Relative Abundance by Subject Taxa 1.00 c Clostridia; o Clostridiales c Bacteroidia; o Bacteroidales c__Gammaproteobacteria; o__Pseudomonadales Other c_Spirochaetes; o_Spirochaetales c__Bacilli; o__Bacillales 0.75 c__Bacilli; o__Lactobacillales c__Coriobacteriia; o__Coriobacteriales Root Relative Abundance c__Gammaproteobacteria; o__Enterobacteriales c__Mollicutes; o__RF39 c__Methanobacteria; o__Methanobacteriales c__Erysipelotrichi; o__Erysipelotrichales 0.50 c__Betaproteobacteria; o__Burkholderiales c__Chloroplast; o__Streptophyta c__Flavobacteriia; o__Flavobacteriales c__Gammaproteobacteria; o__Xanthomonadales c__Gammaproteobacteria; o__Aeromonadales c__Verrucomicrobiae; o__Verrucomicrobiales 0.25 p_Proteobacteria; c_Betaproteobacteria c Actinobacteria; o Bifidobacteriales c__RF3; o__ML615J-28 c__Deltaproteobacteria; o__GMD14H09 c__Alphaproteobacteria; o__Caulobacterales c__4C0d-2; o__YS2 0.00 c__Verruco-5; o__WCHB1-41

4h 5f 5h

3h 4f

Subject

3f