Summarizing heatmap FDR-values: < 0.001 = ***, < 0.01 = **, < 0.1 = * ASV37 [Eubacterium] ruminantium group
ASV37 [Eubacterium] ruminantium group
ASV351 Lachnoclostridium
ASV351 Lachnoclostridium
ASV361 Clostridiu UCG-014
ASV429 Lachnospiraceae
ASV445 Pugmaiobacter
ASV345 Paludicolostrideae
ASV45 Paludicolostrideae
ASV45 Paludicolostridium ASV28 [Eubacterium] corpostanoligenes group
ASV121 [Eubacterium] corpostanoligenes group
ASV131 [Eubacterium] corpostanoligenes group
ASV368 Clostridia vadinBB60 group
ASV368 Clostridia vadinBB60 group
ASV368 Clostridia vadinBB60 group
ASV200 Intestiminonas
ASV10 Scillospiraceae
ASV130 [Eubacterium] XVJanophilum group
ASV365 ASV379 [Actinospiraceae
ASV379 [Actinospiraceae
ASV379 [Actinospiraceae
ASV214 Clostrida vadinBB60 group
ASV214 [Costrida vadinBB60 group
ASV214 [ASV355 Lachnospiraceae
ASV214 [ASV355 Lachnospiraceae
ASV214 [ASV355 Lachnospiraceae ASV136 LIBA1818
ASV146 Clost Butty Ticicoccus
ASV13 Lachnospiraceae NK 4A 36 group
ASV38 Lachnospiraceae NK 4A 36 group
ASV38 Lachnospiraceae NK 4A 36 group
ASV39 Anaerotruncus
ASV34 Anaerotruncus
ASV34 Dores
ASV474 Anaerotruncus
ASV34 Dores
ASV474 Colidextribacter
ASV364 Colidextribacter
ASV364 Colidextribacter
ASV182 GCA-900066575
ASV475 Ruminprocyccaceae Ds ASV92 Lachnospiraceae UC-006
ASV404 Colidextribacter
ASV482 GCA-900066575
ASV175 Ruminococaceae
ASV175 Ruminococaceae
ASV271 Asv218 UCG-005
ASV271 Asv218 UCG-016
ASV37 Lachnospiraceae
ASV196 Lachnospiraceae
ASV197 Lachnospiraceae
ASV197 Lachnospiraceae
ASV174 UCG-016
ASV37 Asv218 UCG-016
ASV37 Lachnospiraceae
ASV174 UCG-016
ASV37 Lachnospiraceae NXAA136 group
ASV7 Lachnospiraceae NXAA136 group
ASV176 Colidextribacter
ASV177 Colidextribacter
ASV177 Colidextribacter
ASV177 Colidextribacter
ASV177 Colidextribacter
ASV178 Lachnospiraceae
ASV177 Colidextribacter
ASV178 Lachnospiraceae
ASV177 Colidextribacter
ASV178 Lachnospiraceae NXAA136 group
ASV49 Lachnospiraceae NXAA136 group
ASV414 Clostrida UCG-014
ASV18 Lachnospiraceae NXAA136 group
ASV418 Colidextribacter
ASV118 Colidextribacter
ASV326 Castribacter
ASV326 Castribacter
ASV31 Lachnospiraceae
ASV3 0.5 Omics features 0.0 -0.5-1.0Confounding status confounded deconfounded SCFA_BA Treated_CCL NR2A

Metadata variables