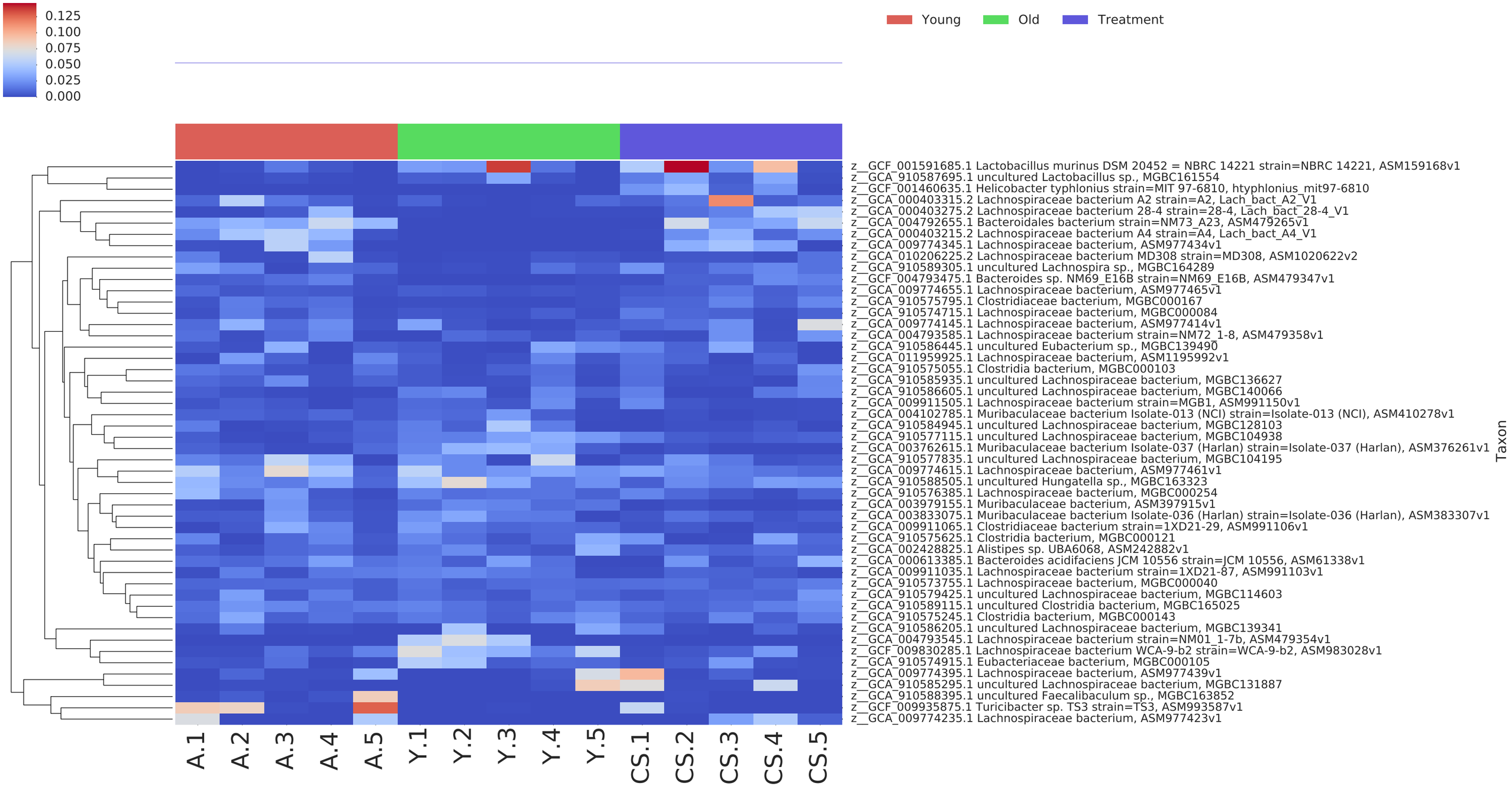
0.125
0.100
0.075
0.050
0.025
0.000
Young
Old
Treatment
z__GCF_001591685.1 Lactobacillus murinus DSM 20452 = NBRC 14221 strain=NBRC 14221, ASM159168v1
z__GCA_910587695.1 uncultured Lactobacillus sp., MGBC161554
z__GCF_001460635.1 Helicobacter typhlonius strain=MIT 97-6810, htyphlonius_mit97-6810
z__GCA_000403315.2 Lachnospiraceae bacterium A2 strain=A2, Lach_bact_A2_V1
z__GCA_000403275.2 Lachnospiraceae bacterium 28-4 strain=28-4, Lach_bact_28-4_V1
z__GCA_004792655.1 Bacteroidales bacterium strain=NM73_A23, ASM479265v1
z__GCA_000403215.2 Lachnospiraceae bacterium A4 strain=A4, Lach_bact_A4_V1
z__GCA_009774345.1 Lachnospiraceae bacterium, ASM977434v1
z__GCA_010206225.2 Lachnospiraceae bacterium MD308 strain=MD308, ASM1020622v2
z__GCA_910589305.1 uncultured Lachnospira sp., MGBC164289
z__GCF_004793475.1 Bacteroides sp. NM69_E16B strain=NM69_E16B, ASM479347v1
z__GCA_009774655.1 Lachnospiraceae bacterium, ASM977465v1
z__GCA_910575795.1 Clostridiaceae bacterium, MGBC000167
z__GCA_910574715.1 Lachnospiraceae bacterium, MGBC000084
z__GCA_009774145.1 Lachnospiraceae bacterium, ASM977414v1
z__GCA_004793585.1 Lachnospiraceae bacterium strain=NM72_1-8, ASM479358v1
z__GCA_910586445.1 uncultured Eubacterium sp., MGBC139490
z__GCA_011959925.1 Lachnospiraceae bacterium, ASM1195992v1
z__GCA_910575055.1 Clostridia bacterium, MGBC000103
z__GCA_910585935.1 uncultured Lachnospiraceae bacterium, MGBC136627
z__GCA_910586605.1 uncultured Lachnospiraceae bacterium, MGBC140066
z__GCA_009911505.1 Lachnospiraceae bacterium strain=MGB1, ASM991150v1
z__GCA_004102785.1 Muribaculaceae bacterium Isolate-013 (NCI) strain=Isolate-013 (NCI), ASM410278v1
z__GCA_910584945.1 uncultured Lachnospiraceae bacterium, MGBC128103
z__GCA_910577115.1 uncultured Lachnospiraceae bacterium, MGBC104938
z__GCA_003762615.1 Muribaculaceae bacterium Isolate-037 (Harlan) strain=Isolate-037 (Harlan), ASM376261v1
z__GCA_910577835.1 uncultured Lachnospiraceae bacterium, MGBC104195
z__GCA_009774615.1 Lachnospiraceae bacterium, ASM977461v1
z__GCA_910588505.1 uncultured Hungatella sp., MGBC163323
z__GCA_910576385.1 Lachnospiraceae bacterium, MGBC000254
z__GCA_003979155.1 Muribaculaceae bacterium, ASM397915v1
z__GCA_003833075.1 Muribaculaceae bacterium Isolate-036 (Harlan) strain=Isolate-036 (Harlan), ASM383307v1
z__GCA_009911065.1 Clostridiaceae bacterium strain=1XD21-29, ASM991106v1
z__GCA_910575625.1 Clostridia bacterium, MGBC000121
z__GCA_002428825.1 Alistipes sp. UBA6068, ASM242882v1
z__GCA_000613385.1 Bacteroides acidifaciens JCM 10556 strain=JCM 10556, ASM61338v1
z__GCA_009911035.1 Lachnospiraceae bacterium strain=1XD21-87, ASM991103v1
z__GCA_910573755.1 Lachnospiraceae bacterium, MGBC000040
z__GCA_910579425.1 uncultured Lachnospiraceae bacterium, MGBC114603
z__GCA_910589115.1 uncultured Clostridia bacterium, MGBC165025
z__GCA_910575245.1 Clostridia bacterium, MGBC000143
z__GCA_910586205.1 uncultured Lachnospiraceae bacterium, MGBC139341
z__GCA_004793545.1 Lachnospiraceae bacterium strain=NM01_1-7b, ASM479354v1
z__GCF_009830285.1 Lachnospiraceae bacterium WCA-9-b2 strain=WCA-9-b2, ASM983028v1
z__GCA_910574915.1 Eubacteriaceae bacterium, MGBC000105
z__GCA_009774395.1 Lachnospiraceae bacterium, ASM977439v1
z__GCA_910585295.1 uncultured Lachnospiraceae bacterium, MGBC131887
z__GCA_910588395.1 uncultured Faecalibaculum sp., MGBC163852
z__GCF_009935875.1 Turicibacter sp. TS3 strain=TS3, ASM993587v1
z__GCA_009774235.1 Lachnospiraceae bacterium, ASM977423v1
Taxon
A.1
A.2
A.3
A.4
A.5
Y.1
Y.2
Y.3
Y.4
Y.5
CS.1
CS.2
CS.3
CS.4
CS.5