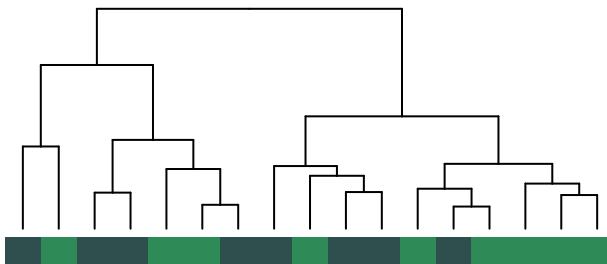


Value



Group  
KO5Dig  
WT5Dig

Rodentibacter pneumotropicus  
Pasteurella\_Rodentibacter caecimuris\_heylii  
Streptococcus acidominimus\_sp.\_MOT-012  
Helicobacter typhlonius  
Prevotellamassilia timonensis\_nov\_93.952%  
Oscillospiraceae\_[G-7] bacterium\_MOT-154  
Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_92.625%  
Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_92.704%  
Muribaculum intestinalе\_nov\_91.106%  
Parabacteroides merdae\_nov\_95.032%  
Clostridium disparicum\_saudicense  
Lachnospiraceae\_[G-11] bacterium\_MOT-178  
Neglectibacter timonensis\_nov\_96.825%  
Bacteroides caecimuris  
Oscillospiraceae\_[G-3] bacterium\_MOT-150  
Rhodoligotrophos multispecies\_sppn51\_2\_nov\_79.912%  
Actinidia eriantha\_nov\_97.291%  
Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_92.641%  
Lactobacillus\_Limosilactobacillus reuteri\_reuteri\_clade\_938  
Lachnospiraceae\_[G-14] bacterium\_MOT-185  
Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_92.641%  
Muribaculaceae\_[G-1] bacterium\_MOT-129  
Muribaculum intestinalе\_nov\_92.258%  
Bacteroides acidifaciens\_acidifaciens  
Bacteroidetes\_[G-3] multispecies\_sppn10\_2\_nov\_87.339%  
Bacteroidetes\_[G-3] multispecies\_sppn13\_2\_nov\_86.422%  
Bacteroides multispecies\_sppn705\_2\_nov\_96.104%  
Eubacteriales\_[G-4] bacterium\_MOT-164  
Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_90.672%  
Parabacteroides distasonis\_nov\_97.624%  
Parabacteroides distasonis\_nov\_96.963%  
Parabacteroides distasonis\_nov\_97.397%  
Parasutterella\_excrementihominis\_nov\_94.231%  
Enterocloster aldenensis  
Parabacteroides merdae\_nov\_95.032%  
Faecalcatena multispecies\_sppn189\_2\_nov\_97.279%  
Rothia nasimirum  
Alistipes putredinis\_nov\_96.529%  
Rhodoligotrophos multispecies\_sppn60\_2\_nov\_79.736%  
Olsenella multispecies\_sppn21\_2\_nov\_95.991%  
Paracoccus carotinifaciens\_hibiscisolii\_markusii\_nototheniae  
Erysipelatoclostridium [Clostridium] innocuum  
Turicimonas muris  
Muribaculaceae\_[G-1] bacterium\_MOT-129\_nov\_86.853%  
Parabacteroides distasonis\_nov\_96.753%  
Parabacteroides merdae\_nov\_95.032%  
Olsenella multispecies\_sppn704\_2\_nov\_96.214%  
Lawsonibacter asaccharolyticus\_nov\_95.281%  
Muribaculaceae\_[G-1] bacterium\_MOT-129\_nov\_87.284%  
Lactobacillus johnsonii  
Muribaculaceae\_[G-1] bacterium\_MOT-129\_nov\_91.540%  
Prevotella sp.\_MOT-128  
Helicobacter ganmani  
Ihubacter massiliensis\_nov\_96.840%  
Muribaculaceae\_[G-1] bacterium\_MOT-129\_nov\_91.757%  
Akermansia muciniphila  
Faecalcatena orotica\_nov\_95.711%  
Cutibacterium acnes  
Staphylococcus ureilyticus  
Ralstonia picketti  
Desulfovibrio fairfieldensis  
Triticum aestivum  
Bacteroidetes\_[G-3] multispecies\_sppn670\_2\_nov\_87.768%  
Parasutterella excrementihominis\_nov\_94.004%  
Parabacteroides distasonis\_nov\_97.186%  
Alistipes putredinis\_nov\_96.104%  
Olsenella multispecies\_sppn107\_2\_nov\_95.982%  
Desulfovibrio fairfieldensis\_nov\_94.433%  
Anaerostipes caccae  
Adlercreutzia caecimuris  
Muribaculum intestinalе  
Adlercreutzia mucosicola  
Muribaculaceae\_[G-1] bacterium\_MOT-129\_nov\_91.087%  
Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_93.521%  
Corynebacterium mastitidis  
Lachnospiraceae\_[G-3] bacterium\_MOT-168  
Parabacteroides goldsteinii  
Staphylococcus saprophyticus\_xylosus  
Erysipelatoclostridium [Clostridium] cocleatum  
Alistipes sp.\_MOT-127  
Alistipes timonensis  
Gemella sp.\_MOT-043  
Robinsoniella peoriensis  
Ileibacterium valens  
Faecalibaculum rodentium  
Escherichia\_Shigella coli\_fergusonii\_flexneri\_sonnei  
Terrisporobacter petrolearius  
Clostridium subterminale\_sulfidigenes\_thiosulfatireducens  
Bacteroides acidifaciens  
Ruthenibacterium lactatiformans  
Bacteroides multispecies\_sppn9\_2\_nov\_96.095%  
Enterococcus faecalis  
Maihella massiliensis\_nov\_92.111%  
Carnobacteriaceae\_[G-1] bacterium\_MOT-198  
Bifidobacterium pseudolongum  
Ligilactobacillus animalis\_apodermi\_murinus  
Erysipelatoclostridium ramosum  
Streptococcus danieliae  
Phocaeicola sartorii  
Erysipelotrichaceae\_[G-1] bacterium\_MOT-189

F0000.S080  
F0000.S075  
F0000.S082  
F0000.S083  
F0000.S078  
F0000.S079  
F0000.S087  
F0000.S084  
F0000.S071  
F0000.S085  
F0000.S081  
F0000.S073  
F0000.S086  
F0000.S076  
F0000.S077  
F0000.S072  
F0000.S074

Samples

Species