

Species

SP123	Celobacterium somerae	SP7	Neisseriaceae [G-1] bacterium_MOT-031	SPN332	Duricardia freteri nov_89.069%	SPN333	Alistipes senegalensis nov_93.089%
SP124	Trepnema denticola	SP70	Muribaculaceae [G-1] bacterium_MOT-129	SPN334	Faecalicatena orotica nov_92.484%	SPN335	Oscillospiraceae [G-3] bacterium_MOT-150_nov_92.931%
SP126	Lactobacillus johnsonii	SP71	Stenotrophomonas [Pseudomonas] hibiscicola	SPN336	Lachnospiraceae [G-6] bacterium_MOT-171_nov_93.515%	SPN337	Actinomyces sp._HMT_169_nov_97.992%
SP127	Enterobacter hormaechei	SP73	Parabacteroides goldsteinii	SPN338	Saccharibacteria [TM7] [G-3] bacterium_HMT_351_nov_94.411%	SPN339	Eubacterium coprostanoligenes nov_91.631%
SP129	Streptococcus gordonii	SP74	Acidovorax temperans	SPN340	Natroniella acetigena nov_80.992%	SPN341	Muribaculaceae [G-1] bacterium_MOT-129_nov_84.929%
SP13	Streptococcus oralis	SP75	Alloprevotella sp._HMT_473	SPN342	Muribaculaceae [G-2] bacterium_MOT-104_nov_88.623%	SPN343	Hydrogenoanaerobacterium saccharovorans nov_90.249%
SP130	Shigella flexneri	SP76	Rhodococcus qingshengii	SPN344	Lachnospiraceae [G-6] bacterium_MOT-171_nov_94.572%	SPN345	Phoece massiliensis nov_90.426%
SP132	Fusobacterium periodonticum	SP77	Faecalibaculum rodentium	SPN346	Neglectibacter timonensis nov_95.325%	SPN347	Butyrivibrio proteoclasticus nov_83.594%
SP133	Eubacteriales [G-1] bacterium_MOT-158	SP78	Bifidobacterium pseudolongum	SPN348	Eubacteriales [G-3] bacterium_MOT-163_nov_84.913%	SPN349	Muribaculaceae [G-1] bacterium_MOT-129_nov_86.089%
SP134	Bifidobacterium longum	SP79	Enterobacter cancerogenus	SPN350	Bacteroidetes [G-3] bacterium_HMT_436_nov_86.585%	SPN351	Duncaniella freteri nov_88.978%
SP135	Pseudoramibacter alactolyticus	SP8	Phocaeicola sartorii	SPN352	Lachnospiraceae [G-5] bacterium_MOT-170_nov_97.904%	SPN353	Phoece massiliensis nov_87.179%
SP136	Agromyces mediolanus	SP80	Lawsonella clevelandensis	SPN354	Oscillospiraceae [G-4] bacterium_MOT-151_nov_95.417%	SPN355	Lachnospiraceae [G-13] bacterium_MOT-181_nov_91.189%
SP137	Alloprevotella tannerae	SP81	Campylobacter sp._HMT_044	SPN356	Eisenbergiella massiliensis nov_84.867%	SPN357	Sporobacter termitidis nov_87.580%
SP138	Peptostreptococcaceae [XII][G-4] bacterium_HMT_369	SP82	Acidovorax ebreus	SPN358	Lachnospiraceae [G-11] bacterium_MOT-177_nov_88.296%	SPN359	Christensenella hongkongensis nov_86.308%
SP139	Lachnoclostridium [Clostridium] symbiosum	SP83	Alistipes sp._MOT-127	SPN360	Lachnospiraceae [G-14] bacterium_MOT-182_nov_90.254%	SPN361	Lachnoclostridium phytofermentans nov_90.417%
SP14	Streptococcus sp._HMT_064	SP84	Escherichia fergusonii	SPN362	Blautia marasmi nov_93.711%	SPN363	Muribaculaceae [G-1] bacterium_MOT-129_nov_83.636%
SP140	Bradyrhizobium pachyrhizi	SP91	Staphylococcus saccharolyticus	SPN364	Butyricoccus pullicaecorum nov_85.093%	SPN365	Selenomonas sp._HMT_137_nov_97.228%
SP141	Peptococcaceae [G-1] bacterium_MOT-146	SP92	Leptotrichia sp._HMT_219	SPN366	Mollicutes [G-2] bacterium_MOT-187_nov_95.481%	SPN367	Oscillospiraceae [G-6] bacterium_MOT-153_nov_83.582%
SP142	Peptidiphaga sp._HMT_183	SP93	Streptococcus cristatus clade_578	SPN368	Eubacterium xylanophilum nov_91.075%	SPN369	Ihubacter massiliensis nov_90.644%
SP143	Leucobacter chromiirestis	SP94	Muribaculaceae [G-2] bacterium_MOT-104	SPN370	Prevotella shahii nov_87.602%	SPN371	Anaerotruncus rubiinfantis nov_92.500%
SP144	Acinetobacter lwoffii	SP95	Streptococcus chosunense	SPN372	Eubacteriales [G-3] bacterium_MOT-163_nov_85.944%	SPN373	Duncaniella freteri nov_89.379%
SP145	Deinococcus geothermalis	SP96	Blautia hominis	SPN374	Anaerotignum faecicola nov_85.396%	SPN375	Lachnospiraceae [G-1] bacterium_MOT-166_nov_95.643%
SP146	Selenomonas flueggei	SP97	Enterococcus casseliflavus	SPN376	Duncaniella freteri nov_85.685%	SPN376	Glucerbacter canis nov_93.305%
SP147	Catonella sp._HMT_164	SP98	Streptococcus sanguinis	SPN377	Eubacteriales [G-4] bacterium_MOT-164_nov_92.373%	SPN378	Parafannyhessea umbonata nov_92.161%
SP148	Prevotella nigrescens	SPN100	Duncaniella freteri nov_91.853%	SPN380	Veillonellaceae [G-1] bacterium_HMT_129_nov_96.356%	SPN381	Adlercreutzia muris nov_97.186%
SP149	Adlercreutzia caecimuris	SPN107	Muribaculaceae [G-2] bacterium_MOT-104_nov_85.686%	SPN382	Oscillospiraceae [G-2] bacterium_MOT-149_nov_94.802%	SPN383	Lachnospiraceae [G-6] bacterium_MOT-171_nov_91.423%
SP15	Acinetobacter johnsonii	SPN108	Anaeromassilibacillus senegalensis nov_93.390%	SPN384	Hydrogenoanaerobacterium saccharovorans nov_88.797%	SPN385	Kineothrix alysoides nov_86.667%
SP151	Filifactor alovis	SPN109	Muribaculaceae [G-1] bacterium_MOT-129_nov_89.431%	SPN390	Lawsonibacter asaccharolyticus nov_90.329%	SPN42	Lachnoclostridium [Clostridium] aminophilum nov_87.318%
SP152	Mucispirillum schaedleri	SPN11	Lacrimispora xylanolytica nov_93.789%	SPN44	Lachnospiraceae [G-1] bacterium_MOT-129_nov_87.576%	SPN45	Duncaniella freteri nov_92.653%
SP156	Methylobacterium brachiatum	SPN118	Duncaniella freteri nov_90.184%	SPN131	Duncaniella freteri nov_92.915%	SPN5	Duncaniella freteri nov_86.290%
SP157	Atlantibacter hermannii	SPN120	Lachnospiraceae [G-7] bacterium_MOT-172_nov_93.279%	SPN132	Leptotrichia hofstadii nov_96.970%	SPN52	Chryseobacterium yeoncheonense nov_97.484%
SP158	Streptococcus acidominimus	SPN126	Alloprevotella sp._HMT_473_nov_89.366%	SPN137	Muribaculaceae [G-1] bacterium_MOT-129_nov_86.640%	SPN53	Pyrinomonas methylaliphatogenes nov_96.809%
SP159	Streptococcus oralis subsp._denticani_clade_058	SPN13	Muribaculaceae [G-1] bacterium_MOT-129_nov_87.576%	SPN142	Marinisporobacter balticus nov_82.692%	SPN58	Alloprevotella sp._HMT_473_nov_89.634%
SP16	Haemophilus haemolyticus	SPN14	Muribaculaceae [G-2] bacterium_MOT-104_nov_89.621%	SPN149	Alistipes senegalensis nov_93.648%	SPN64	Muricomes intestini nov_89.583%
SP161	Aggregatibacter aphrophilus	SPN149	Alistipes senegalensis nov_93.648%	SPN153	Anaerostipes caccae nov_96.328%	SPN67	Muribaculaceae [G-2] bacterium_MOT-104_nov_86.373%
SP164	Corynebacterium aurimucosum	SPN156	Alistipes putredinis nov_94.444%	SPN161	Lachnospiraceae [G-3] bacterium_MOT-168_nov_92.902%	SPN71	Muribaculaceae [G-2] bacterium_MOT-104_nov_88.822%
SP165	Streptococcus oralis subsp._tigurinus_clade_070	SPN161	Lachnospiraceae [G-3] bacterium_MOT-168_nov_92.902%	SPN164	Duncaniella freteri nov_90.612%	SPN75	Lachnospiraceae [G-11] bacterium_MOT-178_nov_89.293%
SP166	Streptococcus mutans	SPN164	Duncaniella freteri nov_90.612%	SPN170	Erysipelatoclostridium [Clostridium] innocuum nov_88.270%	SPN86	Lachnoclostridium [Clostridium] polysaccharolyticum nov_92.083%
SP168	Butyrivibrio sp._HMT_080	SPN174	Muribaculaceae [G-1] bacterium_MOT-129_nov_86.427%	SPN174	Muribaculaceae [G-1] bacterium_MOT-129_nov_86.427%	SPN88	Duncaniella freteri nov_91.039%
SP170	Aggregatibacter sp._HMT_512	SPN175	Alistipes finegoldii nov_93.608%	SPN185	Hathewayia proteolytica nov_83.514%	SPN92	Duncaniella freteri nov_91.429%
SP171	Johnsonella sp._HMT_166	SPN185	Hathewayia proteolytica nov_83.514%	SPN191	Duncaniella freteri nov_93.699%		
SP172	Oscillospiraceae [G-6] bacterium_MOT-153	SPN191	Duncaniella freteri nov_93.699%	SPN194	Muribaculaceae [G-1] bacterium_MOT-129_nov_90.816%		
SP176	Prevotella sp._HMT_304	SPN196	Oscillospiraceae [G-2] bacterium_MOT-149_nov_95.198%	SPN196	Oscillospiraceae [G-2] bacterium_MOT-149_nov_95.198%		
SP177	Lachnospiraceae [G-3] bacterium_MOT-168	SPN199	Beduini massiliensis nov_87.705%	SPN206	Muribaculaceae [G-2] bacterium_MOT-104_nov_86.600%		
SP179	Desulfobulbus sp._HMT_041	SPN207	Phoece massiliensis nov_90.129%	SPN207	Phoece massiliensis nov_90.129%		
SP18	Janibacter melonis	SPN210	Duncaniella freteri nov_87.424%	SPN210	Duncaniella freteri nov_87.424%		
SP180	Prevotella melaninogenica	SPN217	Acutalibacter muris nov_96.289%	SPN217	Acutalibacter muris nov_96.289%		
SP181	Actinomyces sp._HMT_896	SPN218	Oscillospiraceae [G-3] bacterium_MOT-150_nov_93.582%	SPN218	Oscillospiraceae [G-3] bacterium_MOT-150_nov_93.582%		
SP185	Dialister pneumosintes	SPN22	Muribaculaceae [G-2] bacterium_MOT-104_nov_89.400%	SPN22	Muribaculaceae [G-2] bacterium_MOT-104_nov_89.400%		
SP186	Corynebacterium matruchotii	SPN220	Longibaculum muris nov_86.957%	SPN220	Longibaculum muris nov_86.957%		
SP187	Ruthenibacterium lactatiformans	SPN228	Parabacteroides distasonis nov_97.938%	SPN228	Parabacteroides distasonis nov_97.938%		
SP189	Clostridium disporicum	SPN229	Flavonifractor plautii nov_92.308%	SPN229	Flavonifractor plautii nov_92.308%		
SP19	Limosilactobacillus reuteri	SPN23	Duncaniella freteri nov_86.842%	SPN23	Duncaniella freteri nov_86.842%		
SP191	Schaalia sp._HMT_178	SPN238	Muribaculaceae [G-1] bacterium_MOT-129_nov_86.290%	SPN238	Muribaculaceae [G-1] bacterium_MOT-129_nov_86.290%		
SP192	Thermicanus aegyptius	SPN240	Blautia luti nov_94.561%	SPN240	Blautia luti nov_94.561%		
SP194	Duncaniella freteri	SPN242	Muribaculaceae [G-2] bacterium_MOT-104_nov_88.822%	SPN242	Muribaculaceae [G-2] bacterium_MOT-104_nov_88.822%		
SP195	Staphylococcus warneri	SPN246	Eubacteriales [G-4] bacterium_MOT-164_nov_97.655%	SPN246	Eubacteriales [G-4] bacterium_MOT-164_nov_97.655%		
SP196	Rothia dentocariosa	SPN249	Duncaniella freteri nov_88.330%	SPN249	Duncaniella freteri nov_88.330%		
SP197	Actinomyces sp._HMT_448	SPN254	Lachnospiraceae [G-7] bacterium_MOT-172_nov_91.718%	SPN254	Lachnospiraceae [G-7] bacterium_MOT-172_nov_91.718%		
SP198	Leptotrichia buccalis	SPN256	Duncaniella freteri nov_87.071%	SPN256	Duncaniella freteri nov_87.071%		
SP20	Akkermansia muciniphila						
SP201	Actinomyces sp._HMT_414						
SP21	Alishewanella agri						
SP22	Streptococcus sinensis						
SP23	Kosakonia sacchari						
SP24	Erysipelotrichaceae [G-1] bacterium_MOT-189						
SP25	Porphyromonas endodontalis						
SP26	Cuthbertium acnes						