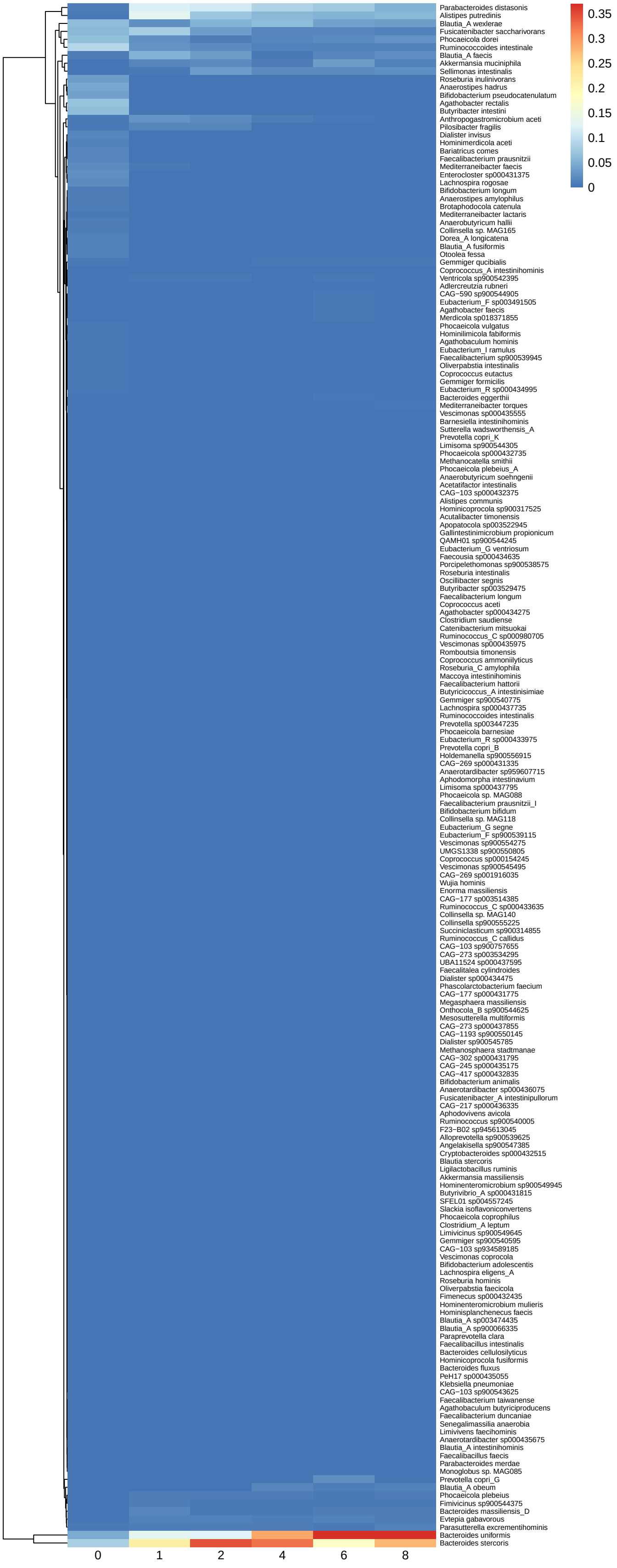
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Phocaeicola dorei
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Parabacteroides merdae
Phocaeicola vulgatus
Acetatifactor intestinalis
Catenibacterium mitsuokai
Lachnospira eligens_A
Clostridium_A leptum
Maccoya intestinihominis
Blautia_A obeum
Roseburia inulinivorans
Mesosutterella multiformis
Coprococcus_A intestinihominis
Lachnospira rogosae
Faecalibacterium prausnitzii
Anthropogastromicrobium aceti
Mediterraneibacter lactaris
Eubacterium_G segne
Agathobaculum hominis
Faecalibacterium sp900539945
Pilosibacter fragilis
Mediterraneibacter faecis
Bariatricus comes
Faecalibacterium prausnitzii_I
Hominimerdicola aceti
Faecalibacterium duncaniae
Romboutsia timonensis
Hominenteromicrobium mulieris
Vescimonas coprocola
CAG−177 sp003514385
Otoolea fessa
Hominisplanchenecus faecis
Vescimonas sp900554275
Enterocloster sp000431375
Agathobaculum butyriciproducens
Roseburia intestinalis
Anaerostipes amylophilus
Limivicinus sp900549645
Anaerotardibacter sp000436075
Faecousia sp000434635
Roseburia hominis
Vescimonas sp000435555
Collinsella sp. MAG118
Faecalibacillus intestinalis
PeH17 sp000435055
Mediterraneibacter torques
Eubacterium_I ramulus
Slackia isoflavoniconvertens
Angelakisella sp900547385
Agathobacter sp000434275
Coprococcus eutactus
Butyrivibrio_A sp000431815
Aphodomorpha intestinavium
SFEL01 sp004557245
Vescimonas sp900545495
Faecalibacterium hattorii
Bifidobacterium pseudocatenulatum
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Eubacterium_F sp003491505
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Faecalibacillus faecis
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Barnesiella intestinihominis
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Klebsiella pneumoniae
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Porcipelethomonas sp900538575
Ligilactobacillus ruminis
Gemmiger formicilis
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Bacteroides massiliensis_D
Faecalitalea cylindroides
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Phocaeicola coprophilus
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Gemmiger qucibialis
Bifidobacterium longum
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Limivivens faecihominis
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Coprococcus sp000154245
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Alloprevotella sp900539625
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Paraprevotella clara
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Sellimonas intestinalis
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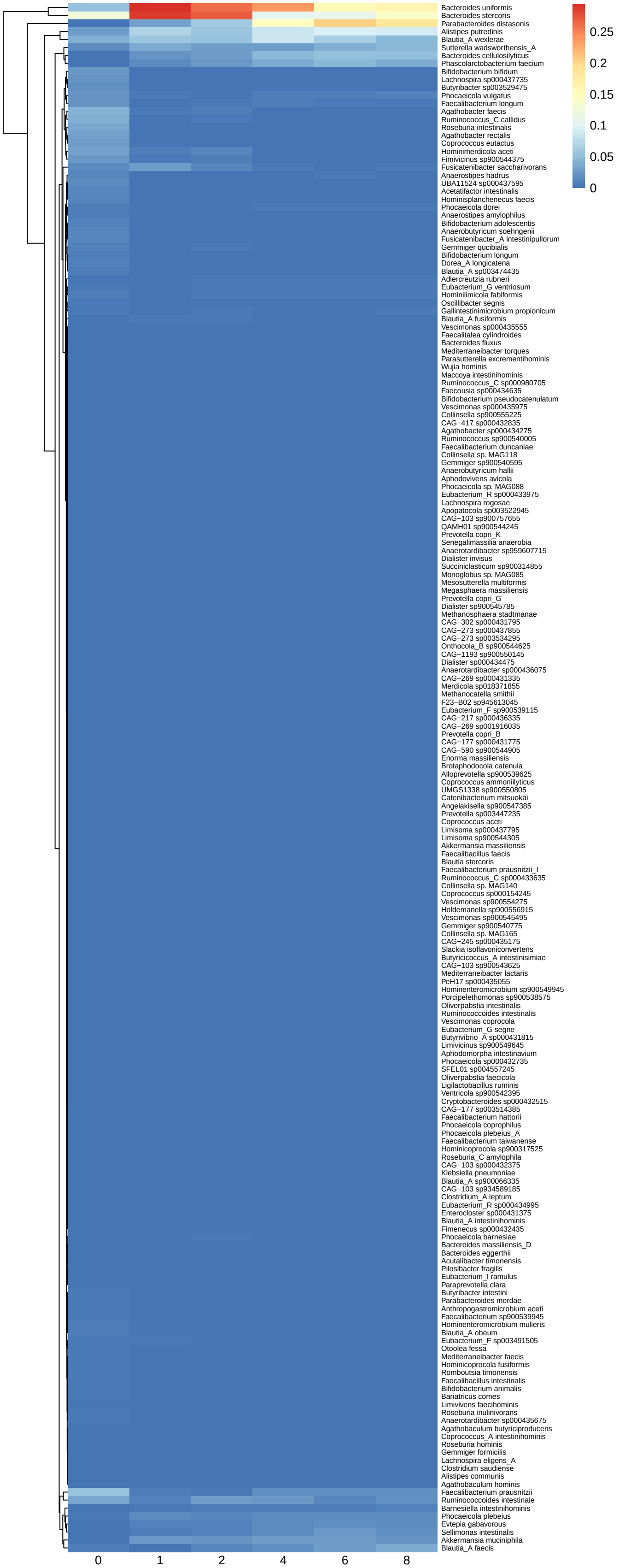
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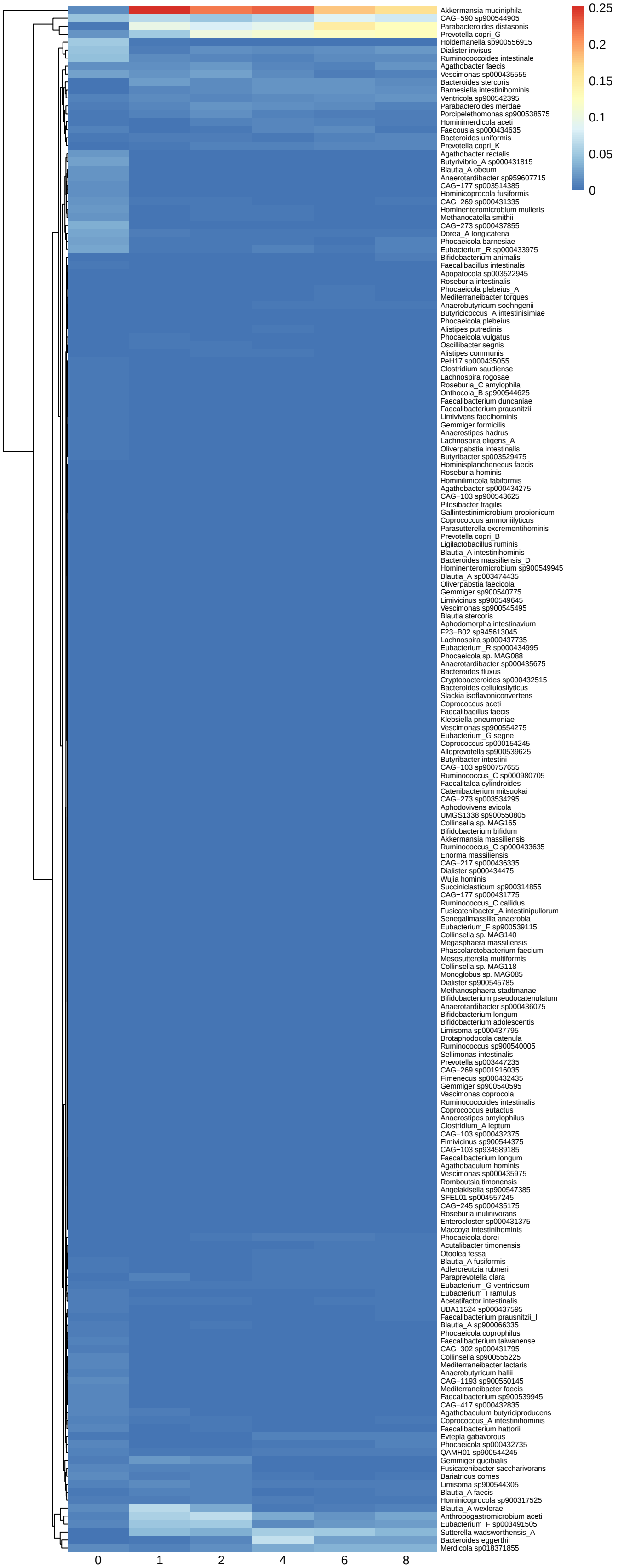
F1


Parabacteroides distasonis
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Blautia_A faecis
Akkermansia muciniphila
Sellimonas intestinalis
Roseburia inulinivorans
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Bifidobacterium pseudocatenulatum
Agathobacter rectalis
Butyribacter intestini
Anthropogastromicrobium aceti
Pilosibacter fragilis
Dialister invisus
Hominimerdicola aceti
Bariatricus comes
Faecalibacterium prausnitzii
Mediterraneibacter faecis
Enterocloster sp000431375
Lachnospira rogosae
Bifidobacterium longum
Anaerostipes amylophilus
Brotaphodocola catenula
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Collinsella sp. MAG165
Dorea_A longicatena
Blautia_A fusiformis
Otoolea fessa
Gemmiger qucibialis
Coprococcus_A intestinihominis
Ventricola sp900542395
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Eubacterium_F sp003491505
Agathobacter faecis
Merdicola sp018371855
Phocaeicola vulgatus
Hominilimicola fabiformis
Agathobaculum hominis
Eubacterium_I ramulus
Faecalibacterium sp900539945
Oliverpabstia intestinalis
Coprococcus eutactus
Gemmiger formicilis
Eubacterium_R sp000434995
Bacteroides eggerthii
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Gallintestinimicrobium propionicum
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Butyribacter sp003529475
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Agathobacter sp000434275
Clostridium saudiense
Catenibacterium mitsuokai
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Aphodomorpha intestinavium
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Ligilactobacillus ruminis
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Butyrivibrio_A sp000431815
SFEL01 sp004557245
Slackia isoflavoniconvertens
Phocaeicola coprophilus
Clostridium_A leptum
Limivicinus sp900549645
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Vescimonas coprocola
Bifidobacterium adolescentis
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Roseburia hominis
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Agathobaculum butyriciproducens
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Limivivens faecihominis
Anaerotardibacter sp000435675
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Parasutterella excrementihominis
Bacteroides uniformis
Bacteroides stercoris
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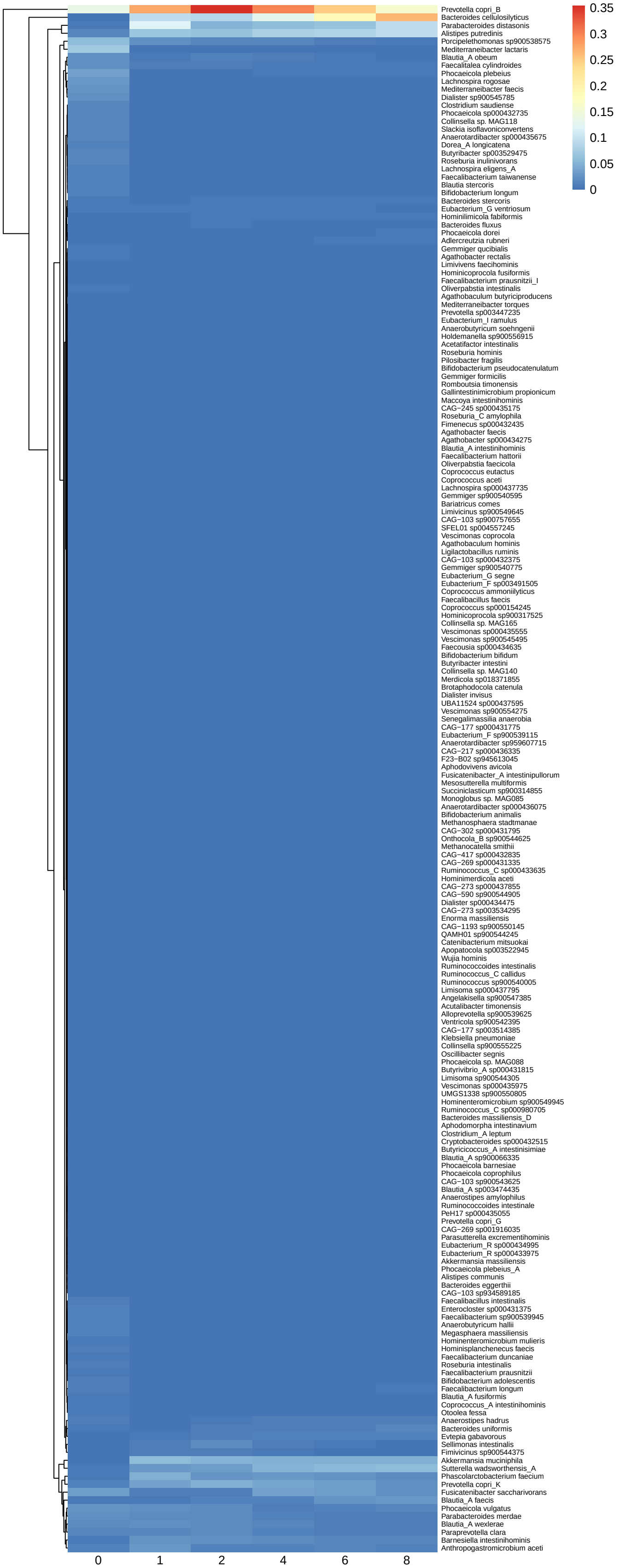
F2


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Bifidobacterium bifidum
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Butyribacter sp003529475
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Faecalibacterium longum
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Roseburia intestinalis
Agathobacter rectalis
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Hominimerdicola aceti
Fimivicinus sp900544375
Fusicatenibacter saccharivorans
Anaerostipes hadrus
UBA11524 sp000437595
Acetatifactor intestinalis
Hominisplanchenecus faecis
Phocaeicola dorei
Anaerostipes amylophilus
Bifidobacterium adolescentis
Anaerobutyricum soehngenii
Fusicatenibacter_A intestinipullorum
Gemmiger qucibialis
Bifidobacterium longum
Dorea_A longicatena
Blautia_A sp003474435
Adlercreutzia rubneri
Eubacterium_G ventriosum
Hominilimicola fabiformis
Oscillibacter segnis
Gallintestinimicrobium propionicum
Blautia_A fusiformis
Vescimonas sp000435555
Faecalitalea cylindroides
Bacteroides fluxus
Mediterraneibacter torques
Parasutterella excrementihominis
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Maccoya intestinihominis
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Roseburia hominis
Hominilimicola fabiformis
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Pilosibacter fragilis
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Romboutsia timonensis
Angelakisella sp900547385
SFEL01 sp004557245
CAG−245 sp000435175
Roseburia inulinivorans
Enterocloster sp000431375
Maccoya intestinihominis
Phocaeicola dorei
Acutalibacter timonensis
Otoolea fessa
Blautia_A fusiformis
Adlercreutzia rubneri
Paraprevotella clara
Eubacterium_G ventriosum
Eubacterium_I ramulus
Acetatifactor intestinalis
UBA11524 sp000437595
Faecalibacterium prausnitzii_I
Blautia_A sp900066335
Phocaeicola coprophilus
Faecalibacterium taiwanense
CAG−302 sp000431795
Collinsella sp900555225
Mediterraneibacter lactaris
Anaerobutyricum hallii
CAG−1193 sp900550145
Mediterraneibacter faecis
Faecalibacterium sp900539945
CAG−417 sp000432835
Agathobaculum butyriciproducens
Coprococcus_A intestinihominis
Faecalibacterium hattorii
Evtepia gabavorous
Phocaeicola sp000432735
QAMH01 sp900544245
Gemmiger qucibialis
Fusicatenibacter saccharivorans
Bariatricus comes
Limisoma sp900544305
Blautia_A faecis
Hominicoprocola sp900317525
Blautia_A wexlerae
Anthropogastromicrobium aceti
Eubacterium_F sp003491505
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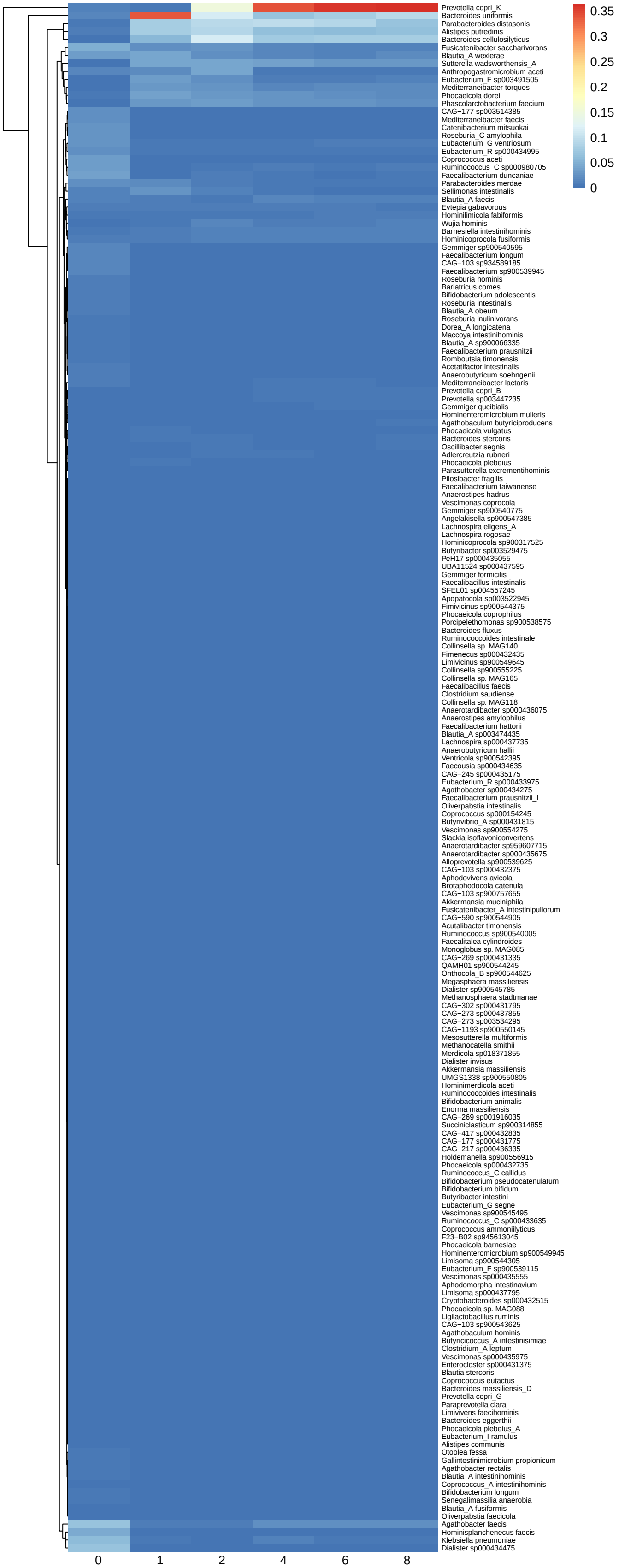
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Faecalitalea cylindroides
Phocaeicola plebeius
Lachnospira rogosae
Mediterraneibacter faecis
Dialister sp900545785
Clostridium saudiense
Phocaeicola sp000432735
Collinsella sp. MAG118
Slackia isoflavoniconvertens
Anaerotardibacter sp000435675
Dorea_A longicatena
Butyribacter sp003529475
Roseburia inulinivorans
Lachnospira eligens_A
Faecalibacterium taiwanense
Blautia stercoris
Bifidobacterium longum
Bacteroides stercoris
Eubacterium_G ventriosum
Hominilimicola fabiformis
Bacteroides fluxus
Phocaeicola dorei
Adlercreutzia rubneri
Gemmiger qucibialis
Agathobacter rectalis
Limivivens faecihominis
Hominicoprocola fusiformis
Faecalibacterium prausnitzii_I
Oliverpabstia intestinalis
Agathobaculum butyriciproducens
Mediterraneibacter torques
Prevotella sp003447235
Eubacterium_I ramulus
Anaerobutyricum soehngenii
Holdemanella sp900556915
Acetatifactor intestinalis
Roseburia hominis
Pilosibacter fragilis
Bifidobacterium pseudocatenulatum
Gemmiger formicilis
Romboutsia timonensis
Gallintestinimicrobium propionicum
Maccoya intestinihominis
CAG−245 sp000435175
Roseburia_C amylophila
Fimenecus sp000432435
Agathobacter faecis
Agathobacter sp000434275
Blautia_A intestinihominis
Faecalibacterium hattorii
Oliverpabstia faecicola
Coprococcus eutactus
Coprococcus aceti
Lachnospira sp000437735
Gemmiger sp900540595
Bariatricus comes
Limivicinus sp900549645
CAG−103 sp900757655
SFEL01 sp004557245
Vescimonas coprocola
Agathobaculum hominis
Ligilactobacillus ruminis
CAG−103 sp000432375
Gemmiger sp900540775
Eubacterium_G segne
Eubacterium_F sp003491505
Coprococcus ammoniilyticus
Faecalibacillus faecis
Coprococcus sp000154245
Hominicoprocola sp900317525
Collinsella sp. MAG165
Vescimonas sp000435555
Vescimonas sp900545495
Faecousia sp000434635
Bifidobacterium bifidum
Butyribacter intestini
Collinsella sp. MAG140
Merdicola sp018371855
Brotaphodocola catenula
Dialister invisus
UBA11524 sp000437595
Vescimonas sp900554275
Senegalimassilia anaerobia
CAG−177 sp000431775
Eubacterium_F sp900539115
Anaerotardibacter sp959607715
CAG−217 sp000436335
F23−B02 sp945613045
Aphodovivens avicola
Fusicatenibacter_A intestinipullorum
Mesosutterella multiformis
Succiniclasticum sp900314855
Monoglobus sp. MAG085
Anaerotardibacter sp000436075
Bifidobacterium animalis
Methanosphaera stadtmanae
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Onthocola_B sp900544625
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Hominimerdicola aceti
CAG−273 sp000437855
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Dialister sp000434475
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Enorma massiliensis
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QAMH01 sp900544245
Catenibacterium mitsuokai
Apopatocola sp003522945
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Ruminococcoides intestinalis
Ruminococcus_C callidus
Ruminococcus sp900540005
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Butyrivibrio_A sp000431815
Limisoma sp900544305
Vescimonas sp000435975
UMGS1338 sp900550805
Hominenteromicrobium sp900549945
Ruminococcus_C sp000980705
Bacteroides massiliensis_D
Aphodomorpha intestinavium
Clostridium_A leptum
Cryptobacteroides sp000432515
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Blautia_A sp900066335
Phocaeicola barnesiae
Phocaeicola coprophilus
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Blautia_A sp003474435
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Eubacterium_R sp000433975
Akkermansia massiliensis
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Bacteroides eggerthii
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Enterocloster sp000431375
Faecalibacterium sp900539945
Anaerobutyricum hallii
Megasphaera massiliensis
Hominenteromicrobium mulieris
Hominisplanchenecus faecis
Faecalibacterium duncaniae
Roseburia intestinalis
Faecalibacterium prausnitzii
Bifidobacterium adolescentis
Faecalibacterium longum
Blautia_A fusiformis
Coprococcus_A intestinihominis
Otoolea fessa
Anaerostipes hadrus
Bacteroides uniformis
Evtepia gabavorous
Sellimonas intestinalis
Fimivicinus sp900544375
Akkermansia muciniphila
Sutterella wadsworthensis_A
Phascolarctobacterium faecium
Prevotella copri_K
Fusicatenibacter saccharivorans
Blautia_A faecis
Phocaeicola vulgatus
Parabacteroides merdae
Blautia_A wexlerae
Paraprevotella clara
Barnesiella intestinihominis
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M5


Parabacteroides distasonis
Bacteroides cellulosilyticus
Agathobacter rectalis
Bifidobacterium pseudocatenulatum
Bacteroides uniformis
Blautia_A wexlerae
Alistipes putredinis
Parasutterella excrementihominis
Phascolarctobacterium faecium
Sellimonas intestinalis
Bariatricus comes
Faecalibacterium prausnitzii
Anaerostipes hadrus
Faecalibacterium duncaniae
Blautia_A fusiformis
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Ruminococcoides intestinale
Gemmiger qucibialis
Blautia_A intestinihominis
Mediterraneibacter torques
Phocaeicola dorei
Barnesiella intestinihominis
Pilosibacter fragilis
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Phocaeicola vulgatus
Lachnospira eligens_A
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Phocaeicola plebeius
Bacteroides stercoris
Faecalibacterium taiwanense
Bacteroides eggerthii
Bacteroides massiliensis_D
Klebsiella pneumoniae
Bacteroides fluxus
Limisoma sp000437795
Mediterraneibacter lactaris
CAG−103 sp934589185
Butyribacter intestini
Romboutsia timonensis
Agathobaculum butyriciproducens
Blautia_A sp003474435
Fimivicinus sp900544375
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Paraprevotella clara
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Coprococcus aceti
Limivicinus sp900549645
Ventricola sp900542395
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Gemmiger sp900540595
Gemmiger sp900540775
Ruminococcoides intestinalis
Faecalibacterium prausnitzii_I
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Fusicatenibacter_A intestinipullorum
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Merdicola sp018371855
Collinsella sp. MAG118
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Clostridium_A leptum
PeH17 sp000435055
Enterocloster sp000431375
Eubacterium_I ramulus
Anaerostipes amylophilus
Slackia isoflavoniconvertens
Lachnospira sp000437735
Roseburia inulinivorans
Faecalibacterium sp900539945
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Faecalibacterium longum
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Acetatifactor intestinalis
Oliverpabstia intestinalis
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Hominilimicola fabiformis
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Roseburia hominis
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Bifidobacterium bifidum
Eubacterium_G ventriosum
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Gemmiger formicilis
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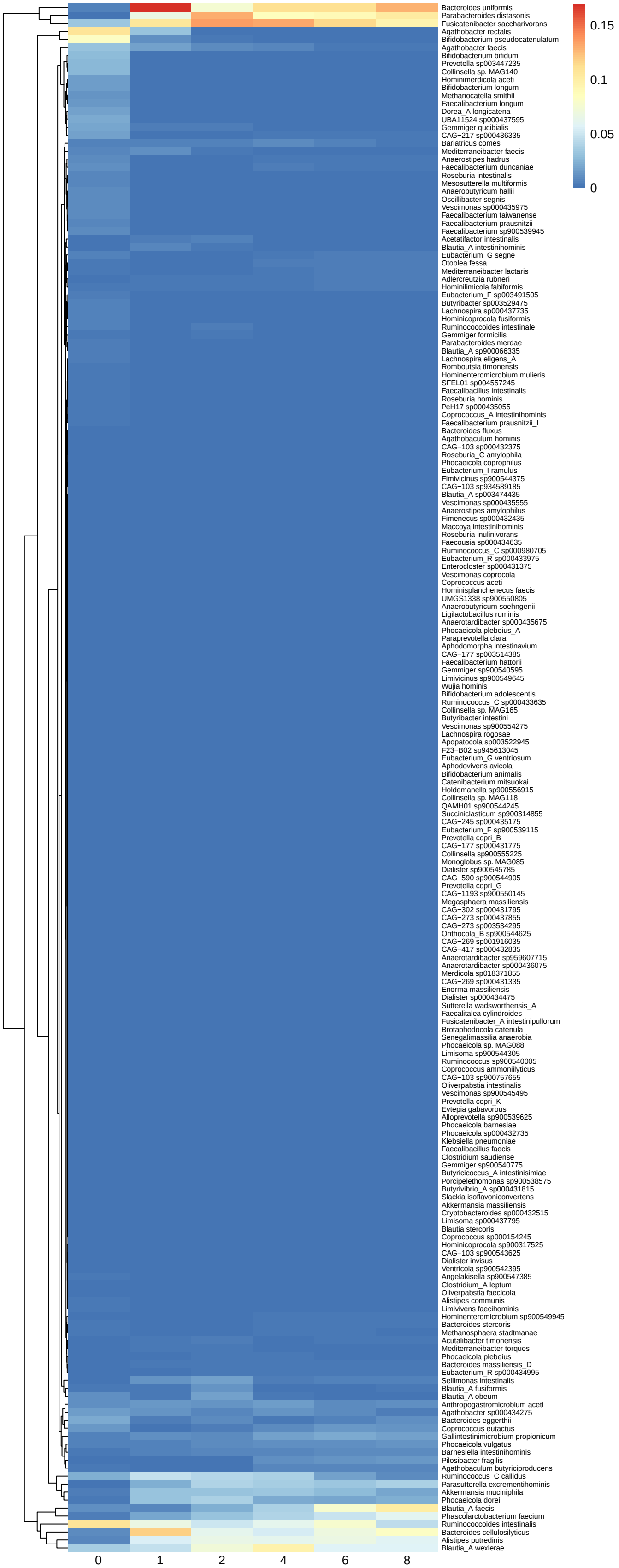
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Sutterella wadsworthensis_A
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Mediterraneibacter torques
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Mediterraneibacter faecis
Catenibacterium mitsuokai
Roseburia_C amylophila
Eubacterium_G ventriosum
Eubacterium_R sp000434995
Coprococcus aceti
Ruminococcus_C sp000980705
Faecalibacterium duncaniae
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Sellimonas intestinalis
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Evtepia gabavorous
Hominilimicola fabiformis
Wujia hominis
Barnesiella intestinihominis
Hominicoprocola fusiformis
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Roseburia intestinalis
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Roseburia inulinivorans
Dorea_A longicatena
Maccoya intestinihominis
Blautia_A sp900066335
Faecalibacterium prausnitzii
Romboutsia timonensis
Acetatifactor intestinalis
Anaerobutyricum soehngenii
Mediterraneibacter lactaris
Prevotella copri_B
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Hominenteromicrobium mulieris
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Bacteroides stercoris
Oscillibacter segnis
Adlercreutzia rubneri
Phocaeicola plebeius
Parasutterella excrementihominis
Pilosibacter fragilis
Faecalibacterium taiwanense
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Vescimonas coprocola
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Oliverpabstia intestinalis
Coprococcus sp000154245
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Vescimonas sp900554275
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Clostridium_A leptum
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Agathobacter rectalis
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Coprococcus_A intestinihominis
Bifidobacterium longum
Senegalimassilia anaerobia
Blautia_A fusiformis
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Agathobacter faecis
Hominisplanchenecus faecis
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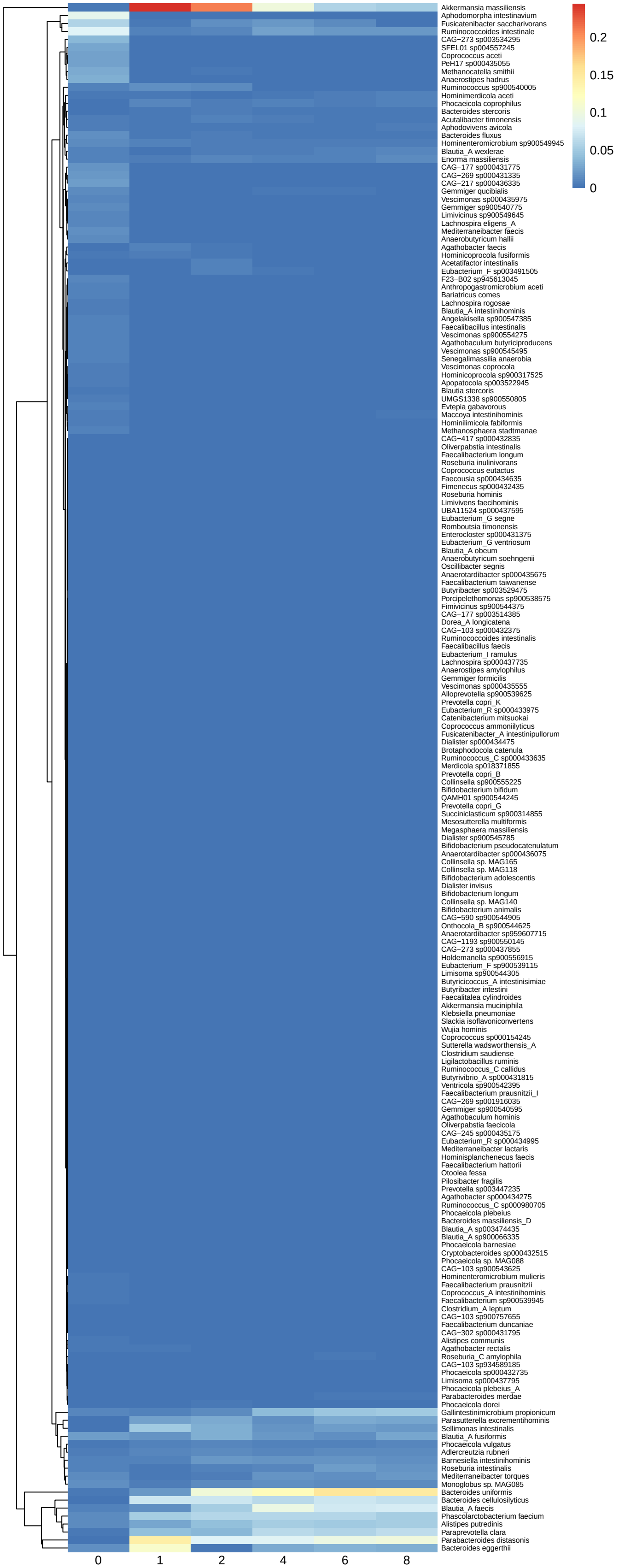
F7


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Akkermansia massiliensis
Sutterella wadsworthensis_A
Sellimonas intestinalis
Blautia_A wexlerae
Dialister invisus
Ruminococcoides intestinale
Agathobacter faecis
Anthropogastromicrobium aceti
Mediterraneibacter torques
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Coprococcus aceti
Bacteroides massiliensis_D
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Coprococcus ammoniilyticus
Phocaeicola sp. MAG088
Ligilactobacillus ruminis
Anaerostipes amylophilus
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Hominenteromicrobium mulieris
Dorea_A longicatena
Anaerobutyricum hallii
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Faecalibacterium prausnitzii
Alistipes communis
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CAG−177 sp003514385
Vescimonas sp900554275
Prevotella copri_G
Phascolarctobacterium faecium
CAG−273 sp000437855
Enorma massiliensis
Prevotella copri_B
Fusicatenibacter_A intestinipullorum
Dialister sp000434475
Anaerotardibacter sp959607715
Succiniclasticum sp900314855
CAG−273 sp003534295
Megasphaera massiliensis
CAG−302 sp000431795
Dialister sp900545785
Methanosphaera stadtmanae
Onthocola_B sp900544625
Methanocatella smithii
Mesosutterella multiformis
CAG−177 sp000431775
UMGS1338 sp900550805
CAG−590 sp900544905
CAG−1193 sp900550145
Aphodovivens avicola
F23−B02 sp945613045
Faecousia sp000434635
Ruminococcus sp900540005
Limisoma sp000437795
Alloprevotella sp900539625
Prevotella copri_K
Ruminococcoides intestinalis
Apopatocola sp003522945
CAG−103 sp900757655
Eubacterium_R sp000433975
Hominenteromicrobium sp900549945
Klebsiella pneumoniae
Cryptobacteroides sp000432515
Catenibacterium mitsuokai
Akkermansia muciniphila
CAG−103 sp900543625
Gemmiger sp900540775
Butyrivibrio_A sp000431815
Phocaeicola coprophilus
Blautia stercoris
Holdemanella sp900556915
Vescimonas sp000435555
Aphodomorpha intestinavium
Roseburia hominis
Lachnospira rogosae
Coprococcus eutactus
Faecalibacterium hattorii
Collinsella sp. MAG140
Coprococcus sp000154245
Collinsella sp. MAG118
Collinsella sp. MAG165
Ventricola sp900542395
Agathobacter sp000434275
Maccoya intestinihominis
Clostridium saudiense
Porcipelethomonas sp900538575
CAG−269 sp001916035
Oscillibacter segnis
Anaerostipes hadrus
Faecalibacillus intestinalis
Roseburia_C amylophila
Bifidobacterium longum
Eubacterium_I ramulus
Enterocloster sp000431375
Lachnospira eligens_A
Faecalibacterium longum
Faecalibacterium sp900539945
CAG−417 sp000432835
Eubacterium_G ventriosum
Hominisplanchenecus faecis
QAMH01 sp900544245
Parabacteroides merdae
Gemmiger formicilis
Phocaeicola sp000432735
Mediterraneibacter faecis
Barnesiella intestinihominis
Bacteroides eggerthii
Merdicola sp018371855
Bariatricus comes
Blautia_A obeum
Phocaeicola plebeius
Blautia_A fusiformis
Blautia_A sp003474435
Adlercreutzia rubneri
0
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F8


Bacteroides uniformis
Parabacteroides distasonis
Fusicatenibacter saccharivorans
Agathobacter rectalis
Bifidobacterium pseudocatenulatum
Agathobacter faecis
Bifidobacterium bifidum
Prevotella sp003447235
Collinsella sp. MAG140
Hominimerdicola aceti
Bifidobacterium longum
Methanocatella smithii
Faecalibacterium longum
Dorea_A longicatena
UBA11524 sp000437595
Gemmiger qucibialis
CAG−217 sp000436335
Bariatricus comes
Mediterraneibacter faecis
Anaerostipes hadrus
Faecalibacterium duncaniae
Roseburia intestinalis
Mesosutterella multiformis
Anaerobutyricum hallii
Oscillibacter segnis
Vescimonas sp000435975
Faecalibacterium taiwanense
Faecalibacterium prausnitzii
Faecalibacterium sp900539945
Acetatifactor intestinalis
Blautia_A intestinihominis
Eubacterium_G segne
Otoolea fessa
Mediterraneibacter lactaris
Adlercreutzia rubneri
Hominilimicola fabiformis
Eubacterium_F sp003491505
Butyribacter sp003529475
Lachnospira sp000437735
Hominicoprocola fusiformis
Ruminococcoides intestinale
Gemmiger formicilis
Parabacteroides merdae
Blautia_A sp900066335
Lachnospira eligens_A
Romboutsia timonensis
Hominenteromicrobium mulieris
SFEL01 sp004557245
Faecalibacillus intestinalis
Roseburia hominis
PeH17 sp000435055
Coprococcus_A intestinihominis
Faecalibacterium prausnitzii_I
Bacteroides fluxus
Agathobaculum hominis
CAG−103 sp000432375
Roseburia_C amylophila
Phocaeicola coprophilus
Eubacterium_I ramulus
Fimivicinus sp900544375
CAG−103 sp934589185
Blautia_A sp003474435
Vescimonas sp000435555
Anaerostipes amylophilus
Fimenecus sp000432435
Maccoya intestinihominis
Roseburia inulinivorans
Faecousia sp000434635
Ruminococcus_C sp000980705
Eubacterium_R sp000433975
Enterocloster sp000431375
Vescimonas coprocola
Coprococcus aceti
Hominisplanchenecus faecis
UMGS1338 sp900550805
Anaerobutyricum soehngenii
Ligilactobacillus ruminis
Anaerotardibacter sp000435675
Phocaeicola plebeius_A
Paraprevotella clara
Aphodomorpha intestinavium
CAG−177 sp003514385
Faecalibacterium hattorii
Gemmiger sp900540595
Limivicinus sp900549645
Wujia hominis
Bifidobacterium adolescentis
Ruminococcus_C sp000433635
Collinsella sp. MAG165
Butyribacter intestini
Vescimonas sp900554275
Lachnospira rogosae
Apopatocola sp003522945
F23−B02 sp945613045
Eubacterium_G ventriosum
Aphodovivens avicola
Bifidobacterium animalis
Catenibacterium mitsuokai
Holdemanella sp900556915
Collinsella sp. MAG118
QAMH01 sp900544245
Succiniclasticum sp900314855
CAG−245 sp000435175
Eubacterium_F sp900539115
Prevotella copri_B
CAG−177 sp000431775
Collinsella sp900555225
Monoglobus sp. MAG085
Dialister sp900545785
CAG−590 sp900544905
Prevotella copri_G
CAG−1193 sp900550145
Megasphaera massiliensis
CAG−302 sp000431795
CAG−273 sp000437855
CAG−273 sp003534295
Onthocola_B sp900544625
CAG−269 sp001916035
CAG−417 sp000432835
Anaerotardibacter sp959607715
Anaerotardibacter sp000436075
Merdicola sp018371855
CAG−269 sp000431335
Enorma massiliensis
Dialister sp000434475
Sutterella wadsworthensis_A
Faecalitalea cylindroides
Fusicatenibacter_A intestinipullorum
Brotaphodocola catenula
Senegalimassilia anaerobia
Phocaeicola sp. MAG088
Limisoma sp900544305
Ruminococcus sp900540005
Coprococcus ammoniilyticus
CAG−103 sp900757655
Oliverpabstia intestinalis
Vescimonas sp900545495
Prevotella copri_K
Evtepia gabavorous
Alloprevotella sp900539625
Phocaeicola barnesiae
Phocaeicola sp000432735
Klebsiella pneumoniae
Faecalibacillus faecis
Clostridium saudiense
Gemmiger sp900540775
Butyricicoccus_A intestinisimiae
Porcipelethomonas sp900538575
Butyrivibrio_A sp000431815
Slackia isoflavoniconvertens
Akkermansia massiliensis
Cryptobacteroides sp000432515
Limisoma sp000437795
Blautia stercoris
Coprococcus sp000154245
Hominicoprocola sp900317525
CAG−103 sp900543625
Dialister invisus
Ventricola sp900542395
Angelakisella sp900547385
Clostridium_A leptum
Oliverpabstia faecicola
Alistipes communis
Limivivens faecihominis
Hominenteromicrobium sp900549945
Bacteroides stercoris
Methanosphaera stadtmanae
Acutalibacter timonensis
Mediterraneibacter torques
Phocaeicola plebeius
Bacteroides massiliensis_D
Eubacterium_R sp000434995
Sellimonas intestinalis
Blautia_A fusiformis
Blautia_A obeum
Anthropogastromicrobium aceti
Agathobacter sp000434275
Bacteroides eggerthii
Coprococcus eutactus
Gallintestinimicrobium propionicum
Phocaeicola vulgatus
Barnesiella intestinihominis
Pilosibacter fragilis
Agathobaculum butyriciproducens
Ruminococcus_C callidus
Parasutterella excrementihominis
Akkermansia muciniphila
Phocaeicola dorei
Blautia_A faecis
Phascolarctobacterium faecium
Ruminococcoides intestinalis
Bacteroides cellulosilyticus
Alistipes putredinis
Blautia_A wexlerae
0.15
0.1
0.05
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F9


Akkermansia massiliensis
Aphodomorpha intestinavium
Fusicatenibacter saccharivorans
Ruminococcoides intestinale
CAG−273 sp003534295
SFEL01 sp004557245
Coprococcus aceti
PeH17 sp000435055
Methanocatella smithii
Anaerostipes hadrus
Ruminococcus sp900540005
Hominimerdicola aceti
Phocaeicola coprophilus
Bacteroides stercoris
Acutalibacter timonensis
Aphodovivens avicola
Bacteroides fluxus
Hominenteromicrobium sp900549945
Blautia_A wexlerae
Enorma massiliensis
CAG−177 sp000431775
CAG−269 sp000431335
CAG−217 sp000436335
Gemmiger qucibialis
Vescimonas sp000435975
Gemmiger sp900540775
Limivicinus sp900549645
Lachnospira eligens_A
Mediterraneibacter faecis
Anaerobutyricum hallii
Agathobacter faecis
Hominicoprocola fusiformis
Acetatifactor intestinalis
Eubacterium_F sp003491505
F23−B02 sp945613045
Anthropogastromicrobium aceti
Bariatricus comes
Lachnospira rogosae
Blautia_A intestinihominis
Angelakisella sp900547385
Faecalibacillus intestinalis
Vescimonas sp900554275
Agathobaculum butyriciproducens
Vescimonas sp900545495
Senegalimassilia anaerobia
Vescimonas coprocola
Hominicoprocola sp900317525
Apopatocola sp003522945
Blautia stercoris
UMGS1338 sp900550805
Evtepia gabavorous
Maccoya intestinihominis
Hominilimicola fabiformis
Methanosphaera stadtmanae
CAG−417 sp000432835
Oliverpabstia intestinalis
Faecalibacterium longum
Roseburia inulinivorans
Coprococcus eutactus
Faecousia sp000434635
Fimenecus sp000432435
Roseburia hominis
Limivivens faecihominis
UBA11524 sp000437595
Eubacterium_G segne
Romboutsia timonensis
Enterocloster sp000431375
Eubacterium_G ventriosum
Blautia_A obeum
Anaerobutyricum soehngenii
Oscillibacter segnis
Anaerotardibacter sp000435675
Faecalibacterium taiwanense
Butyribacter sp003529475
Porcipelethomonas sp900538575
Fimivicinus sp900544375
CAG−177 sp003514385
Dorea_A longicatena
CAG−103 sp000432375
Ruminococcoides intestinalis
Faecalibacillus faecis
Eubacterium_I ramulus
Lachnospira sp000437735
Anaerostipes amylophilus
Gemmiger formicilis
Vescimonas sp000435555
Alloprevotella sp900539625
Prevotella copri_K
Eubacterium_R sp000433975
Catenibacterium mitsuokai
Coprococcus ammoniilyticus
Fusicatenibacter_A intestinipullorum
Dialister sp000434475
Brotaphodocola catenula
Ruminococcus_C sp000433635
Merdicola sp018371855
Prevotella copri_B
Collinsella sp900555225
Bifidobacterium bifidum
QAMH01 sp900544245
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Succiniclasticum sp900314855
Mesosutterella multiformis
Megasphaera massiliensis
Dialister sp900545785
Bifidobacterium pseudocatenulatum
Anaerotardibacter sp000436075
Collinsella sp. MAG165
Collinsella sp. MAG118
Bifidobacterium adolescentis
Dialister invisus
Bifidobacterium longum
Collinsella sp. MAG140
Bifidobacterium animalis
CAG−590 sp900544905
Onthocola_B sp900544625
Anaerotardibacter sp959607715
CAG−1193 sp900550145
CAG−273 sp000437855
Holdemanella sp900556915
Eubacterium_F sp900539115
Limisoma sp900544305
Butyricicoccus_A intestinisimiae
Butyribacter intestini
Faecalitalea cylindroides
Akkermansia muciniphila
Klebsiella pneumoniae
Slackia isoflavoniconvertens
Wujia hominis
Coprococcus sp000154245
Sutterella wadsworthensis_A
Clostridium saudiense
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Butyrivibrio_A sp000431815
Ventricola sp900542395
Faecalibacterium prausnitzii_I
CAG−269 sp001916035
Gemmiger sp900540595
Agathobaculum hominis
Oliverpabstia faecicola
CAG−245 sp000435175
Eubacterium_R sp000434995
Mediterraneibacter lactaris
Hominisplanchenecus faecis
Faecalibacterium hattorii
Otoolea fessa
Pilosibacter fragilis
Prevotella sp003447235
Agathobacter sp000434275
Ruminococcus_C sp000980705
Phocaeicola plebeius
Bacteroides massiliensis_D
Blautia_A sp003474435
Blautia_A sp900066335
Phocaeicola barnesiae
Cryptobacteroides sp000432515
Phocaeicola sp. MAG088
CAG−103 sp900543625
Hominenteromicrobium mulieris
Faecalibacterium prausnitzii
Coprococcus_A intestinihominis
Faecalibacterium sp900539945
Clostridium_A leptum
CAG−103 sp900757655
Faecalibacterium duncaniae
CAG−302 sp000431795
Alistipes communis
Agathobacter rectalis
Roseburia_C amylophila
CAG−103 sp934589185
Phocaeicola sp000432735
Limisoma sp000437795
Phocaeicola plebeius_A
Parabacteroides merdae
Phocaeicola dorei
Gallintestinimicrobium propionicum
Parasutterella excrementihominis
Sellimonas intestinalis
Blautia_A fusiformis
Phocaeicola vulgatus
Adlercreutzia rubneri
Barnesiella intestinihominis
Roseburia intestinalis
Mediterraneibacter torques
Monoglobus sp. MAG085
Bacteroides uniformis
Bacteroides cellulosilyticus
Blautia_A faecis
Phascolarctobacterium faecium
Alistipes putredinis
Paraprevotella clara
Parabacteroides distasonis
Bacteroides eggerthii
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