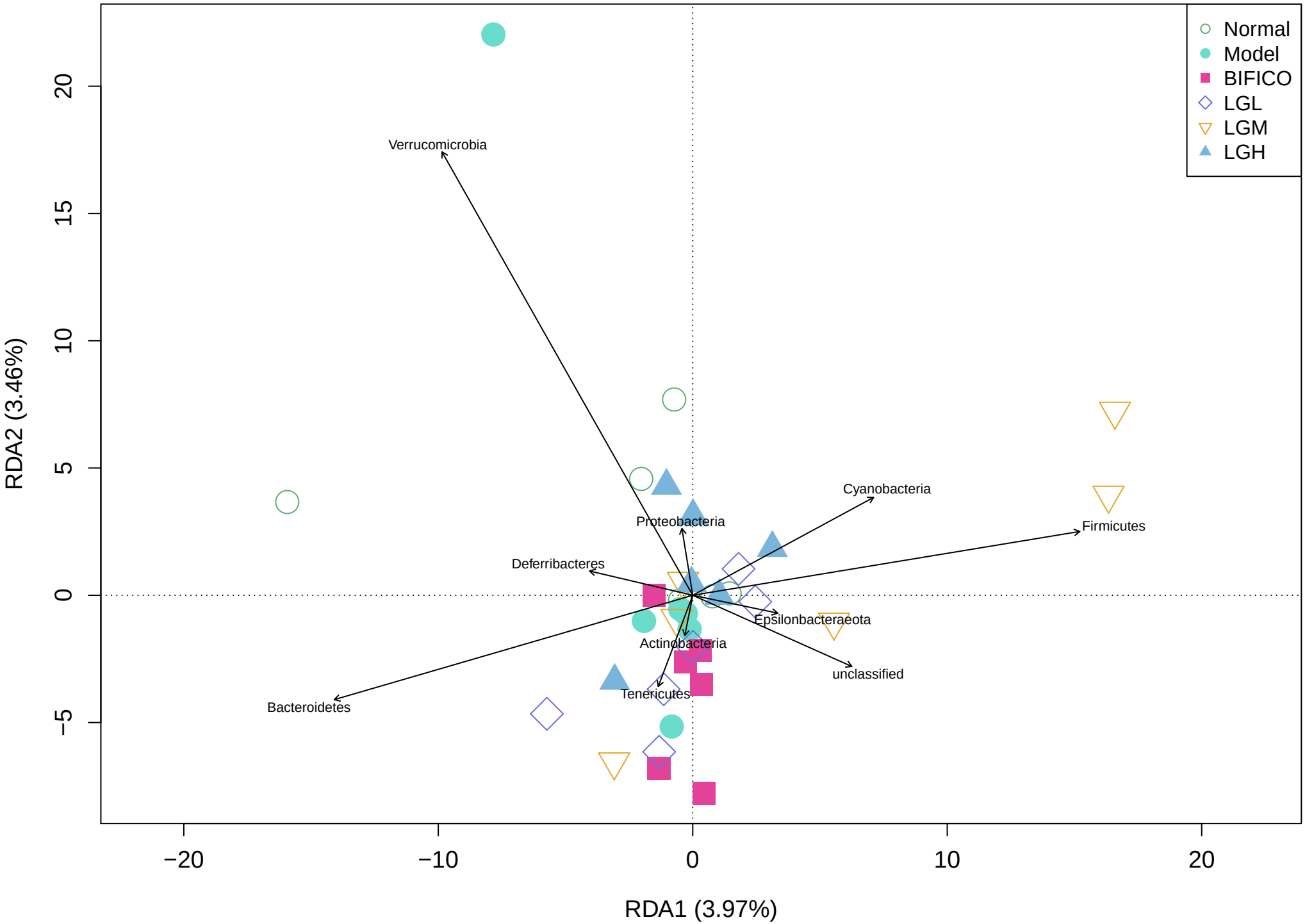
Normal
Model
BIFICO
LGL
LGM
LGH
Verrucomicrobia
Cyanobacteria
Firmicutes
Proteobacteria
Deferribacteres
Epsilonbacteraeota
Actinobacteria
unclassified
Tenericutes
Bacteroidetes
RDA2 (3.46%)
RDA1 (3.97%)
20
15
10
5
0
−5
−20
−10
0
10
20