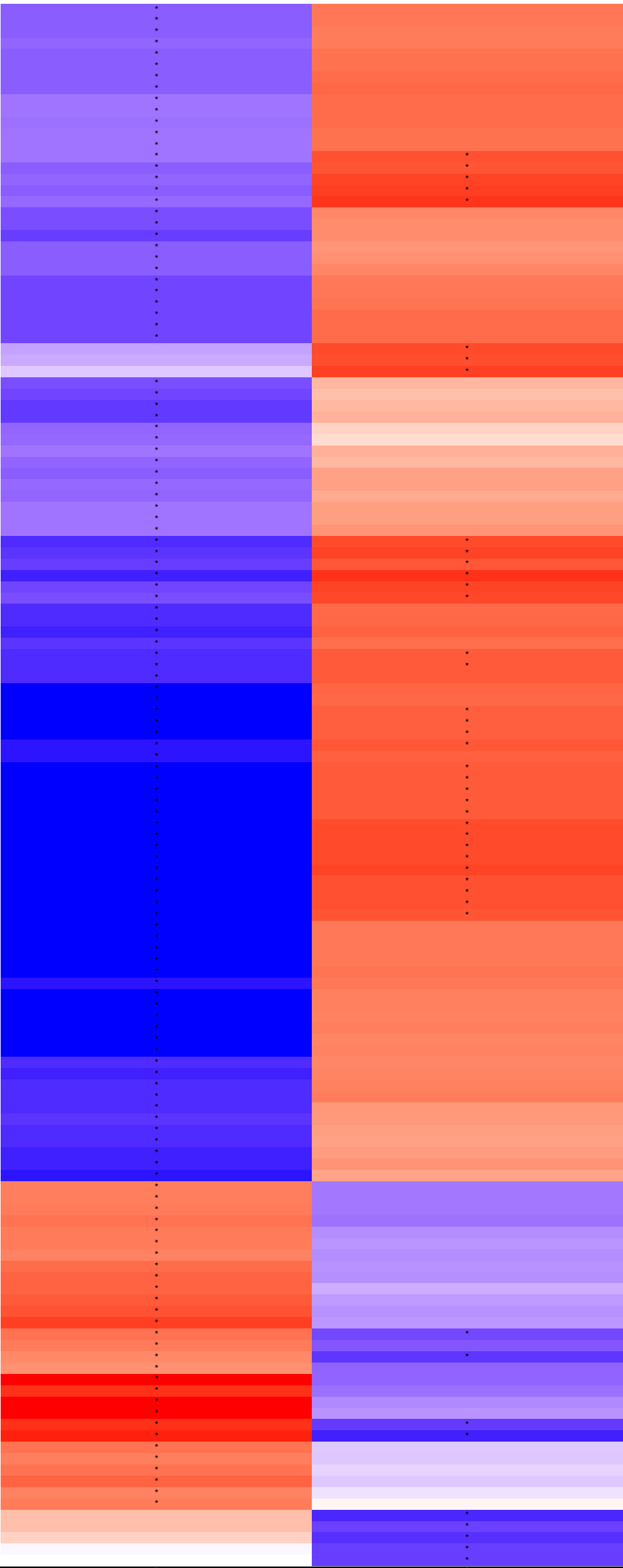


Summarizing heatmap

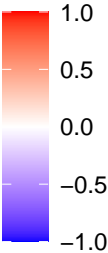
FDR-values: < 0.001 = ***, < 0.01 = **, < 0.1 = *

Omic features

ASV754 Parabacteroides
ASV486 Bacteroides
ASV727 RF39
ASV111 Lachnospiraceae NK4A136 group
ASV447 RF39
ASV154 intestinale
ASV161 Lachnospiraceae NK4A136 group
ASV7 johnsonii
ASV559 obesi
ASV430 bacterium
ASV424 Tuzzerella
ASV402 Enterorhabdus
ASV353 Intestinimonas
ASV605 Clostridia vadinBB60 group
ASV387 massiliensis
ASV526 Clostridia vadinBB60 group
ASV108 Alistipes
ASV28 Jeotgalicoccus
ASV351 [Eubacterium] xylanophilum group
ASV182 Muribaculaceae
ASV361 Christensenellaceae
ASV491 NK4A214 group
ASV173 Colidextribacter
ASV284 Oscillospiraceae
ASV528 cellulossilyticus
ASV135 Muribaculaceae
ASV498 Ruminococcaceae
ASV405 Erysipelotrichaceae
ASV150 lentus
ASV538 A2
ASV588 UCG-005
ASV211 [Eubacterium] coprostanoligenes group
ASV394 Enterorhabdus
ASV286 Rhodospirillales
ASV25 Muribaculaceae
ASV511 thetaiotaomicron
ASV320 Oscillospiraceae
ASV87 Lachnospiraceae
ASV75 Ruminococcus
ASV436 Enterorhabdus
ASV423 UBA1819
ASV427 Lachnoclostridium
ASV403 Oscillospiraceae
ASV589 Muribaculaceae
ASV585 Bilophila
ASV363 Lachnospiraceae
ASV235 Clostridia UCG-014
ASV131 RF39
ASV34 Ruminococcus
ASV51 Colidextribacter
ASV41 [Eubacterium] xylanophilum group
ASV504 Paludicola
ASV157 goldsteinii
ASV313 Paludicola
ASV308 Muribaculaceae
ASV21 Muribaculaceae
ASV191 Colidextribacter
ASV255 Roseburia
ASV102 choerinum
ASV18 Oscillospiraceae
ASV184 Bacteroides
ASV171 Clostridia UCG-014
ASV467 UCG-010
ASV178 Muribaculaceae
ASV517 Ruminococcaceae
ASV198 Colidextribacter
ASV117 Lachnospiraceae NK4A136 group
ASV22 Muribaculaceae
ASV19 Muribaculaceae
ASV63 RF39
ASV163 UCG-010
ASV413 Parasutterella
ASV128 Muribaculaceae
ASV105 uniformis
ASV176 Muribaculaceae
ASV355 Marvinbryantia
ASV227 Ruminococcus
ASV172 Muribaculaceae
ASV168 Muribaculaceae
ASV218 finegoldii
ASV317 Lachnospiraceae
ASV165 caecimuris
ASV58 Muribaculaceae
ASV251 dorei
ASV46 Colidextribacter
ASV169 Oscillospiraceae
ASV181 Alistipes
ASV256 Muribaculaceae
ASV26 Alistipes
ASV80 Lachnospiraceae
ASV208 Muribaculaceae
ASV134 Bacteroides
ASV39 Muribaculaceae
ASV291 Muribaculaceae
ASV213 Muribaculaceae
ASV202 Intestinimonas
ASV194 Lachnospiraceae
ASV155 Muribaculaceae
ASV98 Clostridia UCG-014
ASV381 distasonis
ASV84 Alistipes
ASV271 Lachnospiraceae
ASV32 Muribaculaceae
ASV122 acidifaciens
ASV377 Lachnospiraceae
ASV272 Muribaculaceae
ASV662 [Clostridium] methylpentosum group
ASV4 Lachnospiraceae NK4A136 group
ASV382 Fournierella
ASV344 UCG-005
ASV352 Muribaculaceae
ASV373 GCA-900066575
ASV275 Lachnospiraceae
ASV60 Ruminococcus
ASV269 Muribaculaceae
ASV149 Muribaculaceae
ASV16 pseudolongum
ASV183 Lachnoclostridium
ASV56 Clostridia UCG-014
ASV569 GCA-900066575
ASV239 ilealis
ASV395 Colidextribacter
ASV274 Ruminococcaceae
ASV228 Lachnospiraceae
ASV29 Muribaculaceae
ASV174 Lachnospiraceae
ASV1 Muribaculaceae
ASV282 GCA-900066575
ASV27 Lachnospiraceae NK4A136 group
ASV85 HT002
ASV67 [Eubacterium] coprostanoligenes group
ASV470 UCG-007
ASV47 xylosus
ASV343 Oscillospiraceae
ASV133 Roseburia
ASV383 merdae
ASV8 Lachnospiraceae NK4A136 group
ASV2 Lachnospiraceae NK4A136 group



Ds



Confounding status

- confounded
- * deconfounded

Metadata variables