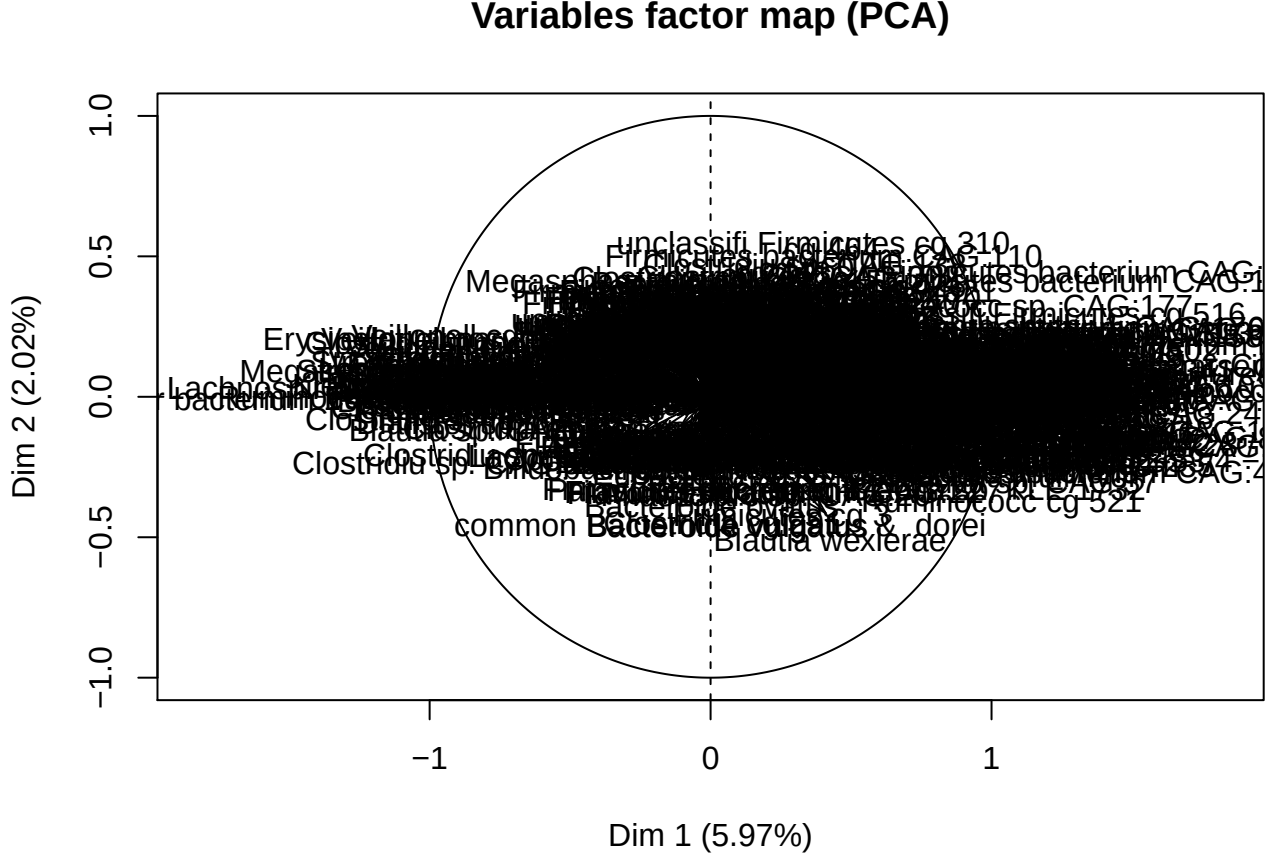
Variables factor map (PCA)
Dim 1 (5.97%)
Dim 2 (2.02%)
−1
0
1
−1.0
−0.5
0.0
0.5
1.0
unclassifi Firmicutes co 310
Megasp
Erysipelotrichaceae
Lachnospi
bacterium
Clostridia sp.
Blautia sp.
Clostridium
common
Bacteroides vulgatus & dorei
Blautia wexlerae
Firmicutes bacterium CAG:110
Firmicutes bacterium CAG:1
CAG:177
Firmicutes co 516
Ruminococc cg 321
CAG:2