Default: no differential taxa (continuous-Stool) g:GMPR+DESeq2 h:GMPR+edgeR i:LDM a:ANCOM-BC b:Beta-binomial c:DACOMP d:DESeq2 e:edgeR f:GLM(quasipoisson) Depth confounding None 1.00 sample=50 0.10 0.05 1.00sample=100 0.20 0.10 0.05 1.00 sample=200 0.20 0.10 0.05 OTU=500 OTU=500 OTU=50 OTU=200 OTU=50 OTU=200 OTU number