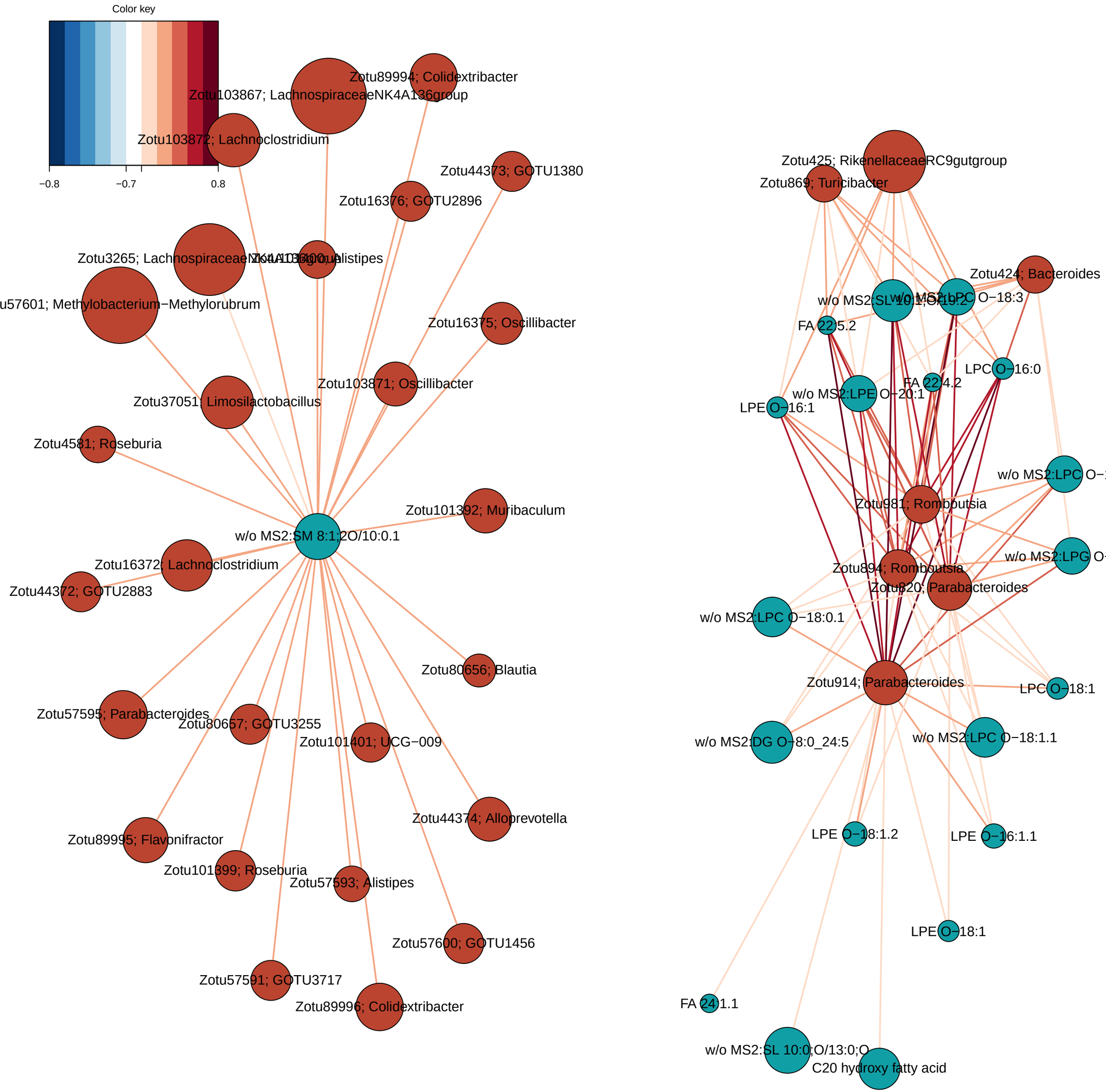
Color key
−0.8
−0.7
0.8
Zotu89994; Colidextribacter
Zotu103867; LachnospiraceaeNK4A136group
Zotu103872; Lachnoclostridium
Zotu44373; GOTU1380
Zotu16376; GOTU2896
Zotu3265; LachnospiraceaeNK4A136group
Zotu103400; Alistipes
u57601; Methylobacterium−Methylorubrum
Zotu16375; Oscillibacter
Zotu103871; Oscillibacter
Zotu37051; Limosilactobacillus
Zotu4581; Roseburia
Zotu101392; Muribaculum
w/o MS2:SM 8:1;2O/10:0.1
Zotu16372; Lachnoclostridium
Zotu44372; GOTU2883
Zotu80656; Blautia
Zotu57595; Parabacteroides
Zotu80657; GOTU3255
Zotu101401; UCG−009
Zotu44374; Alloprevotella
Zotu89995; Flavonifractor
Zotu101399; Roseburia
Zotu57593; Alistipes
Zotu57600; GOTU1456
Zotu57591; GOTU3717
Zotu89996; Colidextribacter
Zotu425; RikenellaceaeRC9gutgroup
Zotu869; Turicibacter
Zotu424; Bacteroides
w/o MS2:SL 10:1;O/19:2
w/o MS2:LPC O−18:3
FA 22:5.2
LPC O−16:0
FA 22:4.2
w/o MS2:LPE O−20:1
LPE O−16:1
w/o MS2:LPC O−18:0
Zotu981; Romboutsia
w/o MS2:LPG O−16:3
Zotu894; Romboutsia
Zotu820; Parabacteroides
w/o MS2:LPC O−18:0.1
Zotu914; Parabacteroides
LPC O−18:1
w/o MS2:DG O−8:0_24:5
w/o MS2:LPC O−18:1.1
LPE O−18:1.2
LPE O−16:1.1
LPE O−18:1
FA 24:1.1
w/o MS2:SL 10:0;O/13:0;O
C20 hydroxy fatty acid