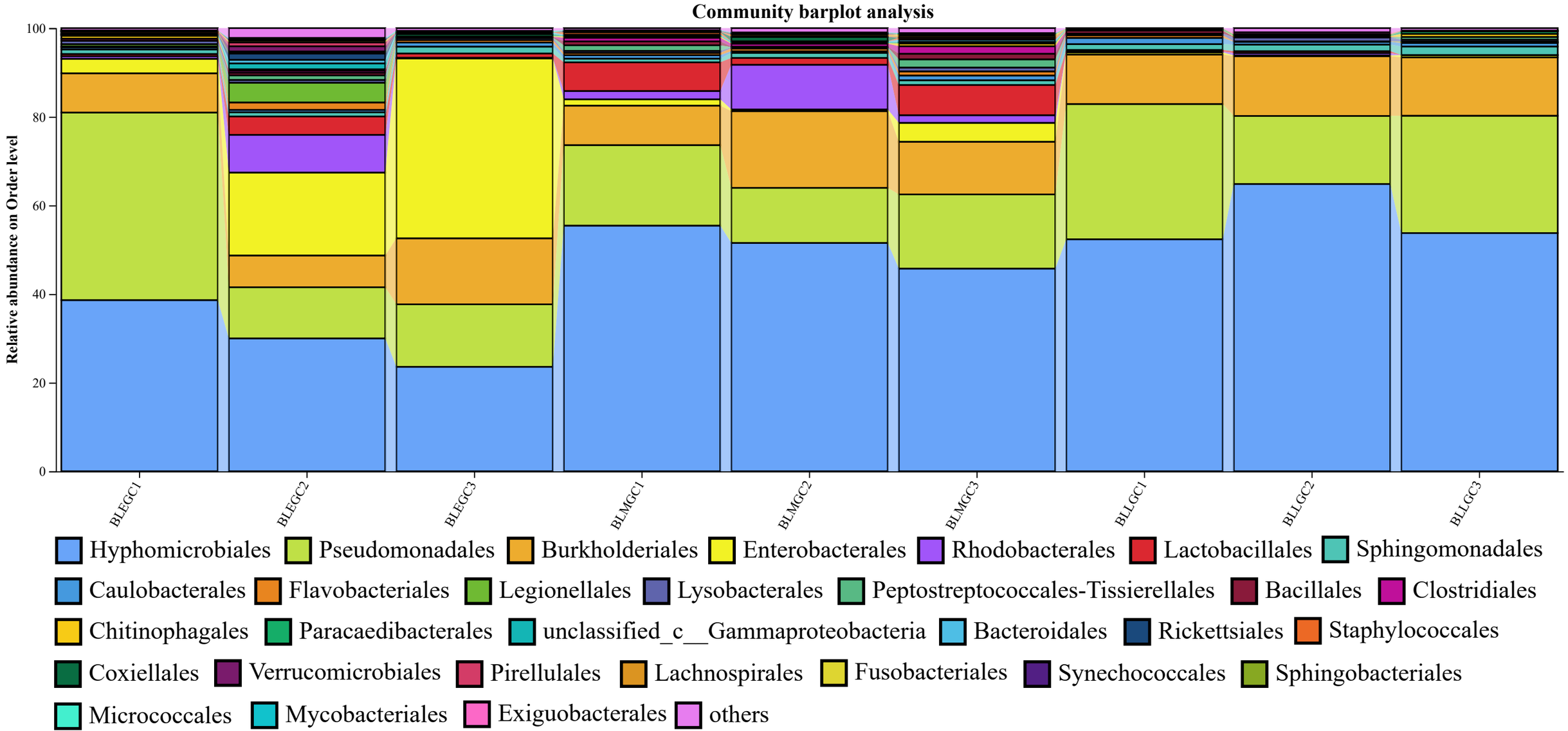

Community barplot analysis
Relative abundance on Order level
100
80
60
40
20
0
BLEGC1
BLEGC2
BLEGC3
BLMGC1
BLMGC2
BLMGC3
BLLGC1
BLLGC2
BLLGC3
Hyphomicrobiales
Pseudomonadales
Burkholderiales
Enterobacterales
Rhodobacterales
Lactobacillales
Sphingomonadales
Caulobacterales
Flavobacteriales
Legionellales
Lysobacterales
Peptostreptococcales-Tissierellales
Bacillales
Clostridiales
Chitinophagales
Paracaedibacterales
unclassified_c__Gammaproteobacteria
Bacteroidales
Rickettsiales
Staphylococcales
Coxiellales
Verrucomicrobiales
Pirellulales
Lachnospirales
Fusobacteriales
Synechococcales
Sphingobacteriales
Micrococcales
Mycobacteriales
Exiguobacterales
others