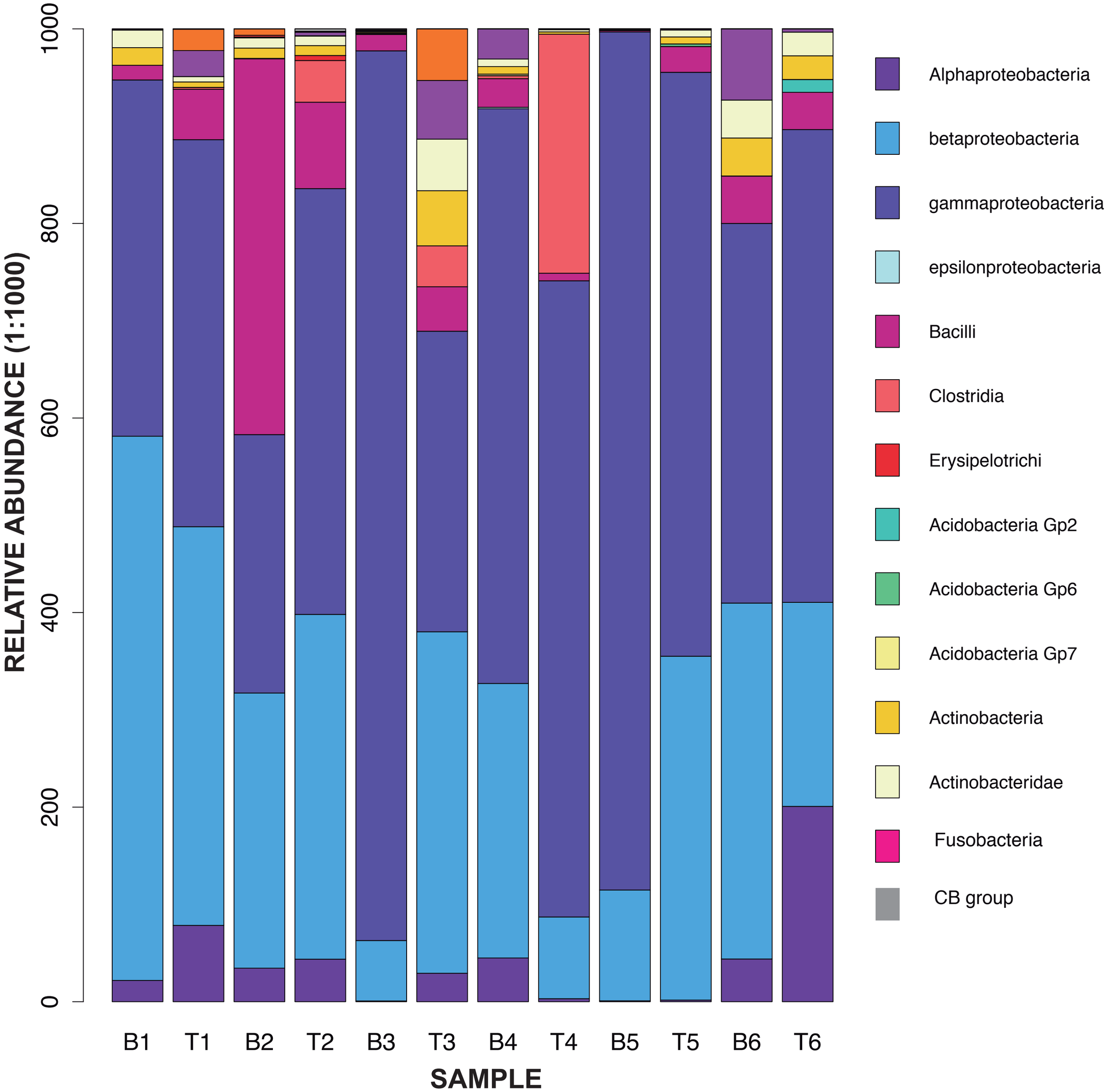
RELATIVE ABUNDANCE (1:1000)
0
200
400
600
800
1000
B1
T1
B2
T2
B3
T3
B4
T4
B5
T5
B6
T6
SAMPLE
Alphaproteobacteria
betaproteobacteria
gammaproteobacteria
epsilonproteobacteria
Bacilli
Clostridia
Erysipelotrichi
Acidobacteria Gp2
Acidobacteria Gp6
Acidobacteria Gp7
Actinobacteria
Actinobacteridae
Fusobacteria
CB group