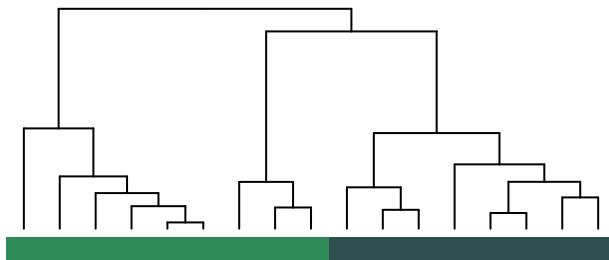


Value



Group  
WT0Defc  
WT3Hlig

Bacteroides acidifaciens  
Alistipes sp.\_MOT-127  
Bacteroides acidifaciens\_acidofaciens  
Faecalibaculum multispecies\_sppn189\_2\_nov\_97.279%  
Ileibacterium valens  
Oscillospiraceae\_[G-3] bacterium\_MOT-150  
Muribaculaceae\_[G-1] bacterium\_MOT-129  
Oscillospiraceae\_[G-7] bacterium\_MOT-154  
Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_89.177%  
Oscillospiraceae\_[G-6] bacterium\_MOT-153\_nov\_94.157%  
Eubacterium ventriosum\_nov\_96.372%  
Eubacteriales\_[G-1] bacterium\_MOT-159\_nov\_94.118%  
Desulfovibrio fairfieldensis\_nov\_94.421%  
multigenus multispecies\_sppn292\_3\_nov\_87.368%  
Lachnospiraceae\_[G-11] bacterium\_MOT-178  
Eubacteriales\_[G-2] bacterium\_MOT-162\_nov\_95.260%  
Eubacteriales\_[G-1] bacterium\_MOT-160  
Duncaniella frateri\_nov\_90.672%  
Muribaculaceae\_[G-1] bacterium\_MOT-129\_nov\_89.224%  
Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_91.810%  
Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_92.625%  
Eubacteriales\_[G-4] bacterium\_MOT-165  
Bifidobacterium pseudolongum  
Lachnospiraceae\_[G-14] bacterium\_MOT-184  
Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_90.693%  
Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_90.929%  
Alistipes putredinis\_nov\_96.753%  
Helicobacter typhlonius  
Paraeggerthelia hongkongensis\_nov\_92.793%  
Duncaniella frateri\_nov\_88.248%  
Longibaculum muris\_nov\_92.094%  
Eubacteriales\_[G-2] bacterium\_MOT-162  
Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_91.323%  
Lactobacillus\_limosilactobacillus\_reuteri\_reuteri\_clade\_938  
Erysipelatoclostridium ramosum\_nov\_87.975%  
Lachnospiraceae\_[G-9] bacterium\_MOT-174\_nov\_95.701%  
Millionella massiliensis\_nov\_92.424%  
Eubacteriales\_[G-4] bacterium\_MOT-164\_nov\_95.485%  
Lachnospiraceae\_[G-12] bacterium\_MOT-179\_nov\_92.534%  
Faecalibaculum orotica\_nov\_95.692%  
Millionella massiliensis\_nov\_87.257%  
Eubacteriales\_[G-4] bacterium\_MOT-165\_nov\_94.357%  
Aminipila butyrlica\_nov\_91.461%  
Culturomica massiliensis\_nov\_91.145%  
Carnobacteriaceae\_[G-1] bacterium\_MOT-198  
Odoribacter splanchnicus\_nov\_93.290%  
Parasutterella excrementimonis\_nov\_93.576%  
Ruminococcus albus\_nov\_92.517%  
Amedibacillus dolichus\_nov\_91.649%  
Bifidobacterium choerinum\_pseudolongum  
Neglectibacter timonensis\_nov\_97.506%  
Bacteroides caecumuris  
Saccharibacteria\_(TM7)\_\*[G-3] bacterium\_HMT\_351\_nov\_95.500%  
Kinotrichia alysoides\_nov\_97.517%  
multigenus multispecies\_sppn87\_2\_nov\_92.225%  
Pseudoflavorifractor phocaensis\_nov\_89.189%  
Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_93.737%  
Oscillospiraceae\_[G-4] bacterium\_MOT-151\_nov\_96.854%  
Lactobacillus intestinalis  
Helicobacter ganmani  
Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_94.577%  
Faecalibaculum rodentium  
Lactobacillus taiwanensis  
Anaerostipes caccae  
Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_92.857%  
Eubacteriales\_[G-3] bacterium\_MOT-163\_nov\_93.468%  
Cutibacterium acnes  
Actinidia eriantha\_nov\_97.291%  
Blautia wexlerae  
Escherichia\_Shigella coli\_fergusonii\_flexneri\_sonnei  
Clostridium disporicum\_saudicense  
Muribaculaceae\_[G-1] bacterium\_MOT-129\_nov\_91.087%  
Muribaculaceae\_[G-1] bacterium\_MOT-129\_nov\_91.991%  
Triticum aestivum  
Enterococcus faecalis  
Eubacteriales\_[G-1] bacterium\_MOT-159  
Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_93.521%  
Lactobacillus johnsonii  
Muribaculum intestine  
Eubacteriales\_[G-4] bacterium\_MOT-164  
Muribaculaceae\_[G-1] bacterium\_MOT-129\_nov\_91.304%  
Streptococcus acidominimus\_sp.\_MOT-012  
Staphylococcus saprophyticus\_xylosus  
Muribaculum intestine\_nov\_87.957%  
Parabacteroides goldsteinii  
Ligilactobacillus animalis\_apodemii\_murinus  
Muribaculaceae\_[G-1] bacterium\_MOT-129\_nov\_91.757%  
Prevotella sp.\_MOT-128  
Erysipelatoclostridium [Clostridium] cocleatum  
Muribaculaceae\_[G-1] bacterium\_MOT-129\_nov\_91.087%  
Ralstonia picketti  
Corynebacterium mastitidis  
Pasteurella\_Rodentibacter caecumuris\_heyliae  
Gemella sp.\_MOT-043  
Muribaculaceae\_[G-1] bacterium\_MOT-129\_nov\_91.540%  
Paracoccus carotinifaciens\_hibiscisolii\_marcusii\_nototheniae  
Akkermansia muciniphila  
Streptococcus danieliae  
Rodentibacter pneumatropicus

F0000.S058 F0000.S054 F0000.S059 F0000.S061 F0000.S062 F0000.S060 F0000.S056 F0000.S055 F0000.S057 F0000.S013 F0000.S011 F0000.S014 F0000.S018 F0000.S015 F0000.S016 F0000.S012 F0000.S017

Samples

species