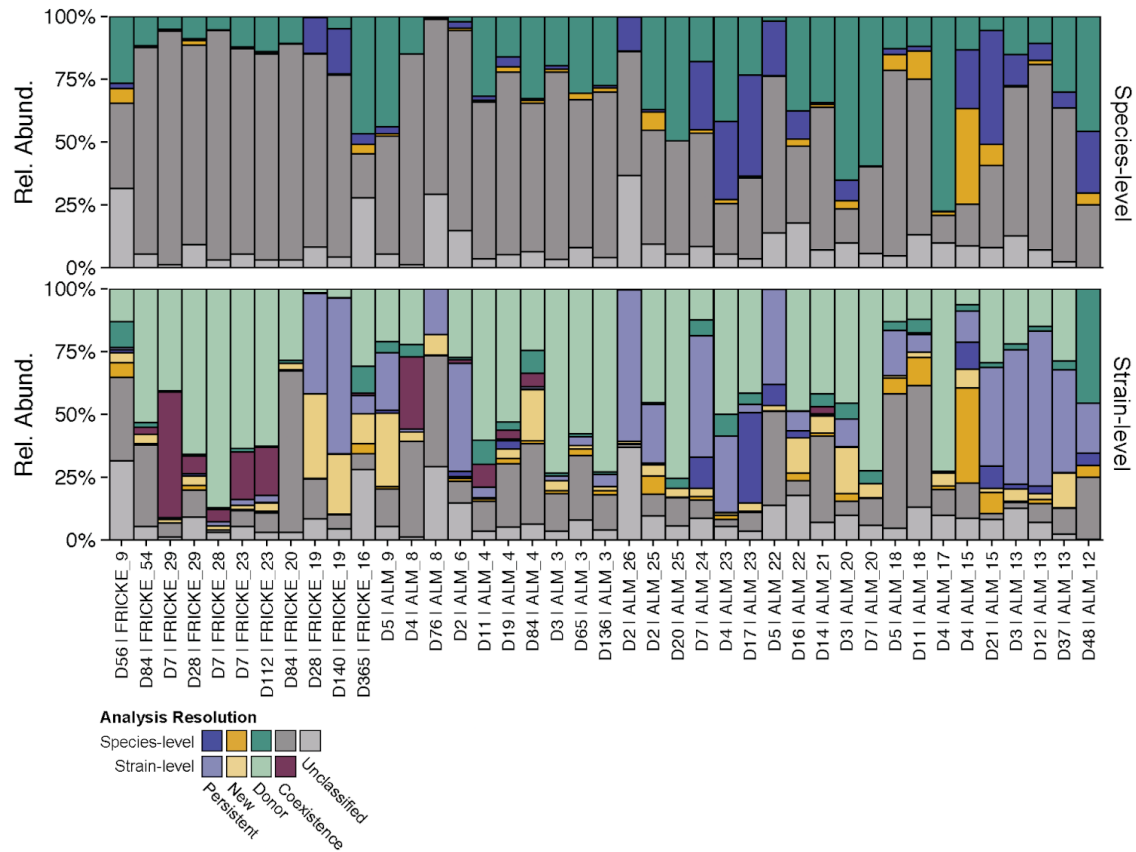


Figure D7. Microbial tracking across individual metagenomic samples of the *rCDI Cohort*.

Donor-derived strains and species (exclusively shared with donor but insufficient resolution for strain prediction) account for large and stable relative abundances across all post-FMT patient samples in the *rCDI Cohort*, whereas contributions of recipient-derived strains are comparatively smaller.