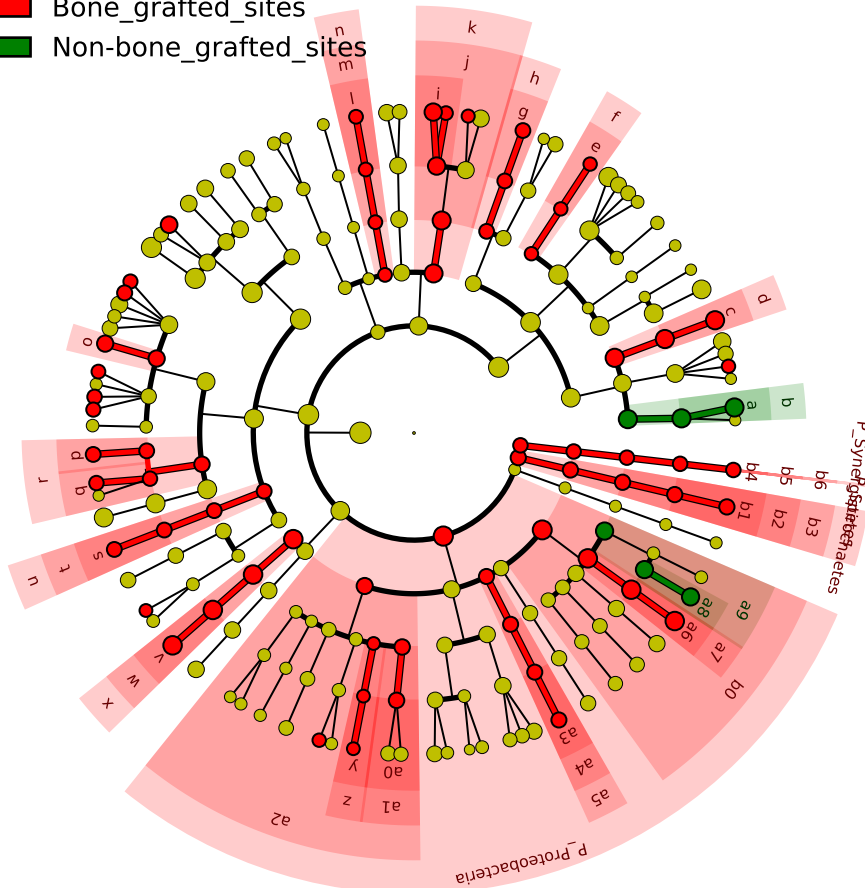


Significantly Represented Taxa

■ Bone_grafted_sites
■ Non-bone_grafted_sites



- a: F_Actinomycetaceae
- b: O_Actinomycetales
- c: F_Propionibacteriaceae
- d: O_Propionibacteriales
- e: F_Pseudonocardiaceae
- f: O_Pseudonocardiales
- g: F_Eggerthellaceae
- h: O_Eggerthellales
- i: F_Prevotellaceae
- j: O_Bacteroidales
- k: C_Bacteroidia
- l: F_Bacteroidaceae
- m: O_Bacteroidales
- n: C_Bacteroidia
- o: F_Peptoniphilaceae
- p: F_Eubacteriaceae
- q: F_Peptostreptococcaceae
- r: O_Eubacteriales
- s: F_Erysipelotrichaceae
- t: O_Erysipelotrichales
- u: C_Erysipelotrichia
- v: F_Fusobacteriaceae
- w: O_Fusobacteriales
- x: C_Fusobacteria
- y: F_Geminicoccaceae
- z: O_Rhodospirillales
- a0: F_Sphingomonadaceae
- a1: O_Sphingomonadales
- a2: C_Alphaproteobacteria
- a3: F_Desulfovibrionaceae
- a4: O_Desulfovibrionales
- a5: C_Deltaproteobacteria
- a6: F_Pasteurellaceae
- a7: O_Pasteurellales
- a8: F_Moraxellaceae
- a9: O_Pseudomonadales
- b0: C_Gammaproteobacteria
- b1: F_Treponemataceae
- b2: O_Spirochaetales
- b3: C_Spirochaetia
- b4: F_Synergistaceae
- b5: O_Synergistales
- b6: C_Synergistia