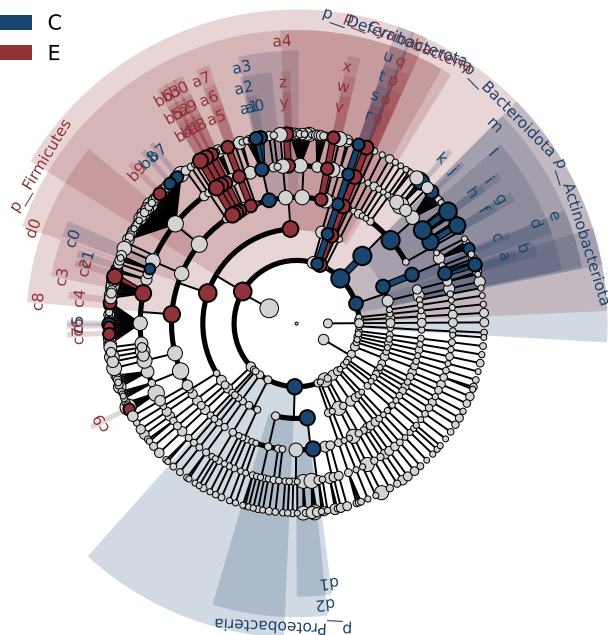


LEfSe Cladogram

C
E



- | | |
|---------------------------------|------------------------------------|
| a : g_CoriobacteriaceaeUCG002 | a4 : c_Bacilli |
| b : f_Atopobiaceae | a5 : c_Clostridia x L6 |
| c : g_Adlercreutzia | a6 : c_Clostridia x L5 |
| d : o_Coriobacteriales | a7 : c_Clostridia x L4 |
| e : c_Coriobacteriia | a8 : g_ClostridiaUCG014 |
| f : g_Bacteroides | a9 : f_Clostridia UCG 014 |
| g : f_Bacteroidaceae | b0 : o_Clostridia UCG 014 |
| h : g_Muribaculaceae | b1 : g_ClostridiavadinBB60group |
| i : f_Muribaculaceae | b2 : f_Clostridia vadinBB60 group |
| j : g_PrevotellaceaeNK3B31group | b3 : o_Clostridia vadinBB60 group |
| k : g_Parabacteroides | b4 : g_Clostridiumsensustricto1 |
| l : o_Bacteroidales | b5 : f_Clostridiaceae |
| m : c_Bacteroidia | b6 : o_Clostridiales |
| n : g_Gastranaerophilales | b7 : g_ASF356 |
| o : f_Gastranaerophilales | b8 : g_Blautia |
| p : o_Gastranaerophilales | b9 : g_GCA900066575 |
| q : c_Vampirivibrionia | c0 : f_Butyricicoccaceae |
| r : g_Mucispirillum | c1 : f_Hungateiclostridiaceae x L6 |
| s : f_Deferribacteraceae | c2 : f_Oscillospiraceae x L6 |
| t : o_Deferribacterales | c3 : f_Oscillospiraceae |
| u : c_Deferribacteres | c4 : f_Ruminococcaceae x L6 |
| v : g_Bacillus | c5 : g_Negativibacillus |
| w : f_Bacillaceae | c6 : g_Paludicola |
| x : o_Bacillales | c7 : g_Ruminococcus |
| y : o_Erysipelotrichales x L6 | c8 : o_Oscillospirales |
| z : o_Erysipelotrichales x L5 | c9 : g_Terrisporobacter |
| a0 : g_HT002 | d0 : c_Clostridia |
| a1 : g_Lactobacillus | d1 : o_Enterobacteriales |
| a2 : f_Lactobacillaceae | d2 : c_Gammaproteobacteria |
| a3 : o_Lactobacillales | |