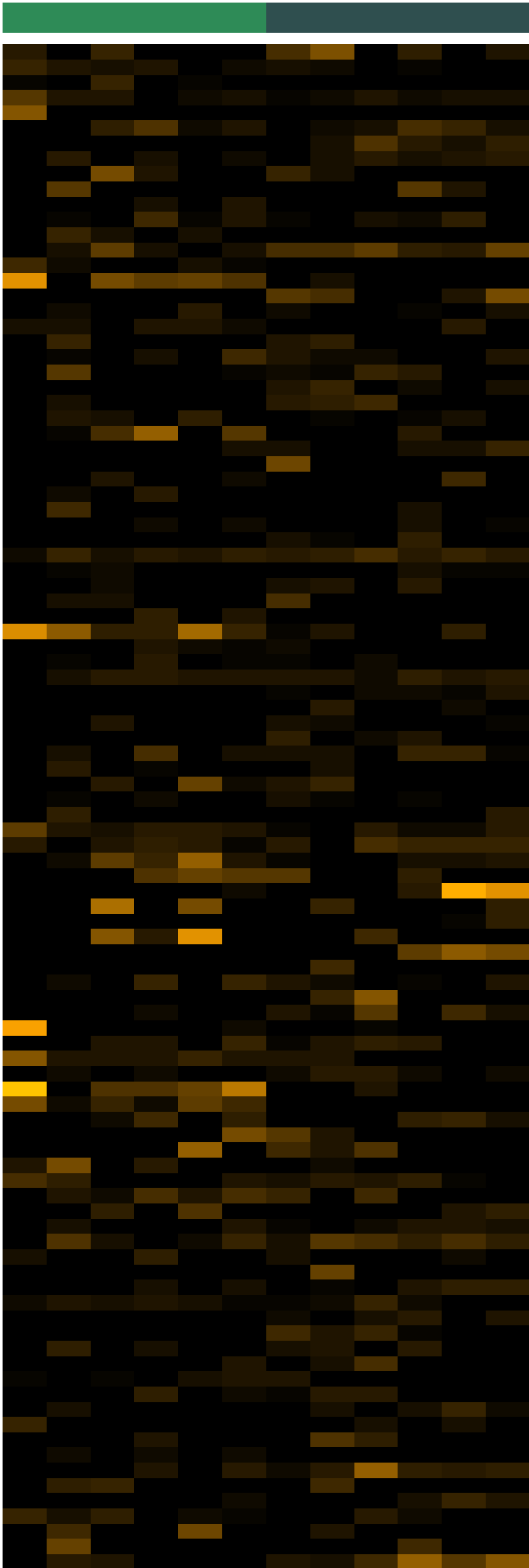




Comparison\_1  
Control  
Amoxicillin



| Species  |
|--|
| Lachnospiraceae_[G-11] bacterium_MOT-178                   |
| Alistipes sp._MOT-127                                      |
| Bacteroides acidofaciens                                   |
| Bacteroides caecimuris                                     |
| Blautia caecimuris   |
| Prevotella sp._MOT-128                                     |
| Lactobacillus taiwanensis                                  |
| Oscillospiraceae_[G-2] bacterium_MOT-149                   |
| Lachnospiraceae_[G-14] bacterium_MOT-183                   |
| Lachnospiraceae_[G-9] bacterium_MOT-174                    |
| Eubacteriales_[G-1] bacterium_MOT-158                      |
| Bacteroides stercorisoris                                  |
| Muribaculum intestinale                                    |
| Lachnospiraceae_[G-14] bacterium_MOT-184                   |
| Parabacteroides goldsteinii                                |
| Akkermansia muciniphila                                    |
| Kineothrix alysoides_nov_95.227%                           |
| Anaerotruncus rubifantis_nov_92.760%                       |
| Muribaculaceae_[G-2] bacterium_MOT-104_nov_91.991%         |
| Mediterraneibacter [Ruminococcus] gnavus_nov_93.424%       |
| Alistipes putredinis_nov_95.887%                           |
| Oscillibacter valericigenes_nov_95.260%                    |
| Kineothrix alysoides_nov_97.279%                           |
| Pseudobutyrvibrio ruminis_nov_91.176%                      |
| Saccharofermentans acetigenes_nov_88.764%                  |
| Lacrimispora indolis_nov_90.724%                           |
| Lawsonibacter asaccharolyticus_nov_97.973%                 |
| Lachnospiraceae_[G-14] bacterium_MOT-185_nov_92.358%       |
| Muribaculaceae_[G-2] bacterium_MOT-104_nov_90.870%         |
| Muribaculaceae_[G-2] bacterium_MOT-104_nov_93.043%         |
| Muribaculaceae_[G-2] bacterium_MOT-104_nov_91.106%         |
| Alistipes putredinis_nov_95.879%                           |
| Odoribacter splanchnicus_nov_93.939%                       |
| Oscillospiraceae_[G-2] bacterium_MOT-149_nov_95.506%       |
| Ruthenibacterium lactatiformans_nov_97.045%                |
| Lachnospiraceae_[G-11] bacterium_MOT-178_nov_97.978%       |
| Lachnospiraceae_[G-14] bacterium_MOT-185_nov_96.599%       |
| Faecalicatena orotica_nov_95.238%                          |
| Prevotellamassilia timonensis_nov_94.168%                  |
| Eisenbergiella massiliensis_nov_96.599%                    |
| Neglectibacter timonensis_nov_97.500%                      |
| Maiihella massiliensis_nov_92.094%                         |
| Neglectibacter timonensis_nov_97.727%                      |
| Lachnospiraceae_[G-11] bacterium_MOT-176_nov_95.946%       |
| Lachnospiraceae_[G-14] bacterium_MOT-184_nov_95.405%       |
| Acetatifactor muris_nov_96.145%                            |
| Oscillospiraceae_[G-2] bacterium_MOT-149_nov_95.056%       |
| Oscillospiraceae_[G-2] bacterium_MOT-149_nov_95.281%       |
| Lachnospiraceae_[G-6] bacterium_MOT-171_nov_95.238%        |
| Eubacteriales_[G-2] bacterium_MOT-162_nov_95.260%          |
| Anaerosporeobacter mobilis_nov_95.000%                     |
| Alistipes putredinis_nov_96.529%                           |
| Phoceia massiliensis_nov_95.682%                           |
| Muribaculum intestinale_nov_93.737%                        |
| Lachnospiraceae_[G-12] bacterium_MOT-179_nov_94.796%       |
| Lachnospiraceae_[G-11] bacterium_MOT-176_nov_97.297%       |
| Oscillospiraceae_[G-2] bacterium_MOT-149_nov_93.919%       |
| Acetatifactor muris_nov_92.551%                            |
| Kineothrix alysoides_nov_97.059%                           |
| Lachnospiraceae_[G-12] bacterium_MOT-179_nov_92.986%       |
| Duncaniella freteri_nov_90.456%                            |
| Kineothrix alysoides_nov_93.682%                           |
| Roseburia faecis_nov_97.964%                               |
| Duncaniella freteri_nov_93.103%                            |
| Muribaculaceae_[G-2] bacterium_MOT-104_nov_93.074%         |
| Lacrimispora xylanolytica_nov_97.285%                      |
| Culturomica massiliensis_nov_93.709%                       |
| Kineothrix alysoides_nov_95.928%                           |
| Lachnospiraceae_[G-10] bacterium_MOT-175_nov_96.372%       |
| Muribaculaceae_[G-2] bacterium_MOT-104_nov_92.191%         |
| Lachnospiraceae_[G-10] bacterium_MOT-175_nov_92.174%       |
| Lachnospiraceae_[G-10] bacterium_MOT-175_nov_96.825%       |
| Muribaculaceae_[G-2] bacterium_MOT-104_nov_89.462%         |
| Eubacterium coprostanoligenes_nov_95.485%                  |
| Muribaculaceae_[G-2] bacterium_MOT-104_nov_91.974%         |
| Lachnospiraceae_[G-12] bacterium_MOT-179_nov_92.534%       |
| Lachnospiraceae_[G-14] bacterium_MOT-184_nov_94.989%       |
| Pseudoflavonifractor capillosus_nov_95.721%                |
| Anaerotruncus lactatifermentans_nov_95.270%                |
| Caecibacterium sporiformans_nov_95.045%                    |
| Alistipes timonensis_nov_97.831%                           |
| Oscillospiraceae_[G-2] bacterium_MOT-149_nov_95.946%       |
| Oscillospiraceae_[G-2] bacterium_MOT-149_nov_94.157%       |
| Alistipes senegalensis_nov_95.228%                         |
| Oscillospiraceae_[G-4] bacterium_MOT-151_nov_91.723%       |
| Lachnospiraceae_[G-11] bacterium_MOT-177_nov_97.523%       |
| Bacteroides acidifaciens_acidofaciens                      |
| Lachnospiraceae_[G-12] bacterium_MOT-179_bacterium_MOT-183 |
| multigenus multispecies_sppn10_2_nov_95.918%               |
| Alistipes multispecies_sppn12_2_nov_96.304%                |
| multigenus multispecies_sppn13_5_nov_94.570%               |
| Eubacteriales_[G-1] multispecies_sppn15_2_nov_97.511%      |
| Bacteroidetes_[G-3] multispecies_sppn2_2_nov_87.554%       |
| Lachnospiraceae_[G-11] multispecies_sppn4_2_nov_96.847%    |
| multigenus multispecies_sppn5_2_nov_97.279%                |
| Bacteroides multispecies_sppn6_2_nov_96.312%               |
| multigenus multispecies_sppn7_2_nov_92.777%                |
| multigenus multispecies_sppn8_3_nov_95.011%                |
| multigenus multispecies_sppn9_2_nov_93.002%                |

F8810.S10  
F8810.S11  
F8810.S12  
F8810.S22  
F8810.S23  
F8810.S24  
F8810.S01  
F8810.S02  
F8810.S03  
F8810.S13  
F8810.S14  
F8810.S15

Samples