



Group  
■ Group A1  
■ Group A2

Culturomica massiliensis\_nov\_93.709%  
 Eubacteriales\_[G-2] bacterium\_MOT-162\_nov\_95.260%  
 Anaerotruncus rubiinfantis\_nov\_92.517%  
 Rhodospirillum rubrum\_nov\_88.036%  
 Lachnospiraceae\_[G-9] bacterium\_MOT-174\_nov\_96.136%  
 Turicimonas muris  
 multigenus multispecies\_sppn13\_5\_nov\_94.570%  
 Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_90.022%  
 Anaerotruncus rubiinfantis\_nov\_93.182%  
 Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_92.873%  
 Alistipes timonensis\_nov\_97.831%  
 Alistipes putredinis\_nov\_95.879%  
 Eisenbergiella massiliensis\_nov\_96.599%  
 Eubacteriales\_[G-1] multispecies\_sppn15\_2\_nov\_97.511%  
 Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_93.043%  
 Neglectibacter timonensis\_nov\_97.500%  
 Anaerotignum lactatifermentans\_nov\_95.270%  
 Clostridiales\_[F-1][G-1] bacterium\_HMT\_093\_nov\_90.337%  
 Petrocella atlantisensis\_nov\_87.810%  
 Ruthenibacterium lactatiformans\_nov\_97.045%  
 Lachnospiraceae\_[G-14] bacterium\_MOT-185\_nov\_96.599%  
 Oscillospiraceae\_[G-2] bacterium\_MOT-149\_nov\_95.270%  
 Coprococcus catus\_nov\_94.570%  
 Acutalibacter muris  
 multigenus multispecies\_sppn10\_2\_nov\_95.918%  
 Roseburia multispecies\_sppn21\_3\_nov\_95.711%  
 Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_90.870%  
 multigenus multispecies\_sppn22\_2\_nov\_95.465%  
 Lachnospiraceae\_[G-14] bacterium\_MOT-184\_nov\_95.405%  
 Anaerotignum lactatifermentans\_nov\_97.523%  
 Roseburia faecis  
 Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_92.208%  
 Lawsonibacter asaccharolyticus\_nov\_97.973%  
 Lachnospiraceae\_[G-11] bacterium\_MOT-177\_nov\_97.523%  
 Lachnospiraceae\_[G-14] bacterium\_MOT-184\_nov\_94.989%  
 Bacteroides acidifaciens\_acidofaciens  
 Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_91.106%  
 Mediterraneibacter [Ruminococcus] gnavus\_nov\_93.424%  
 Oscillospiraceae\_[G-2] bacterium\_MOT-149\_nov\_95.281%  
 Anaerosporebacter mobilis\_nov\_95.000%  
 Oscillospiraceae\_[G-2] bacterium\_MOT-149  
 Oscillospiraceae\_[G-4] bacterium\_MOT-151\_nov\_91.723%  
 Muribaculum intestinale  
 Saccharofermentans acetigenes\_nov\_88.764%  
 Anaerotruncus rubiinfantis\_nov\_92.760%  
 Lachnospiraceae\_[G-6] bacterium\_MOT-171\_nov\_96.606%  
 Lachnospiraceae\_[G-11] multispecies\_sppn4\_2\_nov\_96.847%  
 multigenus multispecies\_sppn9\_2\_nov\_93.002%  
 Bacteroides acidofaciens  
 Kineothrix alysoides\_nov\_93.651%  
 Bacteroides multispecies\_sppn6\_2\_nov\_96.312%  
 Lachnospiraceae\_[G-11] bacterium\_MOT-178  
 Oscillospiraceae\_[G-2] bacterium\_MOT-149\_nov\_95.946%  
 Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_91.991%  
 Phocaea massiliensis\_nov\_95.682%  
 Parabacteroides goldsteinii  
 Alistipes multispecies\_sppn12\_2\_nov\_96.304%  
 Blautia caecimuris\_nov\_96.825%  
 Bacteroides caecimuris  
 Eubacterium coprostanoligenes\_nov\_95.485%  
 Alistipes sp.\_MOT-127  
 Oscillospiraceae\_[G-2] bacterium\_MOT-149\_nov\_95.506%  
 Mithella massiliensis\_nov\_92.094%  
 Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_91.974%  
 Oscillospiraceae\_[G-2] bacterium\_MOT-149\_nov\_95.056%  
 Bacteroides stercorisoris  
 Prevotella sp.\_MOT-128  
 Faecalicatena orotica\_nov\_95.238%  
 Lachnospiraceae\_[G-12] bacterium\_MOT-179\_bacterium\_MOT-18  
 Bacteroidetes\_[G-3] multispecies\_sppn2\_2\_nov\_87.554%  
 Eubacteriales\_[G-1] bacterium\_MOT-158  
 Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_92.191%  
 Duncaniella freteri\_nov\_90.456%  
 Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_93.074%  
 Alistipes putredinis\_nov\_95.887%  
 Lachnospiraceae\_[G-10] bacterium\_MOT-175\_nov\_92.174%  
 multigenus multispecies\_sppn8\_3\_nov\_95.011%  
 Lachnospiraceae\_[G-9] bacterium\_MOT-174  
 Oscillibacter valericigenes\_nov\_95.260%  
 Pseudoflavonifractor capillosus\_nov\_95.721%  
 Muribaculaceae\_[G-2] bacterium\_MOT-104\_nov\_89.462%  
 Lacrimispora xylanolytica\_nov\_97.285%  
 Alistipes putredinis\_nov\_96.529%  
 Lachnospiraceae\_[G-10] bacterium\_MOT-175\_nov\_96.372%  
 multigenus multispecies\_sppn7\_2\_nov\_92.777%  
 Lachnospiraceae\_[G-12] bacterium\_MOT-171\_nov\_95.238%  
 Lachnospiraceae\_[G-12] bacterium\_MOT-179\_nov\_92.534%  
 Lachnospiraceae\_[G-14] bacterium\_MOT-184  
 Lachnospiraceae\_[G-14] bacterium\_MOT-183  
 Lacrimispora indolis\_nov\_90.724%  
 Acetatifactor muris\_nov\_92.551%  
 Muribaculum intestinale\_nov\_93.737%  
 Lachnospiraceae\_[G-11] bacterium\_MOT-176\_nov\_97.297%  
 Kineothrix alysoides\_nov\_95.928%  
 Akkermansia muciniphila  
 Prevotellamasilia timonensis\_nov\_94.168%  
 Duncaniella freteri\_nov\_93.103%  
 Blautia caecimuris

Species



F8810.S10  
 F8810.S11  
 F8810.S12  
 F8810.S22  
 F8810.S23  
 F8810.S24

Samples

