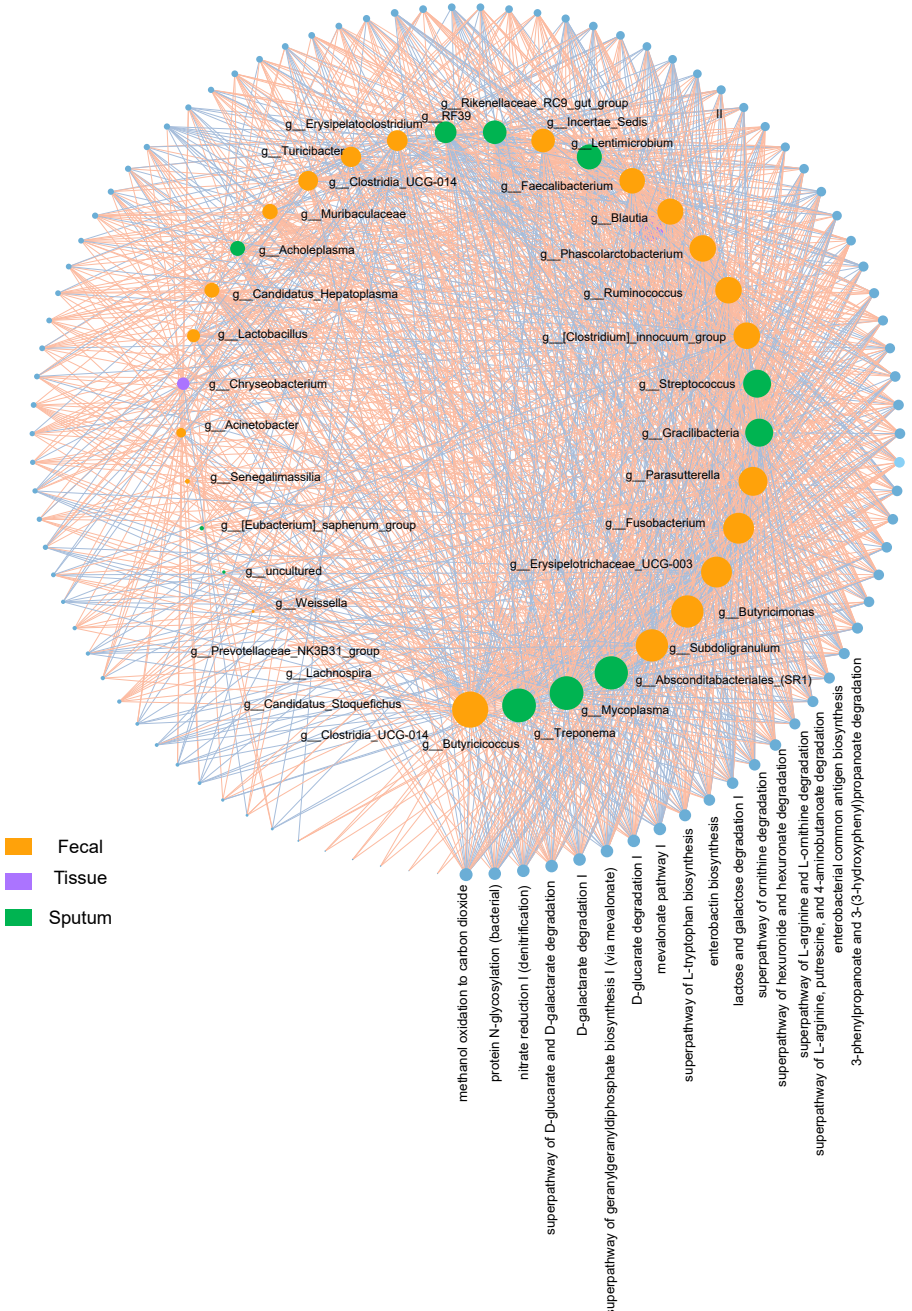
A


g__Rikenellaceae_RC9_gut_group
g__RF39
g__Erysipelatoclostridium
g__Incertae_Sedis
g__Lentimicrobium
g__Turicibacter
g__Clostridia_UCG-014
g__Faecalibacterium
g__Muribaculaceae
g__Blautia
g__Acholeplasma
g__Phascolarctobacterium
g__Candidatus_Hepatoplasma
g__Ruminococcus
g__Lactobacillus
g__[Clostridium]_innocuum_group
g__Chryseobacterium
g__Streptococcus
g__Acinetobacter
g__Gracilibacteria
g__Senegalimassilia
g__Parasutterella
g__[Eubacterium]_saphenum_group
g__Fusobacterium
g__uncultured
g__Erysipelotrichaceae_UCG-003
g__Weissella
g__Butyricimonas
g__Prevotellaceae_NK3B31_group
g__Subdoligranulum
g__Lachnospira
g__Absconditabacteriales_(SR1)
g__Candidatus_Stoquefichus
g__Mycoplasma
g__Clostridia_UCG-014
g__Treponema
g__Butyricicoccus
Fecal
Tissue
Sputum
methanol oxidation to carbon dioxide
protein N-glycosylation (bacterial)
nitrate reduction I (denitrification)
superpathway of D-glucarate and D-galactarate degradation
D-galactarate degradation I
superpathway of geranylgeranyldiphosphate biosynthesis I (via mevalonate)
D-glucarate degradation I
mevalonate pathway I
superpathway of L-tryptophan biosynthesis
enterobactin biosynthesis
lactose and galactose degradation I
superpathway of ornithine degradation
superpathway of hexuronide and hexuronate degradation
superpathway of L-arginine and L-ornithine degradation
superpathway of L-arginine, putrescine, and 4-aminobutanoate degradation
enterobacterial common antigen biosynthesis
3-phenylpropanoate and 3-(3-hydroxyphenyl)propanoate degradation