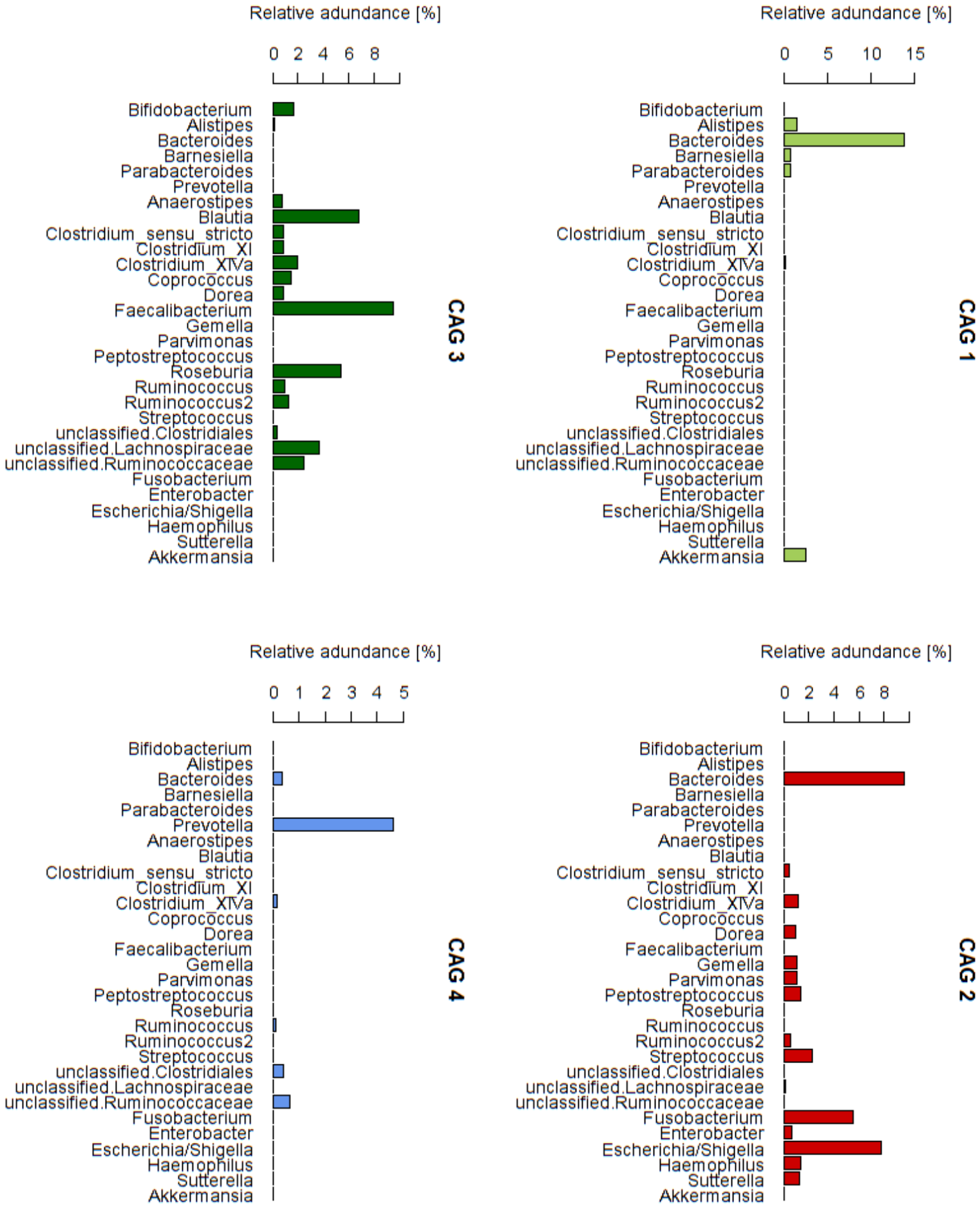
CAG 1
Relative adundance [%]
0 5 10 15
CAG 2
Relative adundance [%]
0 2 4 6 8
CAG 3
Relative adundance [%]
0 2 4 6 8
CAG 4
Relative adundance [%]
0 1 2 3 4 5
Bifidobacterium
Alistipes
Bacteroides
Barnesiella
Parabacteroides
Prevotella
Anaerostipes
Blautia
Clostridium_sensu_stricto
Clostridium_XI
Clostridium_XIVa
Coprococcus
Dorea
Faecalibacterium
Gemella
Parvimonas
Peptostreptococcus
Roseburia
Ruminococcus
Ruminococcus2
Streptococcus
unclassified.Clostridiales
unclassified.Lachnospiraceae
unclassified.Ruminococcaceae
Fusobacterium
Enterobacter
Escherichia/Shigella
Haemophilus
Sutterella
Akkermansia