



- Akkermansia muciniphila  
Lactobacillus johnsonii  
Klebsiella pneumoniae  
Acutalibacter muris  
Erysipelotrichaceae\_[G-1] bacterium\_MOT-189  
Lachnospiraceae\_[G-14] bacterium\_MOT-185  
Enterococcus faecalis  
Adlercreutzia caecimuris  
Oscillospiraceae\_[G-4] bacterium\_MOT-151\_nov\_95.842%  
Duncaniella freteri\_nov\_93.712%  
Oscillospiraceae\_[G-2] bacterium\_MOT-149\_nov\_94.781%  
Lachnospiraceae\_[G-7] bacterium\_MOT-172\_nov\_91.097%  
Duncaniella freteri\_nov\_89.919%  
Kineothrix alysoides\_nov\_89.562%  
Pseudoflavonifractor phocaeensis\_nov\_95.859%  
Lawsonibacter asaccharolyticus\_nov\_91.116%  
Lachnospiraceae\_[G-6] bacterium\_MOT-171\_nov\_93.933%  
Lachnospiraceae\_[G-14] bacterium\_MOT-185\_nov\_96.781%  
Lachnospiraceae\_[G-12] bacterium\_MOT-179\_nov\_87.097%  
Kineothrix alysoides\_nov\_89.792%  
Roseburia intestinalis\_nov\_90.229%  
Eubacteriales\_[G-4] bacterium\_MOT-165\_nov\_92.781%  
Lachnospiraceae\_[G-14] bacterium\_MOT-182\_nov\_90.254%  
Agathobaculum desmolans\_nov\_91.649%  
Lachnospiraceae\_[G-6] bacterium\_MOT-171\_nov\_94.759%  
Lawsonibacter asaccharolyticus\_nov\_91.116%  
Lachnospiraceae\_[G-2] bacterium\_HMT\_096\_nov\_91.632%  
Eubacterium xylanophilum\_nov\_91.075%  
Mollicutes\_[G-2] bacterium\_MOT-187\_nov\_94.892%  
Oscillospiraceae\_[G-3] bacterium\_MOT-150\_nov\_93.125%  
Muricomes intestini\_nov\_89.375%  
Muribaculaceae\_[G-1] bacterium\_MOT-129\_nov\_88.577%  
Lachnospiraceae\_[G-2] bacterium\_MOT-167\_nov\_93.096%  
Muribaculaceae\_[G-1] bacterium\_MOT-129\_nov\_86.640%  
Lachnospiraceae\_[G-11] bacterium\_MOT-176\_nov\_94.898%  
Oscillospiraceae\_[G-2] bacterium\_MOT-149\_nov\_93.319%  
Oscillospiraceae\_[G-3] bacterium\_MOT-150\_nov\_91.511%  
Oscillospiraceae\_[G-2] bacterium\_MOT-149\_nov\_95.010%  
Lachnospiraceae\_[G-7] bacterium\_MOT-172\_nov\_91.097%  
Faecalicatena fissicatena\_nov\_95.407%  
Lachnospiraceae\_[G-14] bacterium\_MOT-185\_nov\_93.348%  
Eubacteriales\_[G-3] bacterium\_MOT-163\_nov\_85.944%  
Lachnospiraceae\_[G-3] bacterium\_MOT-168\_nov\_94.792%  
Faecalicatena fissicatena\_nov\_94.154%  
Acutalibacter muris\_nov\_96.694%  
Eubacteriales\_[G-3] bacterium\_MOT-163\_nov\_89.157%  
Roseburia hominis\_nov\_92.754%  
Lachnospiraceae\_[G-11] bacterium\_MOT-176\_nov\_89.858%  
Muribaculaceae\_[G-1] bacterium\_MOT-129\_nov\_85.887%  
Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_89.022%  
Oscillospiraceae\_[G-3] bacterium\_MOT-150\_nov\_93.582%  
Kineothrix alysoides\_nov\_85.921%  
Anaerotignum faecicola\_nov\_85.287%  
Hathewayia proteolytica\_nov\_83.297%  
Oscillospiraceae\_[G-2] bacterium\_MOT-149\_nov\_93.168%  
Alistipes senegalensis\_nov\_93.089%  
Prevotella shahii\_nov\_87.602%  
Acutalibacter muris\_nov\_96.289%  
Culturomica massiliensis\_nov\_89.817%  
Lachnospiraceae\_[G-6] bacterium\_MOT-171\_nov\_94.572%  
Oscillospiraceae\_[G-1] bacterium\_MOT-147\_nov\_96.674%  
Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_89.000%  
Eubacteriales\_[G-1] bacterium\_MOT-159\_nov\_94.268%  
Hydrogenoanaerobacterium saccharovorans\_nov\_90.041%  
Acetivibrio cellulolyticus\_nov\_83.801%  
Duncaniella freteri\_nov\_89.775%  
Oscillospiraceae\_[G-2] bacterium\_MOT-149\_nov\_95.198%  
Lachnospiraceae\_[G-10] bacterium\_MOT-175\_nov\_90.369%  
Breznakia pachnodae\_nov\_82.824%  
Acetivibrio cellulolyticus\_nov\_83.761%  
Acetivibrio cellulolyticus\_nov\_83.153%  
Glucerbacter canis\_nov\_93.305%  
Erysipelatoclostridium [Clostridium] innocuum\_nov\_88.270%  
Lachnospiraceae\_[G-6] bacterium\_MOT-171\_nov\_94.979%  
Oscillospiraceae\_[G-4] bacterium\_MOT-151\_nov\_94.179%  
Eubacteriales\_[G-4] bacterium\_MOT-164\_nov\_97.228%  
Pseudoflavonifractor capillosus\_nov\_89.897%  
Flavonifractor plautii\_nov\_92.308%  
Lachnospiraceae\_[G-6] bacterium\_MOT-171\_nov\_91.213%  
Sporobacter termitidis\_nov\_87.580%  
Oscillospiraceae\_[G-4] bacterium\_MOT-151\_nov\_92.100%  
Lachnospiraceae\_[G-14] bacterium\_MOT-183\_nov\_97.854%  
Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_89.157%  
Longibaculum muris\_nov\_93.361%  
Lachnospiraceae\_[G-6] bacterium\_MOT-171\_nov\_93.305%  
Christensenella hongkongensis\_nov\_86.308%  
Phoceia massiliensis\_nov\_90.426%  
Roseburia hominis\_nov\_91.476%  
Hydrogenoanaerobacterium saccharovorans\_nov\_88.589%  
Flavonifractor plautii\_nov\_93.555%  
Neglectibacter timonensis\_nov\_94.490%  
Eubacteriales\_[G-3] bacterium\_MOT-163\_nov\_85.825%  
Anaeromassilibacillus senegalensis\_nov\_92.489%  
Duncaniella freteri\_nov\_90.612%  
Oscillospiraceae\_[G-2] bacterium\_MOT-149\_nov\_96.466%  
Anaerotignum aminivorans\_nov\_93.008%  
Acetivibrio cellulolyticus\_nov\_85.776%  
Lachnospiraceae\_[G-11] bacterium\_MOT-177\_nov\_94.606%  
Blautia multispecies\_sppn3\_2\_nov\_94.990%  
Faecalicatena multispecies\_sppn8\_2\_nov\_92.067%

Species

F15387.S41  
F15387.S42  
F15387.S43  
F15387.S44  
F15387.S45  
F15387.S46  
F15387.S47  
F15387.S48  
F15387.S49  
F15387.S50

Samples