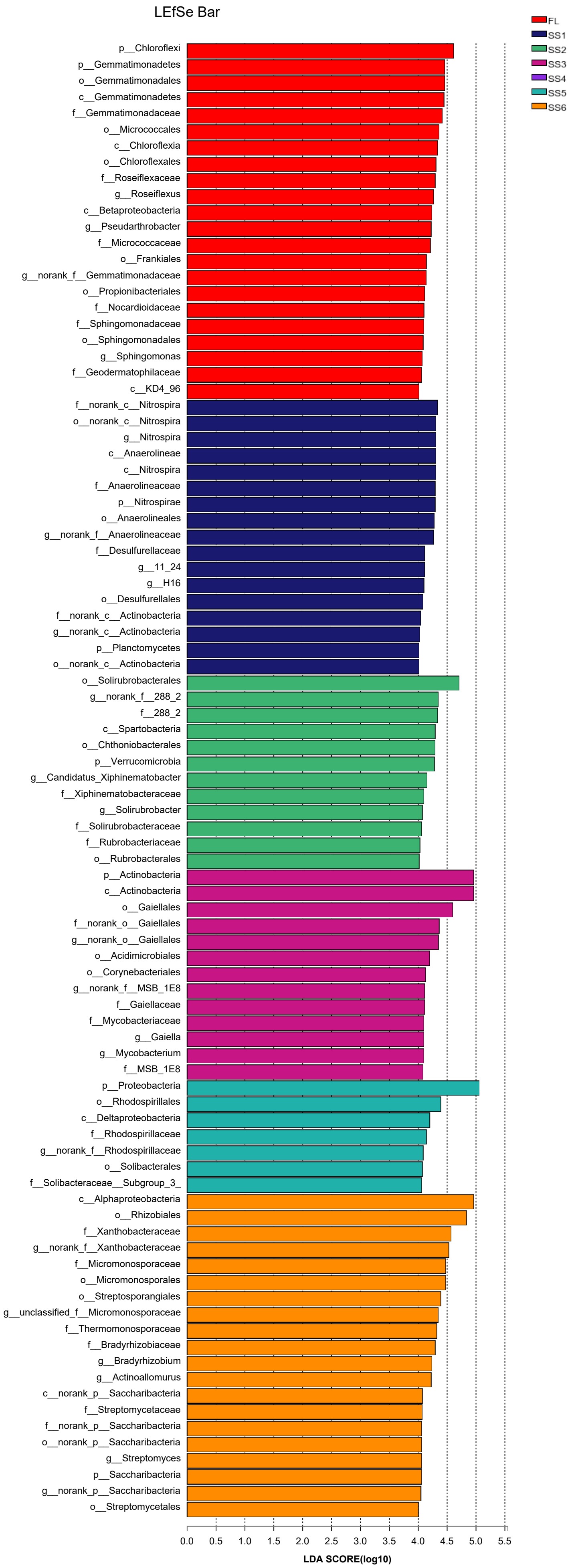
LEfSe Bar
FL
SS1
SS2
SS3
SS4
SS5
SS6
p__Chloroflexi
p__Gemmatimonadetes
o__Gemmatimonadales
c__Gemmatimonadetes
f__Gemmatimonadaceae
o__Micrococcales
c__Chloroflexia
o__Chloroflexales
f__Roseiflexaceae
g__Roseiflexus
c__Betaproteobacteria
g__Pseudarthrobacter
f__Micrococcaceae
o__Frankiales
g__norank_f__Gemmatimonadaceae
o__Propionibacteriales
f__Nocardioidaceae
f__Sphingomonadaceae
o__Sphingomonadales
g__Sphingomonas
f__Geodermatophilaceae
c__KD4_96
f__norank_c__Nitrospira
o__norank_c__Nitrospira
g__Nitrospira
c__Anaerolineae
c__Nitrospira
f__Anaerolineaceae
p__Nitrospirae
o__Anaerolineales
g__norank_f__Anaerolineaceae
f__Desulfurellaceae
g__11_24
g__H16
o__Desulfurellales
f__norank_c__Actinobacteria
g__norank_c__Actinobacteria
p__Planctomycetes
o__norank_c__Actinobacteria
o__Solirubrobacterales
g__norank_f__288_2
f__288_2
c__Spartobacteria
o__Chthoniobacterales
p__Verrucomicrobia
g__Candidatus_Xiphinematobacter
f__Xiphinematobacteraceae
g__Solirubrobacter
f__Solirubrobacteraceae
f__Rubrobacteriaceae
o__Rubrobacterales
p__Actinobacteria
c__Actinobacteria
o__Gaiellales
f__norank_o__Gaiellales
g__norank_o__Gaiellales
o__Acidimicrobiales
o__Corynebacteriales
g__norank_f__MSB_1E8
f__Gaiellaceae
f__Mycobacteriaceae
g__Gaiella
g__Mycobacterium
f__MSB_1E8
p__Proteobacteria
o__Rhodospirillales
c__Deltaproteobacteria
f__Rhodospirillaceae
g__norank_f__Rhodospirillaceae
o__Solibacterales
f__Solibacteraceae__Subgroup_3_
c__Alphaproteobacteria
o__Rhizobiales
f__Xanthobacteraceae
g__norank_f__Xanthobacteraceae
f__Micromonosporaceae
o__Micromonosporales
o__Streptosporangiales
g__unclassified_f__Micromonosporaceae
f__Thermomonosporaceae
f__Bradyrhizobiaceae
g__Bradyrhizobium
g__Actinoallomurus
c__norank_p__Saccharibacteria
f__Streptomycetaceae
f__norank_p__Saccharibacteria
o__norank_p__Saccharibacteria
g__Streptomyces
p__Saccharibacteria
g__norank_p__Saccharibacteria
o__Streptomycetales
0.0
0.5
1.0
1.5
2.0
2.5
3.0
3.5
4.0
4.5
5.0
5.5
LDA SCORE(log10)