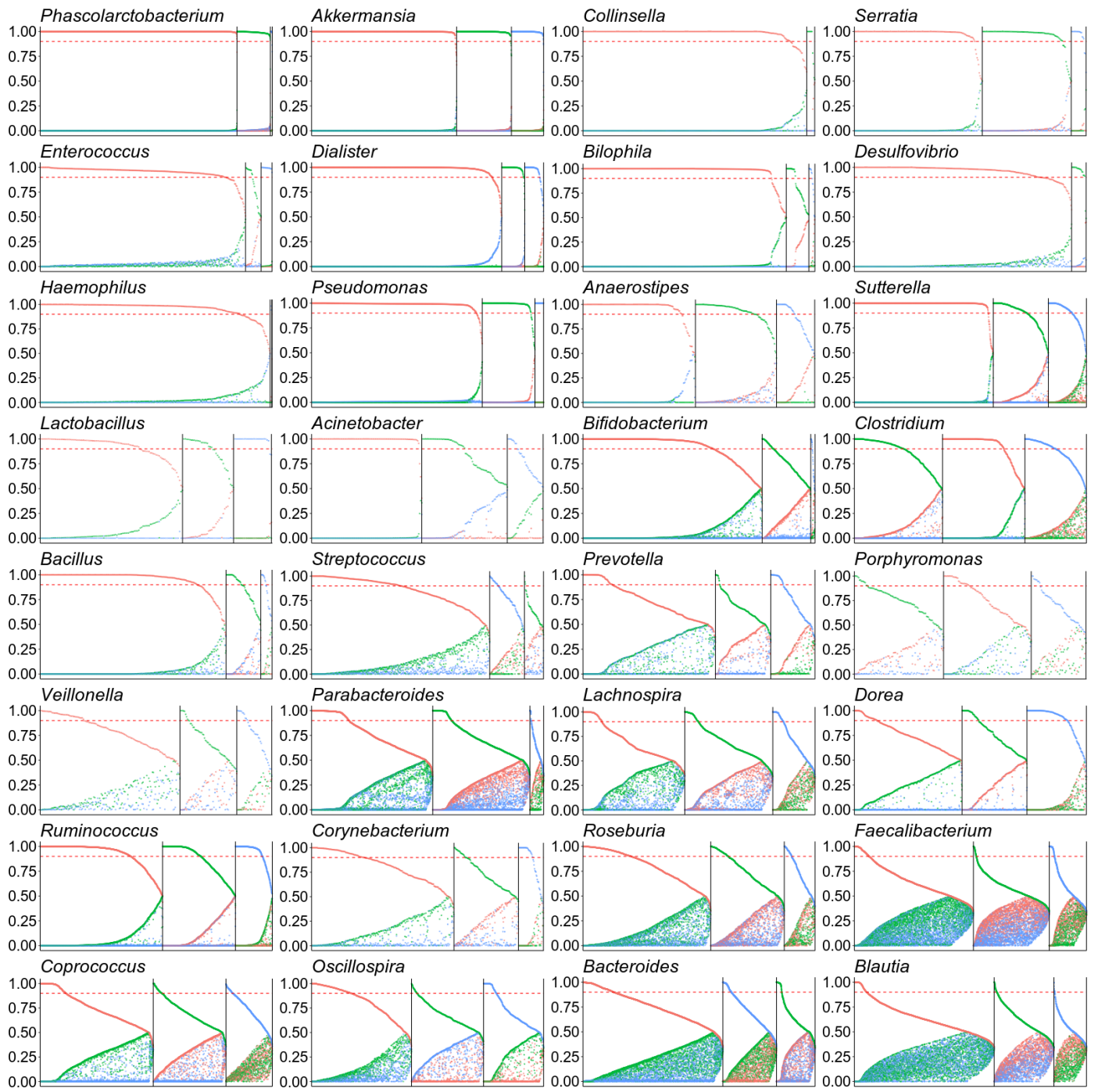


Within-group abundance



Microbiome samples