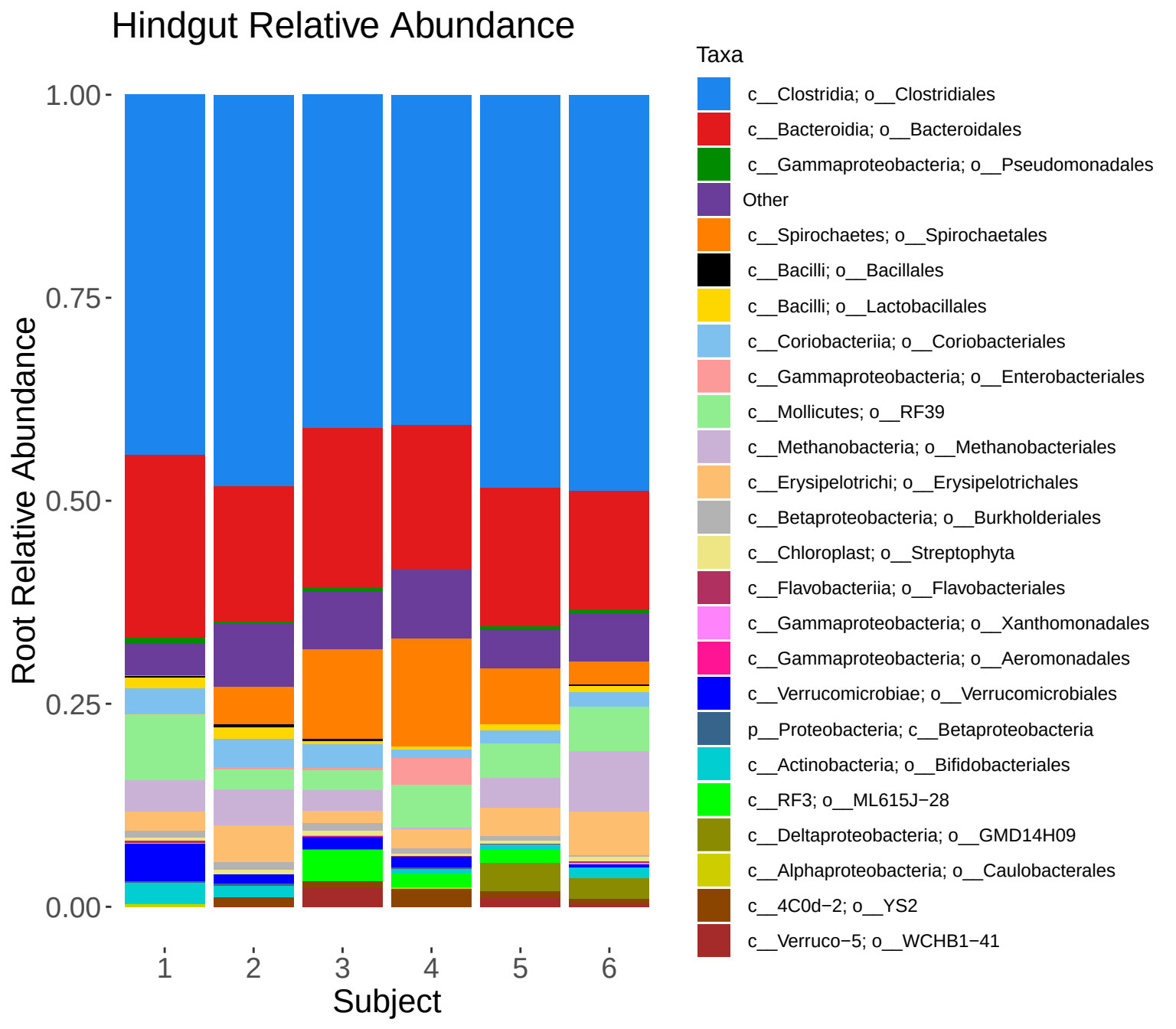
Hindgut Relative Abundance
Root Relative Abundance
1.00
0.75
0.50
0.25
0.00
1
2
3
4
5
6
Subject
Taxa
c__Clostridia; o__Clostridiales
c__Bacteroidia; o__Bacteroidales
c__Gammaproteobacteria; o__Pseudomonadales
Other
c__Spirochaetes; o__Spirochaetales
c__Bacilli; o__Bacillales
c__Bacilli; o__Lactobacillales
c__Coriobacteriia; o__Coriobacteriales
c__Gammaproteobacteria; o__Enterobacteriales
c__Mollicutes; o__RF39
c__Methanobacteria; o__Methanobacteriales
c__Erysipelotrichi; o__Erysipelotrichales
c__Betaproteobacteria; o__Burkholderiales
c__Chloroplast; o__Streptophyta
c__Flavobacteriia; o__Flavobacteriales
c__Gammaproteobacteria; o__Xanthomonadales
c__Gammaproteobacteria; o__Aeromonadales
c__Verrucomicrobiae; o__Verrucomicrobiales
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
c__Verruco−5; o__WCHB1−41