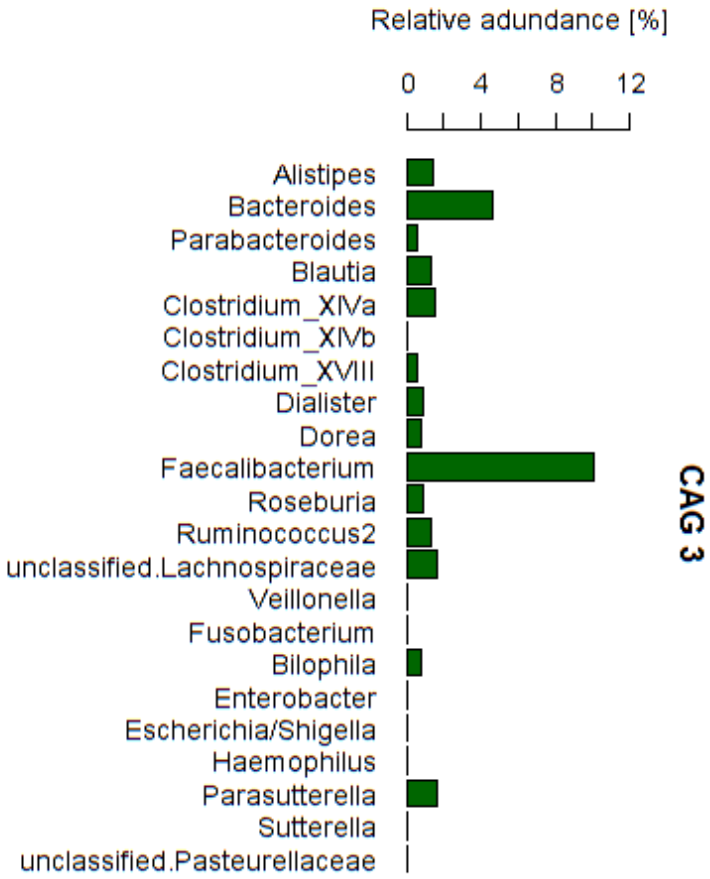
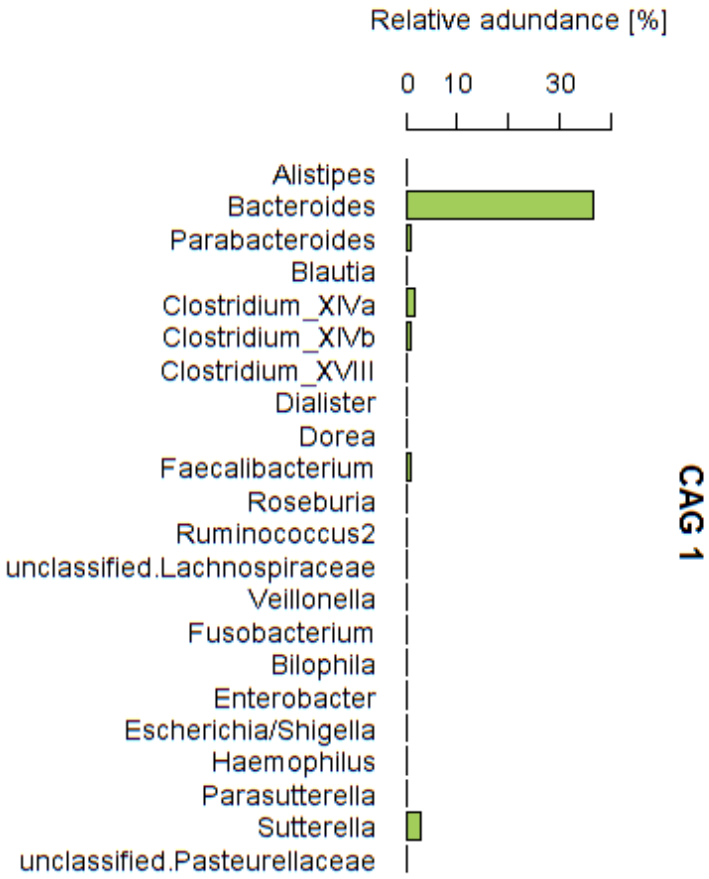
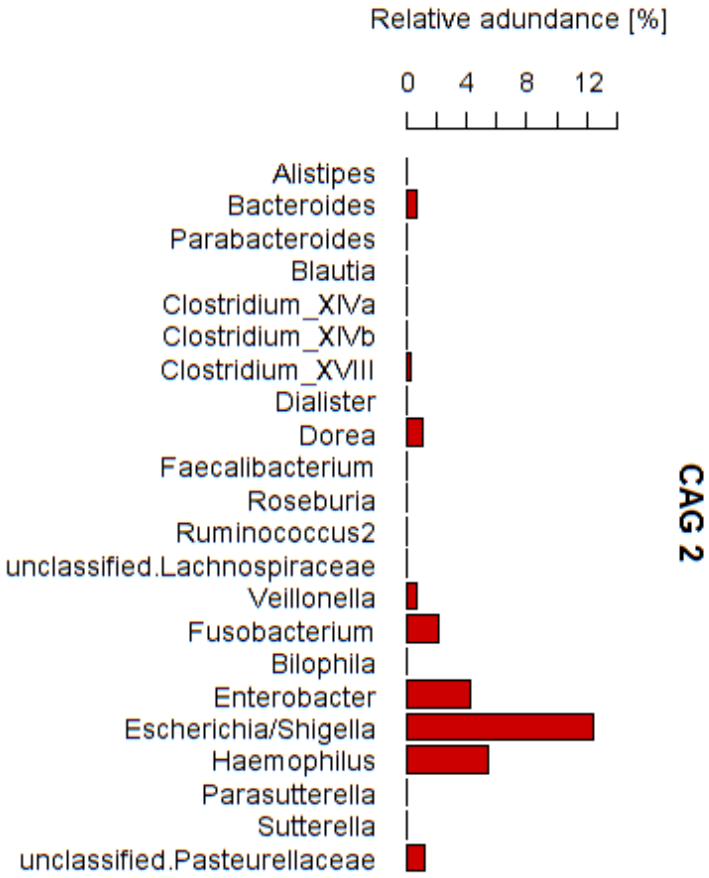

CAG 3
Relative adundance [%]
0
4
8
12
Alistipes
Bacteroides
Parabacteroides
Blautia
Clostridium_XIVa
Clostridium_XIVb
Clostridium_XVIII
Dialister
Dorea
Faecalibacterium
Roseburia
Ruminococcus2
unclassified.Lachnospiraceae
Veillonella
Fusobacterium
Bilophila
Enterobacter
Escherichia/Shigella
Haemophilus
Parasutterella
Sutterella
unclassified.Pasteurellaceae


CAG 1
Relative adundance [%]
0
10
30
Alistipes
Bacteroides
Parabacteroides
Blautia
Clostridium_XIVa
Clostridium_XIVb
Clostridium_XVIII
Dialister
Dorea
Faecalibacterium
Roseburia
Ruminococcus2
unclassified.Lachnospiraceae
Veillonella
Fusobacterium
Bilophila
Enterobacter
Escherichia/Shigella
Haemophilus
Parasutterella
Sutterella
unclassified.Pasteurellaceae


CAG 2
Relative adundance [%]
0
4
8
12
Alistipes
Bacteroides
Parabacteroides
Blautia
Clostridium_XIVa
Clostridium_XIVb
Clostridium_XVIII
Dialister
Dorea
Faecalibacterium
Roseburia
Ruminococcus2
unclassified.Lachnospiraceae
Veillonella
Fusobacterium
Bilophila
Enterobacter
Escherichia/Shigella
Haemophilus
Parasutterella
Sutterella
unclassified.Pasteurellaceae