| **Taxonomic rank** | **Taxon nomenclature** | **Confidence** | **Assigned sequence ID** |
| --- | --- | --- | --- |
| Phylum | Firmicutes | 99.91% | 253da3f2003c66ffda7b498233410fff |
| Phylum | Firmicutes | 96.10% | 7fbe53c790cbcd631feb2385259f3d7c |
| Phylum | Firmicutes | 70.32% | 7a33a753e13714f569153a2aea6ba0da |
| Phylum | Firmicutes | 71.46% | f3e123f539d5e059ea469966e93ee1d2 |
| Phylum | Firmicutes | 85.43% | 1ea10acc964b92acd02d2cc710c7f1a9 |
| Phylum | Planctomycetota | 99.75% | 11dbebac87effb545c46f12fc9bf1bdf |
| Class | Actinobacteria | 100.00% | efa9ce315ac589dc6ade7b9ebac4369c |
| Class | Actinobacteria | 100.00% | 6b17b05ca45be96f2b55bdf906a5add9 |
| Class | Actinobacteria | 100.00% | 65e535c36d6521a4ea0fe901286b7576 |
| Class | Actinobacteria | 100.00% | 9eda2a27ded06ad4b4d06f2f711f0994 |
| Class | Actinobacteria | 100.00% | a263c18d4a4de4e0cf135867d44f2690 |
| Class | Alphaproteobacteria | 100.00% | 4a0568cf762a6322dd5a616966a1a087 |
| Class | Alphaproteobacteria | 72.15% | 71dc96303b95e7accb97a47b67bc5121 |
| Class | Bacilli | 73.30% | 29c6eed065ea39b8cdbadabc9c313ce7 |
| Class | Bacilli | 89.69% | fc7b56c6486b86a0bdcd06d9411a2104 |
| Class | Bacilli | 99.95% | 3530bdff4eb5e9a58f504d0aabc09a9c |
| Class | Bacilli | 99.51% | 9cc4f199a38f66b5f99ea9bcf7e826df |
| Class | Bacteroidia | 76.46% | 3a22f07584cb527c88dad2699b3ccd7f |
| Class | Bacteroidia | 93.05% | 501e399249cdc78810dd1295a01d8283 |
| Class | Bacteroidia | 99.91% | b0acedf596553995dd3f29698a1b43f6 |
| Class | Bacteroidia | 95.99% | 396fa5cfca955a6fe00c4e159b3a28ff |
| Class | Clostridia | 71.06% | e54925a3e2a45f33f4786ea409e6f944 |
| Class | Clostridia | 99.98% | 960bee4448d441982de56dd40dc27e61 |
| Class | Clostridia | 99.86% | aa9f612cc2f06078f6546645e0c75d35 |
| Order | Bacillales | 99.99% | 146c009a9b43159681cab218664cd33a |
| Order | Bacteroidales | 90.47% | f10f746b408c288c444287c82f1065fd |
| Order | Bacteroidales | 99.96% | 1b5e41869c65fa5b95e774b1181b24a5 |
| Order | Bacteroidales | 100.00% | 484f5d212e192cba5a911d7d58049c91 |
| Order | Bacteroidales | 98.10% | 5903d79f355344d6791225ea8db53d51 |
| Order | Bacteroidales | 99.99% | 909d61d2bb4f182fce9de92daa6f7f6e |
| Order | Bacteroidales | 88.70% | 99d77514fa3ad18fc3d0f8b80c6c8381 |
| Order | Bacteroidales | 100.00% | 647cd5675b6d460a1cb5f9bdd95c4ddb |
| Order | Bacteroidales | 93.44% | d3f8b1fd896eeebbe33eaeed7652c026 |
| Order | Burkholderiales | 91.66% | 0d7dd08feac8112c2f4b62848259d7ba |
| Order | Burkholderiales | 99.91% | 4b9907d0d7360ba0e165a57f821efb5a |
| Order | Burkholderiales | 82.37% | 18e647f8b0de1de1ec4c72c4e246e7fd |
| Order | Burkholderiales | 95.96% | 28772d0fbad7418208c159c8d079c4b2 |
| Order | Burkholderiales | 95.33% | e88798c66675796257cf451be8e31f37 |
| Order | Burkholderiales | 80.80% | 606b9e5879fd864e9239ab608b357a99 |
| Order | Corynebacteriales | 100.00% | 7e699578d9780020f65ecda24cf0fdd2 |
| Order | Corynebacteriales | 100.00% | 383bc92c177f0122ab732ed319c22a8f |
| Order | Corynebacteriales | 100.00% | 71836fdd193fe28dead40a8a142c14f9 |
| Order | Cyanobacteriales | 87.67% | 93f1a18a772918bd3d213d1512a35c79 |
| Order | Enterobacterales | 77.00% | 4d4d335a27350cfd004b50251de14285 |
| Order | Oscillospirales | 99.98% | 9cc4c4f184fd4c19e45477eb274ee1ad |
| Order | Oscillospirales | 99.98% | 198f5ede0084932df126c791f08c110d |
| Order | Oscillospirales | 99.99% | 8f2e1acf33d07bc3c1102a212d1ed84f |
| Order | Peptostreptococcales-Tissierellales | 99.42% | ede15ea5a91f1735ced6743546a12bcb |
| Order | Peptostreptococcales-Tissierellales | 99.38% | 2a47ead13657081c95cdf81226400cca |
| Order | Peptostreptococcales-Tissierellales | 99.53% | 1d5caaf93d7110be8beecc91174ccce2 |
| Order | Peptostreptococcales-Tissierellales | 90.25% | 815e19694c88b2d2579bdb3af3fe90f4 |
| Order | Peptostreptococcales-Tissierellales | 92.27% | 88708b317fdc27c32ef95bdac29d67ce |
| Order | Pseudomonadales | 99.99% | 706e47aa32d531e34ef4f935a1063ad4 |
| Order | Pseudomonadales | 99.99% | 5763850e98314c5982ed1035b6336026 |
| Order | Pseudomonadales | 99.94% | da4570babff7068127046567d6369a15 |
| Order | Pseudomonadales | 99.99% | e7eada8a3cabceca4e74a9f423806877 |
| Order | Rhizobiales | 100.00% | 7790628b89f1ecd074db2fe82ddee2c5 |
| Order | Rhizobiales | 100.00% | 87d392c2fde21132952563813a96c460 |
| Order | Saccharimonadales | 79.26% | a28932b4d963b654ebe06e3152ad935c |
| Order | Veillonellales-Selenomonadales | 99.99% | cc203b329f1d32b65ce6fcd0595efa6c |
| Family | Acetobacteraceae | 99.99% | 7a132550afcc1cde92f6efb622f0ae94 |
| Family | Acetobacteraceae | 99.99% | c667c37d92b5535a5ed8c83fbe7bff84 |
| Family | Acetobacteraceae | 100.00% | 18fb27931a7b9ce56bd00f72882093a4 |
| Family | Anaerovoracaceae | 99.13% | 222c97f2c4304944dff5c5f23090e19c |
| Family | Anaerovoracaceae | 87.65% | e7b5c9127d038fb62014fec7f7f0ad67 |
| Family | Bacillaceae | 92.84% | 64bf96af3d9ce29f6ff964c61b44aaeb |
| Family | Bacillaceae | 80.09% | 3d4feeaa282af9c22113f495ce4fda1c |
| Family | Beijerinckiaceae | 84.34% | 8a9b3caa4db1cf39b21510aa1c7c52f8 |
| Family | Beijerinckiaceae | 98.06% | 3150f5f6c7cf0b3b2200bf138a85d7e4 |
| Family | Beijerinckiaceae | 89.19% | bfc18a8bdbfd91db176188aa53fcdc75 |
| Family | Beijerinckiaceae | 74.17% | ae478c9810e0e3e77b697bf3864dd44f |
| Family | Chitinophagaceae | 100.00% | 690df2a02aeb7ea9eaaef59a86daca50 |
| Family | Chitinophagaceae | 98.99% | c40e14e3704d29d1340405272d5d9300 |
| Family | Clostridiaceae | 90.23% | e70c0a260c2701906f60e18716f56548 |
| Family | Comamonadaceae | 99.80% | 257a1243bdd05af46c7ee13c115c6434 |
| Family | Comamonadaceae | 98.60% | 32df36ad88aad6935a9a9755642ccf76 |
| Family | Dysgonomonadaceae | 77.20% | 1801eeb5e5b51f925728d2e8c4fa7879 |
| Family | Dysgonomonadaceae | 75.39% | 49cde44a92721d7b01d40f3e096b4b1c |
| Family | Dysgonomonadaceae | 73.50% | 32c3ff2e3375315c4be9eec811880a23 |
| Family | Lachnospiraceae | 100.00% | 6d8843cc13b2c7df43ef15fead3e993a |
| Family | Lachnospiraceae | 99.99% | 00fedfdbdd1deb6f2d1a0875e588f4d4 |
| Family | Lachnospiraceae | 100.00% | 373ab3522292a3f4d24fbffd6c574be9 |
| Family | Lachnospiraceae | 98.09% | a2338591cbc926fee73cdcba125e1ca7 |
| Family | Lachnospiraceae | 100.00% | f34c4987b03016ed8bd9a8722777b3c6 |
| Family | Lachnospiraceae | 99.97% | f3b748b5b184b6b5ee63916374c7843a |
| Family | Methyloligellaceae | 90.73% | 5aeeedcc346d51a5864777810c3366de |
| Family | Microbacteriaceae | 97.16% | e2dd8a22765600f618d6a225c02ca550 |
| Family | Microbacteriaceae | 93.37% | fe0afa86ff31aef0112b6b15981c3a9d |
| Family | Microbacteriaceae | 85.43% | 352c07a8447f9719a7c4a458f4f67bf2 |
| Family | Microbacteriaceae | 95.42% | 9d91a387896e7d64733a6a730d09d73d |
| Family | Microbacteriaceae | 87.66% | fee488f23b5a9456394b8035f511b796 |
| Family | Microbacteriaceae | 97.54% | b1f85723da8ba81075e07597d57cc8fa |
| Family | Microbacteriaceae | 76.43% | 69762327dde8d91ed7521f52baf63ea3 |
| Family | Microbacteriaceae | 74.25% | 0268e70b929405a02e50d026609a8373 |
| Family | Microbacteriaceae | 96.37% | 06274452709107af1e637eff724e5db1 |
| Family | Microbacteriaceae | 96.08% | 97cd98e1fd4fd91d33cfe82f4e8317dc |
| Family | Micrococcaceae | 97.00% | f844f7557efbabb489aa51ae6658ca4d |
| Family | Neisseriaceae | 99.47% | 3b942627249a38b6cd63640cd9b93ef5 |
| Family | Neisseriaceae | 99.23% | a0edab59f2a8c9bf65805794f5483cd6 |
| Family | Neisseriaceae | 99.69% | 91b24afbcc8c6f2131124b3c1f5afa75 |
| Family | Orbaceae | 99.71% | f5b6d0c1d9b774c9e724f3a2838bad76 |
| Family | Orbaceae | 99.71% | 7e76798fd0739209c546df4bb3d9d16a |
| Family | Orbaceae | 73.43% | b74043a198c51bf850e71ed4ff5d9782 |
| Family | Oscillospiraceae | 77.57% | 7865765e93e7ad9ef770967e37c773ca |
| Family | Pasteurellaceae | 99.98% | b39b26b8f2c9e77d7fface3175a97a94 |
| Family | Peptostreptococcaceae | 100.00% | 0648a9113ab225823e27ba206b3dfe32 |
| Family | Rhizobiaceae | 100.00% | 2ddc2ecff882ad2074de7bffe8071c69 |
| Family | Rhodocyclaceae | 76.25% | 14f3fcb2ce38e4cdda8df1078ae02db6 |
| Family | Rhodocyclaceae | 80.78% | 54bb585ec883aeff630f449f66c0f573 |
| Family | Ruminococcaceae | 93.39% | e3bff8d84c49163ed4759982a5dbe56e |
| Family | Ruminococcaceae | 99.56% | 0c286d525e1c4620dad1b09eb6eef64c |
| Family | Ruminococcaceae | 81.95% | 55b2e23d7d141891085e10297ddf573f |
| Family | Ruminococcaceae | 99.58% | 578ef62a530d2d281693c563330a8923 |
| Family | Ruminococcaceae | 97.88% | c9af8d1b236646c43d4f08a10f0a7660 |
| Family | Saccharimonadaceae | 81.91% | 9fce97e6139db1f0c0932e01a5d972ca |
| Family | Sandaracinaceae | 74.86% | f292658fb7e2524cca771808c3062abc |
| Family | Sphingomonadaceae | 82.60% | ca1e70bcf21190ca7924886e9b0ddcc0 |
| Family | Sutterellaceae | 72.04% | d0ebe3a949ee33a863de4cd928fc7909 |
| Family | Sutterellaceae | 74.23% | 44bb9ec517942ed8c82ce24e3890f111 |
| Family | Tannerellaceae | 99.99% | e5c61a1521d944dc477589e9ce5a2236 |
| Family | Weeksellaceae | 92.10% | 964ea87c411631553e928a2406b1bb89 |
| Family | Weeksellaceae | 92.10% | 90acce5030dc9fb4130b80e0d538347b |
| Family | Weeksellaceae | 100.00% | ed22fd63f4c809514b80cd549ba0515e |
| Family | Weeksellaceae | 99.99% | 32bf7d0c4687d594c369853e8fadc54f |
| Family | Weeksellaceae | 73.82% | f22c01a5efcbfbd3b0c882e1c70ec47d |
| Family | Weeksellaceae | 92.58% | 20d43041362dcad8949e18f870e2f79d |
| Family | Weeksellaceae | 90.48% | 34fa726650e74f6ef85823568061ff32 |
| Family | Weeksellaceae | 89.81% | 0592028b5f9e7551f23ceb7f2fce601d |
| Family | Williamwhitmaniaceae | 99.28% | 05c76092a706fbf3b8dba19de355faf5 |
| Family | Wohlfahrtiimonadaceae | 100.00% | 680b454926edae42a844729b12bdbd78 |
| Family | Wohlfahrtiimonadaceae | 100.00% | d910e2c373fc2fcff28b6d554de6f46f |
| Family | Wohlfahrtiimonadaceae | 100.00% | b06e1f8e14ceb64ec7d8b056f7cd1e51 |
| Family | Wohlfahrtiimonadaceae | 100.00% | 465fe775b0a01383fe5e4c8a8de6e230 |
| Family | Xanthobacteraceae | 99.54% | 1588ef3f5b0996361691ccbf13a9a0a0 |
| Family | Xanthomonadaceae | 87.93% | 12e16b0083b3698ef9c6bd2b902e21cf |
| Genus | Clostridium innocuum group | 99.96% | 414b92eba277238e76b12c687fc90cbd |
| Genus | Clostridium innocuum group | 99.85% | a2abb2730495e527c36c3a18c2f9ccdb |
| Genus | Abditibacterium | 99.79% | a8b1c40d5e7b6612d50d9e9eddc21581 |
| Genus | Abditibacterium | 79.54% | 42e510570ccfbdb27713acfe4fc66c6d |
| Genus | Abditibacterium | 100.00% | 32056940e4c843906baf38572bef891c |
| Genus | Abditibacterium | 99.92% | b9506b516407519482da7a7f814f8774 |
| Genus | Acholeplasma | 74.07% | 9dd6b6cf482a592935e53cfd33993f25 |
| Genus | Achromobacter | 73.27% | 8cb783bb8a374250cad8f28889b158c9 |
| Genus | Acinetobacter | 99.09% | 7bd362a445b5a7ef8c24818b0963c7e4 |
| Genus | Acinetobacter | 98.47% | 18387e9cf1f471d949814462f1f3e43b |
| Genus | Acinetobacter | 98.94% | 0882ed2f3b40de8d8c1319b94074fd73 |
| Genus | Acinetobacter | 99.11% | 8f28784533cde436c39673c5b55cefd5 |
| Genus | Acinetobacter | 100.00% | 1b01b7a711ee4a484960b5d189521b28 |
| Genus | Akkermansia | 99.67% | 051c594e7ca64f7414a75016500ef30a |
| Genus | Alistipes | 100.00% | a76a007d7c92707d153c4433606ae339 |
| Genus | Alistipes | 100.00% | 758e4d0cb9b9de8e32b12e6228c1ee99 |
| Genus | Alkanindiges | 96.99% | e0ed4f477d969e8c6a20b4d6da36ce58 |
| Genus | Anaerolinea | 94.86% | 4bcee44d376eb2c03d5a201c83b4f506 |
| Genus | Arthrobacter | 77.72% | 4cee0c7e19503e767114a210e4abd6e6 |
| Genus | Aurantisolimonas | 71.48% | 4b110641915c04315f236af262550974 |
| Genus | Bacteroides | 99.99% | 5e188cd382c47fd8a7abadf2f47c9b99 |
| Genus | Bacteroides | 100.00% | c41290aa09743e6c4f369ae7f1cfee19 |
| Genus | Bacteroides | 100.00% | d96ee4ebcd458f092546c5b97755bb3b |
| Genus | Bacteroides | 99.53% | a6c13e16024131345b05007716792f50 |
| Genus | Bacteroides | 100.00% | 933f5136a5cca1b98fae46e5c5d99683 |
| Genus | Bacteroides | 100.00% | 18eda008fc98bd54a9b9bd10a18b4fea |
| Genus | Bacteroides | 99.99% | a7e992401860b4b0c37491ffb0976f79 |
| Genus | Bacteroides | 100.00% | 1db1711d9a1d7e897ee874fff05fbc3d |
| Genus | Bacteroides | 100.00% | cdf13cca19d9f8175b066b8846ae6eda |
| Genus | Bacteroides | 100.00% | 5cba65b9b9d17bb7f130073ded039473 |
| Genus | Bacteroides | 100.00% | 23611d31ad49cffe0b20b2fa9931aca4 |
| Genus | Bacteroides | 99.99% | c8d9df2433f193acc7f752c205f35614 |
| Genus | Bacteroides | 98.81% | 22db4b09d645312b71a5107f53e71be8 |
| Genus | Bacteroides | 99.74% | 86dfc62fc898213cb03251d110ffc226 |
| Genus | Bacteroides | 100.00% | d5a2371fac20c903880c092a4aabae06 |
| Genus | Bacteroides | 99.53% | 292bd05b6cbcb3ed898c552c2e7eec26 |
| Genus | Bacteroides | 99.69% | 287449441f2df1d221e571eeb4635578 |
| Genus | Bacteroides | 100.00% | eda905e5483c24a1bc3b4157d5284225 |
| Genus | Bacteroides | 99.54% | b9dc4a70dd2cccba4a054abfff63b1a2 |
| Genus | Bacteroides | 100.00% | a095ad1f8a168d65e5b35ed1d31a6c73 |
| Genus | Bacteroides | 100.00% | d3510f58db56383e0c60b55f0374ff4e |
| Genus | Bacteroides | 99.64% | 926e6c460261607f5ed84d208ecc431c |
| Genus | Bacteroides | 99.54% | 8340355293c3b3db2e3796f867f14613 |
| Genus | Bacteroides | 100.00% | 158f8a976142c01d0613d520c266e26c |
| Genus | Bacteroides | 99.99% | c8d6611db6c36a061e7eaceef6630fed |
| Genus | Bacteroides | 100.00% | 7b64b71541e360493144cf77aa00e44d |
| Genus | Bacteroides | 100.00% | ab1d20253205afc4550550b509be7e27 |
| Genus | Bacteroides | 100.00% | acf9fc36df7011a88f5dcf52ec5932dc |
| Genus | Bacteroides | 100.00% | bd72ffb15bce38631319ee1136310731 |
| Genus | Bacteroides | 100.00% | d3deeb017a9596b51b5cf88f34f014c9 |
| Genus | Bacteroides | 99.97% | bf383e6f51525e128f953bc29be72f70 |
| Genus | Bacteroides | 100.00% | 114a27fc1bd54ac6eee0b0e588ade19f |
| Genus | Bacteroides | 99.60% | a21ef53629bfb746f7d1439df0b2f4a7 |
| Genus | Bacteroides | 99.74% | 9ea1561a0632fc6ac8077aa0abf9cc18 |
| Genus | Bacteroides | 98.77% | 310792c5a7a0db4922cb1ee7f050165a |
| Genus | Bdellovibrio | 98.34% | 66f2b76c265f049ebf0766766fe7fce0 |
| Genus | Belnapia | 81.66% | 7a8f821d11ff91d3e921ff73ccab97b7 |
| Genus | Brachybacterium | 99.58% | 032fb7be4a3b9bcd2f5129a3d6914601 |
| Genus | Brachybacterium | 99.66% | f0b76d3dc006662c65c77ba934a3ae21 |
| Genus | Brachybacterium | 99.65% | 7596489a8f7cf7913e0e379658cc8fd2 |
| Genus | Brevundimonas | 99.98% | c5692042e40b7c07ffdf88d1b34e2444 |
| Genus | Brevundimonas | 99.34% | 2f781c54d85b54711c326d8de4698c0d |
| Genus | Butyricimonas | 88.92% | 6cb8d0aa2bd317b56a60d9956dc4871e |
| Genus | Cerasicoccus | 97.89% | 84c3d3b75a03636048a79413fa608178 |
| Genus | Chitinophaga | 98.05% | a772833519ccea08ee25dd0245c978d0 |
| Genus | Christensenellaceae R-7 group | 80.44% | 21e9bbb33812178f353b8b0055d4b41e |
| Genus | Christensenellaceae R-7 group | 92.67% | f5cbb6744f6e0a7e4dd22914fa739522 |
| Genus | Chryseobacterium | 99.99% | 16910584212df2dda3c68970a6af9d96 |
| Genus | Chthoniobacter | 99.68% | e42283c39c6ad4ead261cfa1f50c4c86 |
| Genus | Comamonas | 99.80% | f3fa144a2da7a593ff95c7d76de90e66 |
| Genus | Comamonas | 99.52% | fb9e987bede20b32e6c5ec64faa60947 |
| Genus | Comamonas | 99.90% | d16dc578f0226ea2cad1a86fdec3bfc3 |
| Genus | Comamonas | 98.09% | f2fd12da9bb31a70242fb2f7717426f0 |
| Genus | Comamonas | 99.15% | 2e51dc22cda8d25c1e26f06911709988 |
| Genus | Comamonas | 95.69% | 3bce9c35a120acdb24adaafafa5e83e0 |
| Genus | Comamonas | 99.72% | 4b43b948c89fc07c3c7d4c185a2b5cf0 |
| Genus | Comamonas | 94.85% | 2f4884f17d0d89983aae13eb2725b347 |
| Genus | Cutibacterium | 99.89% | f4f44aed827e8f00132c866312e51673 |
| Genus | Cutibacterium | 96.78% | a3b16882af684210c244602e7414de41 |
| Genus | Cutibacterium | 96.94% | 13d05bd3516320874529c76aba55a25d |
| Genus | Cutibacterium | 96.78% | 1ceb8120b377df8daec6164b58c85d19 |
| Genus | Deinococcus | 100.00% | 9162ba36abe7b0974378cb73a1591c9e |
| Genus | Deinococcus | 100.00% | 104054faa0fdcaf8b823d12327a6d062 |
| Genus | Desulfomonile | 93.27% | be54d8139445f3102db25822c137d0d1 |
| Genus | Desulfovibrio | 70.58% | 81baa8bd7dd144b6a66acc0ab2544ad0 |
| Genus | Desulfovibrio | 82.93% | 4f1fc441a0a4d1e1c981055de7ad98f2 |
| Genus | Desulfovibrio | 97.28% | e550d2bef5bcd847433f4a3ecc16ee28 |
| Genus | Desulfovibrio | 93.60% | ae29f03f5e6bd5ca6acfcb52376440fa |
| Genus | Desulfovibrio | 96.98% | c36758e304004381f4eb31ee775610ae |
| Genus | Desulfovibrio | 97.18% | 3ea9ab8ca78c2a1bf51f00e6864022a5 |
| Genus | Desulfovibrio | 97.89% | 567b3ad5d63585acfddea51cbc83c3dd |
| Genus | Desulfovibrio | 76.23% | 13a5e5a7bb05aef08c74a39d4aae3679 |
| Genus | Desulfovibrio | 99.96% | efe2f098fac9ae9c7a642b1def6e43cd |
| Genus | Desulfovibrio | 99.99% | 12d78a003c34256dc06198e306ccaff3 |
| Genus | Desulfovibrio | 99.37% | 89c236e1f1b50df3eaa0e3a36263eea0 |
| Genus | Dickeya | 99.17% | f7bbc998592423dd9904210b6774b0d4 |
| Genus | Dickeya | 98.92% | 12e9e4f6fb389c4290f0ee863f8c113c |
| Genus | Dickeya | 98.97% | 5816a5a31b0721f3d9182a4b46bf0233 |
| Genus | Dickeya | 99.18% | 1a1b0b41f134baf3d27c04430449587e |
| Genus | Dickeya | 99.07% | b190b0eeb47f70ff5bd5015f8b7e0325 |
| Genus | Dickeya | 98.98% | 63b6c1b4693668853e1c49bba7dc633d |
| Genus | Dickeya | 98.92% | 5c6288e670d73c94afab9f276cd9d10d |
| Genus | Dickeya | 98.80% | ed1cea3f2bff616a394a529cf51ef7d1 |
| Genus | Dickeya | 98.97% | 7aebc79c7d4b918604b94658655985b1 |
| Genus | Dickeya | 99.04% | b82f08b60b9c647f180d146400d10d6d |
| Genus | Dickeya | 98.85% | 09c826143d80b4df453f3d9edb89504a |
| Genus | Dickeya | 99.21% | b4258c5f3b98e0cbb7b8f14e199de9ee |
| Genus | Dickeya | 98.98% | d2e8a901416ed47c060c35eeb55fb524 |
| Genus | Dickeya | 98.85% | 46be1f393740e225b54dea4a56deab41 |
| Genus | Dickeya | 98.98% | 932c27e80104c0b6a1c710de59a00509 |
| Genus | Dickeya | 99.04% | 10c5b6ced4704399dd1b6cb6ac5b2d4c |
| Genus | Dickeya | 98.91% | ce5b1709104079bd5f848ab09f91f94d |
| Genus | Dickeya | 99.21% | f5feaba919eaf5cd786c6eb810f93c74 |
| Genus | Dickeya | 98.98% | 8b0a60ffbec2356a2b6c58d7442f1ff5 |
| Genus | Dickeya | 99.73% | 90bb43cdff07599e144986641e6d87ef |
| Genus | Dickeya | 97.38% | 8c7daa31f83011d21e741ed42524dfc0 |
| Genus | Dickeya | 96.35% | 87b6549fdc0d1539a9c0925137df2227 |
| Genus | Dickeya | 95.66% | d5824a7364ff6f1b27afa0abdcce917a |
| Genus | Dickeya | 95.68% | aaae3b8a244183500532a64530f9981d |
| Genus | Dickeya | 96.88% | 1fcd83f168af6be5b53af14ae1109076 |
| Genus | Dickeya | 99.94% | 2c5a44525aa567d58309bc325e1a2271 |
| Genus | Dickeya | 96.90% | 42f09af5b5656a4e9368caf96105939b |
| Genus | Dickeya | 97.37% | 156b73af5133ce1ea970cdb65d9c4ef0 |
| Genus | Dickeya | 96.86% | 9c9bca39f28dfe379725805dbf9ca5bf |
| Genus | Dickeya | 99.93% | 854c80c931bb31efad21179d6e906f73 |
| Genus | Dickeya | 99.92% | b5243d9b27c7e3342ccabf896e8e294e |
| Genus | Dickeya | 93.12% | 9f05bc4e4842d4044330b4f0a31c9ad4 |
| Genus | Dickeya | 93.05% | e897b63e724576b15e8fae9997375ac6 |
| Genus | Dickeya | 99.80% | 902af8e753f3d1e173877ec5dbfb8b9f |
| Genus | Dickeya | 91.89% | 6b5fb8a72a275ba8ca4433e61401da51 |
| Genus | Dickeya | 99.76% | 2af3ad172274a5518fa2aa9752f59208 |
| Genus | Dickeya | 99.76% | 0b6ce48800006c5a2ed4de1bef97809d |
| Genus | Dickeya | 99.94% | 32982489e1f4ff57a4f9c0db426e946f |
| Genus | Dickeya | 99.92% | 6e46dd59ddccb8f4627e7514d9c0c94d |
| Genus | Dickeya | 99.93% | 540ea6e7e293f7a98b70db095a6fd69e |
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| Genus | Dickeya | 99.93% | 548d86de11be7c91b82c09764c550afb |
| Genus | Dickeya | 92.87% | 5905eb4d97dc3ff63b965ae7d0032f78 |
| Genus | Dickeya | 91.81% | 14e22462351a6b4e05e07862bdaac947 |
| Genus | Dickeya | 99.94% | 845c9a08c4eae050c4794cf87a2cf36e |
| Genus | Dickeya | 96.84% | 52b57562beacab3bb982462a1dfcaa9c |
| Genus | Dickeya | 96.26% | b07240161241df091dd76c975a8a5ed8 |
| Genus | Dickeya | 99.37% | 25bb14137555f3dc23d279956d9439b0 |
| Genus | Dickeya | 99.80% | 90cac311949869433916d61d4d003739 |
| Genus | Dickeya | 99.76% | 0f613af3332c5de5f134b14cd98ccc7f |
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| Genus | Dickeya | 99.93% | b7ebd78530f2b7daf9a7c1c5bb2e7601 |
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| Genus | Dickeya | 99.99% | 17926028f316f4c77aab0fee503b0c63 |
| Genus | Dickeya | 91.59% | a55a52e092abe231d2a454d3f68b08e3 |
| Genus | Dickeya | 99.99% | 08a68ff746f1ee276ddd0bcb74957364 |
| Genus | Dickeya | 99.99% | 8d41f3efdaf29cdb350313cde9607646 |
| Genus | Dickeya | 97.38% | ea6df2ec8cf6ffc36f6893487c492de9 |
| Genus | Dickeya | 96.18% | d2036f2af04334c6cf94cb97ca7e8fdb |
| Genus | Dickeya | 91.21% | fb036c0d8e8ea8296b2e123858865f21 |
| Genus | Dickeya | 96.87% | b463c525f6a2e706a3a19565b077a1ee |
| Genus | Dickeya | 99.82% | 55331187a1fb7343d3f4495c13e45581 |
| Genus | Dickeya | 99.87% | 9e61f53d4c32ccb9020a1daee6665e0f |
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| Genus | Dickeya | 99.67% | 8ef50fa0ba0ee6d9118bc0561cd66637 |
| Genus | Dickeya | 87.57% | 82d5f6ad08b6938a30748284091ac35f |
| Genus | Dickeya | 99.94% | d339e115feaf2b837f4ba0a3e9a1e32a |
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| Genus | Dickeya | 96.33% | afc194e05db06a14af390cf8e8fbc3c6 |
| Genus | Dickeya | 99.83% | d19ac461827be06b714fa05e9385be2a |
| Genus | Dickeya | 99.95% | 37c252b36d89c9bd582af5c677135bee |
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| Genus | Dickeya | 99.99% | 5adf958f8fad900bbde5533d9de2cb63 |
| Genus | Dickeya | 89.38% | ae6c892a775a08eec4edf1f5d2999cd9 |
| Genus | Dickeya | 95.68% | 2967d96439f2edf0a6dc8388b5e15ff3 |
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| Genus | Dickeya | 99.80% | b913ae4679333f768c06fbab1ccb4fd8 |
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| Genus | Dickeya | 97.03% | a1cb7bf9d3750f7b24a8a0d85799e285 |
| Genus | Dickeya | 99.95% | 976ea920b2ad406f27592d5210c4c970 |
| Genus | Dickeya | 100.00% | 9b68f4273510302c1e9749bd6acbcb70 |
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| Genus | Dickeya | 99.86% | 5f80cc01dee3e5cc34ae23b91fe7bd70 |
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| Genus | Dickeya | 99.92% | c4d287b969fc960fd42fadc52d67f6cb |
| Genus | Dickeya | 89.38% | f4a04fd1916c08a640034e89df1e1711 |
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| Genus | Dickeya | 78.67% | 9e0fe3d51bc3385d4372b5bf3b403c2f |
| Genus | Dickeya | 99.87% | 24412eb4b5283e883a98069d276c31d1 |
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| Genus | Dickeya | 98.85% | e5f4669610c292d9667a3e1cd44dc35e |
| Genus | Dickeya | 80.83% | 8126cbd8e1b9d8d87c5f5067a3444ee9 |
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| Genus | Dickeya | 99.80% | 22de124d008c2d430a42c2d28c41d016 |
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| Genus | Dickeya | 99.93% | b7df8702ed1a005e695e2e23fc3fada3 |
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| Genus | Dickeya | 99.99% | f5a8e66e1131e75278f79081b8dee562 |
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| Genus | Dickeya | 99.59% | 8fb241fa7f0086816cd6df96d9df0a64 |
| Genus | Dickeya | 100.00% | d49dc6018fd0652f2b38abf50bffbf20 |
| Genus | Dickeya | 99.87% | 3895f5e785528717e1269abbfc55fe62 |
| Genus | Dickeya | 99.90% | 9c2d0e123489e8cba045913716cb31d3 |
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| Genus | Dickeya | 99.99% | 851838d3ad344b85118853a08e368012 |
| Genus | Dickeya | 100.00% | dd325e46dd5e323edbd7a255a1f8ad8b |
| Genus | Dickeya | 99.70% | ecf11db855a0b8a86bc4b76d664a625c |
| Genus | Dickeya | 99.75% | c09dbbde80fa953a95ff989d9a98ac6d |
| Genus | Dickeya | 99.86% | 54c9038e6fca86c7fe5b0c6e7b9086bc |
| Genus | Dickeya | 99.80% | 2b092a764f9b91a166641335a7f78425 |
| Genus | Dickeya | 99.80% | 67feec896ffe5ecc49d2201e8d49d3c9 |
| Genus | Dickeya | 99.83% | 56935a7841b116267cbe2b6eec30150a |
| Genus | Dickeya | 99.50% | 9ec4e00bdca2c03a59e75961c409582a |
| Genus | Dickeya | 100.00% | 468ccc88626371fafa2c3b55f2841b3c |
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| Genus | Dickeya | 99.93% | 5987edf79aac41d88b9fbd3dd4b06312 |
| Genus | Dickeya | 99.99% | 8671f8c6c129d335f8e90441c70e585b |
| Genus | Dickeya | 99.95% | 72bb9959bf4ab8f8268542850cbb4db4 |
| Genus | Dickeya | 99.97% | ea62f7865a4528c4da3346cf117a159a |
| Genus | Dickeya | 100.00% | f1d7d20b410d0c7faed44781cd4be0fb |
| Genus | Dickeya | 90.20% | eadf0f06fcd00584c561932237df18a2 |
| Genus | Dickeya | 99.94% | f21376ca252a44b98126370aa90e4a89 |
| Genus | Dickeya | 99.96% | 0057f898c9b2be6104e9bbdc7391d906 |
| Genus | Dickeya | 100.00% | d7e93323ec4b51845dfde2029bbfa652 |
| Genus | Dickeya | 99.92% | 1280efc3a833fedf1c825ee64f9a4352 |
| Genus | Dickeya | 99.97% | a1fac3af0e70908f0531ecae2ce2cd08 |
| Genus | Dickeya | 100.00% | dd922bce2ad1fcb9725c5910b6ac5d06 |
| Genus | Dickeya | 98.59% | 50ded981f6f6ee7e90d1d9698b7e7856 |
| Genus | Dickeya | 98.32% | 132ee9a62bfea39ed7682f2a0d76a03c |
| Genus | Dickeya | 99.98% | 84fe0d94a883eec69e02f510e2dabf66 |
| Genus | Dickeya | 99.97% | f2b48b1f50eb992c87337bf66b4e00ab |
| Genus | Dickeya | 99.91% | b8b775adb739a5202f25799a0ae11751 |
| Genus | Dickeya | 99.92% | 1fdb225ee43fe8d8e519f24ea5393bbb |
| Genus | Dickeya | 99.86% | 6fc73926144b8f14ece282c0b4c290e9 |
| Genus | Dickeya | 98.19% | e9116479bcc4c0b1704d2b78088281ba |
| Genus | Dickeya | 100.00% | 28e597bb4ae539e50425b2f677376677 |
| Genus | Dickeya | 71.02% | 46ca262d7383c9310827898ebe1262c0 |
| Genus | Dickeya | 98.90% | 59f3b41e35e3517e4a23433696cb4f9e |
| Genus | Dickeya | 97.91% | 3d16fd257fe0cc052b9adb51ac7ad7ab |
| Genus | Dickeya | 99.98% | 2b1871b0de13e643ffe4f928f7376d0d |
| Genus | Dickeya | 94.18% | 29fb866371329293fe542fca01ae4e2a |
| Genus | Dickeya | 99.99% | d69a22b95a96480981bf3de5125bc730 |
| Genus | Dickeya | 98.81% | b364cb5cdd6ca4515be42d5dad9d1133 |
| Genus | Dickeya | 85.16% | 822ece07f6fc633fe3b8e2f91d4f3002 |
| Genus | Dickeya | 99.96% | 881882cd7e3f8e991dd3c262ded68eda |
| Genus | Dickeya | 99.94% | c46262744d3758b696457b3f992513bd |
| Genus | Dickeya | 99.96% | 6acfebdaf3aa6eefe16b69b72b807ba5 |
| Genus | Dickeya | 99.97% | 28798fd6480a210d1f12752fa0ceaea9 |
| Genus | Dickeya | 95.95% | 2b71f1d78cc73aeb59b71a492ed65e0a |
| Genus | Dickeya | 99.77% | 9ba8630944964156e026a1c35a084f54 |
| Genus | Dickeya | 99.33% | 9f7da9a147d27e111f16aa3a3aa6f3f9 |
| Genus | Dickeya | 100.00% | fc20f2408d96948f2f8ceec4b4c3d8a8 |
| Genus | Dickeya | 77.40% | 387a298b37315afd1dc299ea249d9068 |
| Genus | Dickeya | 99.96% | d8c3d2c5b5a4b11e4556c5c1dcd8cb8b |
| Genus | Dickeya | 94.85% | 61d0dde81752d419b28d780b50bb5bc0 |
| Genus | Dickeya | 97.67% | 04905942d2cb6967fe2b639ac07575d6 |
| Genus | Dickeya | 97.90% | 9619665ed4739a655237b7679c9571e1 |
| Genus | Dickeya | 97.88% | e25327ca3926e2129b07119f0395bb1c |
| Genus | Dickeya | 97.93% | d7d3ee8c00cbb60f9e782d6544da1487 |
| Genus | Dickeya | 98.68% | 1870fde9255f0f07ac2e1f1841f05df4 |
| Genus | Dickeya | 99.93% | dce884fb6c68cfe37f49e7c19a33a14f |
| Genus | Dickeya | 90.59% | de2f051cbee340d98e8a6331a05adca2 |
| Genus | Dickeya | 97.97% | fe13e1f1dca6413c709c3c43d84a382d |
| Genus | Dickeya | 85.92% | b7f93079338737cdf3edc927f1a7da85 |
| Genus | Dickeya | 98.75% | f3184245604f437ed6acbccbc696ed6d |
| Genus | Dickeya | 99.89% | b61ec58eab6b019f8fa2f2c684ea5a24 |
| Genus | Dickeya | 99.89% | 39ac916eea2d4ba3bbdd6b9dfa513a05 |
| Genus | Duganella | 97.08% | f4e40369f5329aaf0f4657a71f5b72aa |
| Genus | Dysgonomonadaceae | 99.98% | f32bc1276481c00317ea446841c9c1c1 |
| Genus | Dysgonomonadaceae | 99.92% | a58a5676864ef541fd709afccd2da700 |
| Genus | Dysgonomonas | 72.78% | 7c3a11d94d8e884cc4f81e4d077d3a06 |
| Genus | Dysgonomonas | 96.54% | 073dafa7ac10668669aef84557e28408 |
| Genus | Dysgonomonas | 99.89% | a97854dbaf7e50381532f153178a02ff |
| Genus | Dysgonomonas | 97.56% | a0ed91ce73911275c3e563ec5e71c048 |
| Genus | Dysgonomonas | 71.56% | 17003bda04897c7708ece03da0bf3a72 |
| Genus | Dysgonomonas | 90.12% | 59a4dc39cb7943b574f37c9e01e784be |
| Genus | Enterococcus | 99.09% | 199837dfbc969128a5356c896cc188fd |
| Genus | Enterococcus | 96.62% | 9f94b5065f41b5d16571ecc4b6160083 |
| Genus | Enterococcus | 87.54% | 8a96ea4d4e901a03cc6d11dfdb022301 |
| Genus | Enterococcus | 96.73% | 938b4c15e7a286a3888af9af1eeb8e17 |
| Genus | Enterococcus | 96.66% | 0d0a33d21e09d83e4099b2c87b9f9def |
| Genus | Enterococcus | 96.51% | 65b91a923ca1c53f399bcfb3858e4e33 |
| Genus | Enterococcus | 96.46% | 3f01aeccbd92c1305593af043afeeead |
| Genus | Enterococcus | 99.08% | b6e8242a62bad762c1f00e78fcb21b85 |
| Genus | Enterococcus | 97.03% | d2b5cab3d5f3f69cfc9b0efb05e9e99d |
| Genus | Enterococcus | 99.14% | 6851d0df97039d1a69d6652e744e4dd6 |
| Genus | Enterococcus | 99.15% | 730e79b87611add37f746ae6b10d8887 |
| Genus | Enterococcus | 96.62% | a42f225d259885ec2a655fcd2c76517d |
| Genus | Enterococcus | 98.12% | a41826073f966772c3da62c4fef6bb05 |
| Genus | Enterococcus | 99.18% | 441d02bad57b8f8ce75c531c4f1076d3 |
| Genus | Enterococcus | 86.69% | a623d46842a0fb8daa7a450325041c70 |
| Genus | Enterococcus | 96.94% | 1186974e6f12400931d731722dab6396 |
| Genus | Enterococcus | 96.70% | bf4e822b5cbad1528c56753f9028cba2 |
| Genus | Erysipelothrix | 79.13% | 39c689425ce1fa80244d9f1a917b38a0 |
| Genus | Erysipelothrix | 75.77% | c484d59e3bd2f49133322b3bd6e93230 |
| Genus | Erysipelothrix | 75.69% | 109757324af150ae73272f83e3f92fec |
| Genus | Erysipelothrix | 72.83% | df5aa14309c52af70ce772972f2dea25 |
| Genus | Fusobacterium | 100.00% | d8d18fcd2755ef7ee6732a4cf80518ab |
| Genus | Fusobacterium | 100.00% | 18bd3f294f8e05c6624bf70f32a8b2fa |
| Genus | Fusobacterium | 99.81% | 01e5ce19471349234481709ade192476 |
| Genus | Fusobacterium | 99.73% | ca4a24cfc21810cd7fd154b57e121232 |
| Genus | Fusobacterium | 100.00% | d1b6e59a665f83abce1aae8639ac471f |
| Genus | Fusobacterium | 99.84% | 44c3c4bb3e7506f83737ff8242a8f6ff |
| Genus | Fusobacterium | 100.00% | 9fc10607c800d7d0e2e76a79095355c7 |
| Genus | Fusobacterium | 100.00% | a618338c08d51dbd4d36529b5826b105 |
| Genus | Fusobacterium | 99.98% | bf9f408f5e596843b03f7db2e9d0d6ac |
| Genus | Fusobacterium | 77.75% | b0ac52a1af7c7e6791eb728682803640 |
| Genus | Halomonas | 80.14% | f2885a4af47dbf047bddeba62cefcaa4 |
| Genus | Halomonas | 80.05% | 2aec735001d67636017d54cce4560b99 |
| Genus | Halomonas | 80.14% | e4dd77e2dc6224b1181b2a436de0a1d0 |
| Genus | Halomonas | 81.28% | 4b4c087cb5c3d3f59c149d7bb833332b |
| Genus | Halomonas | 85.72% | 4b1144d1d3227bfbf5d5f18c8a8e3360 |
| Genus | Halomonas | 85.73% | 15c711bf391c646d21821134b06f34c9 |
| Genus | Halomonas | 80.01% | fe0dd2ac7168a45141b715e9aa5637a8 |
| Genus | Halomonas | 80.03% | b9026efe7ee754994ed400770bd093dc |
| Genus | Halomonas | 95.49% | 69e9026a1387c26e29b7a951aa662111 |
| Genus | Halomonas | 76.23% | 92395a6ba4b4df5120bc6af34fa55db2 |
| Genus | Halomonas | 85.72% | 60f07087e973378fd332a4fbe67ca188 |
| Genus | Halomonas | 80.08% | 4181df311dfbef48b5356d92e4f8c429 |
| Genus | Halomonas | 85.71% | 5401c677ebb34ee57eeed2ad200b7ef9 |
| Genus | Halomonas | 82.49% | 325de694f4ead8e5f2e5fc7383009087 |
| Genus | Halomonas | 76.14% | 176a0861d0ff72280001be4f5ec0848e |
| Genus | Hungatella | 74.75% | 6b1a990ebb6c54c8988ea4402a17e7bd |
| Genus | Hymenobacter | 100.00% | aa3fce77ee5be0d8db418accc84e6f66 |
| Genus | Hymenobacter | 100.00% | a48a1788a348a68b6b357c734fc2e39b |
| Genus | Hymenobacter | 100.00% | 603054bc1ce41ae95c8c7d72cb83e983 |
| Genus | Hymenobacter | 100.00% | aff517d66380fd46745322f1c28da51f |
| Genus | Hymenobacter | 99.63% | cfd235b0294cda484c7facc3332baab4 |
| Genus | Ignatzschineria | 100.00% | 6eeb19ad217e44a1d9f1c3672d8574a3 |
| Genus | Ignatzschineria | 100.00% | 3dce234105be5d54cc7debf7a042b8b7 |
| Genus | Lachnoclostridium | 88.96% | ac26d8d57d7654c96f74205dc420af0f |
| Genus | Lachnoclostridium | 98.74% | 4ab33b955c716644aabf94626958578e |
| Genus | Larkinella | 99.99% | 1e1818e5a691045150ce2891094343c1 |
| Genus | Leifsonia | 75.95% | 20f07cce575680b11d6eab72bec556ab |
| Genus | Leifsonia | 76.76% | 18138f98122247583038ce20bca21bdb |
| Genus | Leifsonia | 76.02% | b0272b459b5cca183f773cd664a797aa |
| Genus | Leifsonia | 75.99% | 33c2a753fdebe4b775d8b6a559c9096d |
| Genus | Massilia | 99.93% | 5d40daebcf82458d96910b832ed36d72 |
| Genus | Massilia | 99.94% | 388fe08cf2d177a3c9461415654e6e13 |
| Genus | Massilia | 99.52% | ee8a6db5fa4bcd4ebd4bc80fa4d36f5e |
| Genus | Massilia | 99.93% | 00d427ac8a16adac492f4dc9b9fbe3f5 |
| Genus | Massilia | 99.93% | 37842e73bd8807bfb35697716ed24d1a |
| Genus | Massilia | 99.94% | 43d97df6025dab34da5e0008d05d2124 |
| Genus | Massilia | 93.92% | 2216387acc4003ac0e3a822ed1dd3161 |
| Genus | Massilia | 99.73% | aadd73ad2159adc5c994e4a24e063c51 |
| Genus | Massilia | 99.93% | 611c8255a915554c488ea8ae85d3c189 |
| Genus | Massilia | 99.93% | f330c64120a6eeacf01f84b15f011793 |
| Genus | Methylobacterium-Methylorubrum | 99.99% | 14a7bf8c7f97b442f5e1d17c13008cd5 |
| Genus | Methylobacterium-Methylorubrum | 99.99% | 9591e5d7df6bf4bcfd339190d4b96b9f |
| Genus | Methylobacterium-Methylorubrum | 99.98% | 137f90af9f988d4c445bbb486c233455 |
| Genus | Methylobacterium-Methylorubrum | 99.99% | 7a54c8e445b842218a489b7d5e309d66 |
| Genus | Methylobacterium-Methylorubrum | 99.72% | fe07e7e889d8550f185e555d053f4e89 |
| Genus | Methylobacterium-Methylorubrum | 99.99% | b6b0d5873ba8e72d2da89bd6e1b4fbe3 |
| Genus | Methylobacterium-Methylorubrum | 89.09% | 06e1d2b37c8d7d93e56b0c03609ba282 |
| Genus | Microbacterium | 90.76% | d0aa71b351fe588bb6afd318d7b55929 |
| Genus | Muribaculaceae | 98.45% | 03e56adf77dbb807dd7a3233f14d67f8 |
| Genus | Mycoplasma | 100.00% | 0ab15634508b35ab615a9a783d73a0ff |
| Genus | Mycoplasma | 100.00% | 52aba174e3196ef122a1634627f6fccf |
| Genus | Mycoplasma | 100.00% | f99c3247705c1575e5d82aa650d69e71 |
| Genus | Mycoplasma | 100.00% | e072d37ef26b63b4c42ae94fda6f0e75 |
| Genus | Mycoplasma | 84.47% | 8658bb42034e86f201706654854b2e7e |
| Genus | Mycoplasma | 100.00% | 32a9c4656b7347e020ebcbbd7f4eb7ea |
| Genus | Mycoplasma | 97.82% | 384e51fb942135e27f3619684602e583 |
| Genus | Mycoplasma | 73.50% | c599dbe0a9ec5a6b518d9783bcd8ae3d |
| Genus | Mycoplasma | 98.63% | 7ac14211b9ef9a04443cd102424def3a |
| Genus | Mycoplasma | 100.00% | b2875945263efeb4e5d718fd7f9d1891 |
| Genus | Mycoplasma | 100.00% | de911a9f2476987336ea1cf57291cafb |
| Genus | Mycoplasma | 100.00% | a7ee3c34ab63af470019a6d074b3b469 |
| Genus | Mycoplasma | 100.00% | d79ee5904d6e739fdd0f690a2b6a74d6 |
| Genus | Mycoplasma | 96.28% | 192e6a05b4a5bdbe5beff1981ac4f4cd |
| Genus | Mycoplasma | 100.00% | b544d241645b3389f04cc34d11ceb87c |
| Genus | Mycoplasma | 100.00% | 74237c2d49233df86a17287b3f374634 |
| Genus | Myroides | 96.82% | c25bc05550bb917dd748db44c709d000 |
| Genus | Nocardioides | 82.93% | dfbae5e8db097cbff8a74f13bc4d1d1d |
| Genus | Nocardioides | 74.69% | c891127854d195e317617c0895ea9099 |
| Genus | Nocardioides | 97.39% | 143f7dc8f7dcff4a96f4215b970cd515 |
| Genus | Novosphingobium | 96.35% | 925945142c3b81fd5acbeaf25c2e4e2d |
| Genus | Ochrobactrum | 74.42% | 85230871537c0ac0f3830a6fd8f0ac4e |
| Genus | Orbus | 73.16% | 455a29fccff0feb1028123d91049bb2b |
| Genus | Parabacteroides | 79.89% | fdf6dcca91d76fd126fbf2596f83e44b |
| Genus | Parabacteroides | 100.00% | 18079a1b9bf9912bb00df2dd0278ca4e |
| Genus | Parabacteroides | 98.77% | 0e2d9449dbac054e26be5d0a90f4bb44 |
| Genus | Parabacteroides | 76.79% | 0308a8b7d5e490b32b830494f8d1aa75 |
| Genus | Parabacteroides | 74.94% | 9c1eab4bfef33ed26dda2af48eb4936e |
| Genus | Parabacteroides | 99.56% | 9d61059f98f80b6a7846a8f91c6799a3 |
| Genus | Parabacteroides | 99.97% | 5298714075c9f104bc2fb4a206d1b4c0 |
| Genus | Parabacteroides | 99.99% | aed2b1076a0b54a82e6d99c4a934120f |
| Genus | Paracoccus | 76.06% | fe4a71a0eb5d6313020de2ce9a40b6e0 |
| Genus | Pedobacter | 97.35% | f69df7e15b2b39cca43c1827e690cb6e |
| Genus | Pedobacter | 97.36% | e94874a901c66cc2c6fd462fa5917e43 |
| Genus | Peptoniphilus | 99.00% | 7b9e74c11b16b47bbe61014bd88e2e5f |
| Genus | Phascolarctobacterium | 71.00% | afd363d43bc712be45fef64eda13c90a |
| Genus | Phascolarctobacterium | 71.09% | 47854c3d41ff687b172d936b506b2cce |
| Genus | Proteiniphilum | 98.48% | b1e750d53d3885024286a761dd7dba5f |
| Genus | Pseudomonas | 91.97% | cc756225c8492a3d79c5e89e5fff2471 |
| Genus | Pseudomonas | 95.93% | e2ff5b16430ae09e7dc0d5153136b10a |
| Genus | Pseudomonas | 94.16% | b680cd497192af0ea33d2e8fc3819826 |
| Genus | Pseudomonas | 94.21% | 85cb3c2a3f213b6ceb0ee8de01c5e0cd |
| Genus | Pseudomonas | 88.75% | 078c918e119457a547826ab073a1d26c |
| Genus | Pseudomonas | 83.44% | 2cbb81553aeb8e5a48f01e66d0c9c525 |
| Genus | Pseudomonas | 99.74% | 1372622ea06ef763ddc0d5e8fd3d3baa |
| Genus | Pseudomonas | 96.98% | defd221ecacc2d712fe430cf36dcd11b |
| Genus | Pseudomonas | 99.33% | 44c60a05e738384017a1ca2e794193c2 |
| Genus | Pseudomonas | 83.14% | 021526c443deaa5473532f913c0d43a2 |
| Genus | Pseudomonas | 97.64% | c9cfbf65b7fa97ed7793a74113e4ffe8 |
| Genus | Pseudomonas | 99.28% | 3969b9981cf74bee3c2c25c29d018b97 |
| Genus | Pseudomonas | 86.44% | 81c67751d12c75b1abdd7c1cd9d4e703 |
| Genus | Pseudomonas | 99.38% | dc7843b15857015c23e9d4c9692ae099 |
| Genus | Pseudomonas | 84.26% | 26bb12e1123a26f057dffed705b71c10 |
| Genus | Pseudomonas | 90.37% | ce9021f7e18d6be23e490ecd2baf5010 |
| Genus | Pseudomonas | 99.15% | 245d9882b797d1135e5411999435fb9b |
| Genus | Psychrobacter | 78.14% | e9bd5988ee46e7ec3086b4f289711a47 |
| Genus | Quadrisphaera | 70.02% | 882a7b37ed41c9ffee77d0cb77188e45 |
| Genus | Quadrisphaera | 74.26% | 84d7faf5a45d65a188ae22b2543858b5 |
| Genus | Raoultibacter | 100.00% | dfe423b354081242abab139e680e11ce |
| Genus | Rickettsia | 95.66% | db569ba071fa72bbbf5144aeed358e31 |
| Genus | Rickettsiella | 99.40% | 7597608464ebcbbae223e774cee2e5cb |
| Genus | Rikenellaceae RC9 gut group | 71.57% | d8c3753d0570fde45b2da69703144200 |
| Genus | Rikenellaceae RC9 gut group | 70.64% | abd575ced35313bf5719f46b9a20ff88 |
| Genus | Rs-K70 termite group | 98.21% | 0a28577d488c120007d7420f41c635bf |
| Genus | Segetibacter | 75.42% | 24f36b254921d9ed2eb835fa5e6b7956 |
| Genus | Sphingobium | 97.23% | e92a8a6c81cfad6d8b172d83b4f15fe7 |
| Genus | Sphingomonas | 99.14% | 6581f2d5e43d72db1159f34180e4e38f |
| Genus | Sphingomonas | 98.92% | 722c5a3e58aef125b5228f82c93a4a78 |
| Genus | Sphingomonas | 97.68% | ef7a610e2fd06583f8e06eea9c838656 |
| Genus | Sphingomonas | 98.80% | 5f62f98b0f63a30d74ca16e10d2ac747 |
| Genus | Sphingomonas | 99.11% | a2aecc9bebf3eeb50ed24c76abc14ebb |
| Genus | Sphingomonas | 99.00% | 33b857811e238cfb419e32319cde15e3 |
| Genus | Sphingomonas | 99.37% | b680ed3b7845117d84d5a48baca96467 |
| Genus | Sphingomonas | 94.64% | 846e6e659197b4b61f9384eee84b03a4 |
| Genus | Sphingomonas | 97.45% | df6f4ec33db8c91759b275e4616cc58b |
| Genus | Sphingomonas | 99.32% | 046c19a3f3b77ec3ca4fcff2ff1f0122 |
| Genus | Sphingomonas | 99.67% | 1e925e3cb374d5f3f713921546d9a9b8 |
| Genus | Sphingomonas | 95.78% | f92ccc633c2ae9e143bc8ab5fd24f925 |
| Genus | Sphingomonas | 96.95% | 78172c3cfd995f25dc7973c7ce6a18c0 |
| Genus | Sphingomonas | 94.13% | d9ea9edfef835f4a423c3ac8372916a1 |
| Genus | Sphingomonas | 97.99% | 98b5d49d5280ce2d4170c8eca9957730 |
| Genus | Sphingomonas | 98.64% | b3937d174b147a918dab2e0c7110e132 |
| Genus | Sphingomonas | 98.98% | a577b1bd0da6db79dc9c6f884f0c21ad |
| Genus | Sphingomonas | 98.79% | 0ee12308b78a6055cfad57ca4a9387d5 |
| Genus | Sphingomonas | 98.99% | 841f9c9fb165504c5901476c99f1b1f2 |
| Genus | Stenotrophomonas | 99.75% | 995ae13c4a44ad1587e90e6a98a43b4d |
| Genus | Stenotrophomonas | 79.40% | 3fa1963f5948e0adf6c63b598327aca6 |
| Genus | Stenotrophomonas | 92.96% | 758ba8f879ed18aa6b0482dda3edd1d8 |
| Genus | Streptococcus | 73.40% | df4071ef4063d5281fb77a39eb5adade |
| Genus | Subdoligranulum | 71.50% | 25fb46275aa72ac83b2224977db58844 |
| Genus | Subdoligranulum | 79.77% | 15de4605965aa110667be31f12c46807 |
| Genus | Sutterella | 81.09% | 53957d34e541d94cfad66cd808072d1f |
| Genus | Sutterella | 92.33% | 0b02ec4c4a84ae3f397c67549aa9fef5 |
| Genus | Tepidibacter | 90.83% | 8e1f5234121901a30fb2b60178f1c5db |
| Genus | Tepidibacter | 99.31% | 6b19cb93f276b4ea1dfa5abe4499c6b4 |
| Genus | Tepidibacter | 92.08% | 759af98cbffc00513d478d0ba626f0e1 |
| Genus | Tepidibacter | 99.89% | a6ee2b7b11fa2b20b0dff02a865ea3ac |
| Genus | Tepidibacter | 99.61% | 651a8fcce302d9cf706a3d2bf51ee056 |
| Genus | Terrimicrobium | 76.05% | 2c37a9aed679101f7e9e4e7ba2ce41af |
| Genus | Terrimicrobium | 85.32% | 7cde17aeb3ec849e7998860a83007103 |
| Genus | Tissierella | 98.89% | 2abb797cf686cd283db3dbb158cf38f3 |
| Genus | Tissierella | 98.22% | 316b30b1c638c43cb64a2f9ffcee838a |
| Genus | Tissierella | 98.56% | bde6f4615633a3fba0a9aaa34c89cd8a |
| Genus | Ureaplasma | 99.60% | e399bdf2b31b8c4329f5f0de91cac663 |
| Genus | Vagococcus | 77.40% | e2c691d384947d5a9a5d252a8365f815 |
| Genus | Vagococcus | 96.05% | 50a3aa3d6ed5d8c20a9b93ccd65776c9 |
| Genus | Vagococcus | 96.04% | dbcb3b62741eaf5ea1976fa1004ff28a |
| Genus | Vagococcus | 81.52% | 88b03d8a564be644c04b662c08b6c73c |
| Genus | Vagococcus | 78.09% | 3c709ff2361b972cfc3e7eabcd8505d8 |
| Genus | Vagococcus | 81.52% | 251809f182ebccec382ec903f77f80ed |
| Genus | Vibrio | 74.75% | 06a55bd9e16cc555cb6bc6593f005888 |
| Genus | Weissella | 100.00% | 0679e095cfba30268ade72503333b97e |
| Genus | Zymobacter | 91.58% | 6083285ba95147a7dfd1d6874215aca9 |
| Genus | Zymobacter | 91.49% | 817b083b30b59a2bae9a2210c6b47479 |
| Genus | Zymobacter | 91.61% | fd4eef00430f9c23e329d0369f79acbe |
| Species | *Clostridium colinum* | 90.82% | 4a1022127250417f1dc068e4fa0706c7 |
| Species | *Pseudomonas geniculata* | 100.00% | a6c7d7013c0b7746d7dc0aee6d790d75 |
| Species | *Pseudomonas geniculata* | 84.27% | d373494891190e2f0a15b59577b85a74 |
| Species | *Pseudomonas geniculata* | 95.39% | 500a304512fc42d696945d732402e5a4 |
| Species | *Pseudomonas geniculata* | 78.81% | d08a2fab1f257917d7607a2ac9a48571 |
| Species | *Pseudomonas geniculata* | 96.20% | 9d7613fd6079d49a36370760d770d073 |
| Species | *Pseudomonas geniculata* | 98.26% | 225f9057832ffc48a7edfd9bc56ad79c |
| Species | *Pseudomonas geniculata* | 99.99% | 352533bd5d8174a34be0de814f2aa048 |
| Species | *Pseudomonas geniculata* | 100.00% | 15be7db67bce9df6ccc3a80c6b159d50 |
| Species | *Pseudomonas geniculata* | 99.98% | 3a8e0dd0cb90030dd187932141bdce73 |
| Species | *Pseudomonas geniculata* | 97.03% | 1d3a5d213690a05a8efd246fa00bfa1c |
| Species | *Acidibacter ferrireducens* | 84.48% | 1993c26531463e52fb6e0a886aa7535e |
| Species | *Acidobacteria bacterium* | 71.00% | b5cbd05e82621a94a33ad29663c21720 |
| Species | *Acinetobacter indicus* | 75.25% | 673684063906ad7b66687748bfddca9b |
| Species | *Acinetobacter towneri* | 71.53% | f9d57a301e88538d48cecc989504809e |
| Species | *Acinetobacter towneri* | 70.50% | 7c1ff39ca09125c431127cf90ddfa748 |
| Species | *Aerococcus sp.* | 99.88% | 2a43bb21878f18d4f9d48af175cebd6a |
| Species | *Akkermansia sp.* | 96.94% | bd04d37aa807e55e40d50413e68dbc0b |
| Species | *Alistipes finegoldii* | 97.80% | b1f10bc3618c8846527530eed578a2d8 |
| Species | *Alistipes finegoldii* | 99.82% | ac441333d6657e0ca0cabe718fed2bce |
| Species | *Alistipes finegoldii* | 99.82% | 816d84070ba4669c88e67397e13459c8 |
| Species | *Alistipes finegoldii* | 96.76% | f921b410630a1bbb698f00336987cd09 |
| Species | *Alistipes finegoldii* | 85.22% | db2e67d19139ff81fdead54cb267f67a |
| Species | *Alistipes finegoldii* | 95.45% | 71e84cfa2c715370ecd9159c4bde1ee5 |
| Species | *Alistipes finegoldii* | 91.59% | f092d34fa67b2a1d54af887e43185aa9 |
| Species | *Alistipes finegoldii* | 99.85% | 186fdd7365c0441cfb31e2c563fb143c |
| Species | *Alistipes finegoldii* | 81.42% | 0f8caf0dbe6e7c3a7fa68f03e72eb6c7 |
| Species | *Alosa alosa* | 70.80% | 0ef2dda873865841cb6f66d235e8719d |
| Species | *Anaerococcus provencensis* | 98.09% | f785f17975aab570186cc46a4383d96a |
| Species | *Apibacter sp.* | 96.81% | d33824ffc0dcf9798c9685867ae1e16a |
| Species | *Arachis hypogaea* | 82.40% | 2383d49241941eadb2df119d26c95c59 |
| Species | *Asterionellopsis glacialis* | 85.45% | 188ca78d0aa1b87be6827209d89559fb |
| Species | *Bacteroides acidifaciens* | 81.59% | ab6fbea079385cca432d72958cb1d145 |
| Species | *Bacteroides acidifaciens* | 81.57% | c9a7f72e762cf0d5d66b1e104c176370 |
| Species | *Bacteroides acidifaciens* | 81.57% | 793d6c799f748781300362ecdeb3c0fc |
| Species | *Bacteroides acidifaciens* | 81.75% | b30c4194a9076646bfb8b0426199f3d6 |
| Species | *Bacteroides acidifaciens* | 81.62% | 198922977e8f5bff56e6a4519ebefb3b |
| Species | *Bacteroides acidifaciens* | 85.01% | a62beb8c113721b0362097c476ddf049 |
| Species | *Bacteroides acidifaciens* | 80.40% | c37054e2a145e9c1c86c2c56f44bc535 |
| Species | *Bacteroides acidifaciens* | 84.93% | f81aced774d43b235403a18ad69c0a1b |
| Species | *Bacteroides acidifaciens* | 80.23% | 71cfff45411941c0d590d55d5f117b37 |
| Species | *Bacteroides acidifaciens* | 87.12% | f9a59ee5bc92e02c6a4ca912d83dee10 |
| Species | *Bacteroides caccae* | 92.64% | d850fbe2521f7556b7986f166eade5b0 |
| Species | *Bacteroides caccae* | 92.51% | 1be415d932c000799145d96a417efe74 |
| Species | *Bacteroides cellulosilyticus* | 71.10% | 4e53c1fff423dccdfed9a19ce62cb2aa |
| Species | *Bacteroides cellulosilyticus* | 73.96% | 2fa5e71e021d230e5c931e61cae1df3a |
| Species | *Bacteroides cellulosilyticus* | 90.74% | 6a07dfe6101042ff335160ff345240e0 |
| Species | *Bacteroides coprosuis* | 100.00% | 5c1e210a30d163215d5b46c8c4856d57 |
| Species | *Bacteroides coprosuis* | 100.00% | cd30eaf05513011519af593732a113dc |
| Species | *Bacteroides coprosuis* | 100.00% | 66094e0f7adb205367785086d82455da |
| Species | *Bacteroides coprosuis* | 91.43% | f1df06b97039304bfbe6b60db7476ed2 |
| Species | *Bacteroides coprosuis* | 98.54% | e785e49cc1c2d07fe4a5b0899c810652 |
| Species | *Bacteroides coprosuis* | 100.00% | 0a2cfd7fbdad05a700722e169effa2ae |
| Species | *Bacteroides coprosuis* | 100.00% | 319010b5fd47897dbbfbb04bcf6c0431 |
| Species | *Bacteroides massiliensis* | 99.96% | b389ad0295416a01c53e03783cb7fb7a |
| Species | *Bacteroides oleiciplenus* | 78.23% | 76d423038ac860b92f703372d3c0eb7e |
| Species | *Bacteroides propionicifaciens* | 93.36% | fe0964272d3b6b751ebf7fddb6d65121 |
| Species | *Bacteroides sp.* | 70.89% | 1a7fe9f20f242128320c67fff5ca3e3f |
| Species | *Bacteroides sp.* | 70.75% | 665a5063e21bdaf3ee449c6b5749ff28 |
| Species | *Bacteroides sp.* | 70.16% | 84515c0744a56d264e09b877484ebbaa |
| Species | *Bacteroides sp.* | 73.91% | c78d2e53b6696f90a8de6ca7b818a9c6 |
| Species | *Bacteroides sp.* | 72.29% | f73fee2936db6fb5ec5ef8a4783634c2 |
| Species | *Bacteroides sp.* | 77.33% | 372bc20413cfbbd7395f37a33c831229 |
| Species | *Bacteroides sp.* | 74.44% | f849b0b1e3e22e1676eba6fc2cbb884d |
| Species | *Bacteroides sp.* | 74.00% | 2ba44c718850998dc9b7a412644956ea |
| Species | *Bacteroides sp.* | 73.31% | d11df811cd73ea3afff2d4bbf45bad06 |
| Species | *Bacteroides stercoris* | 97.28% | 964ae73dd60ab555978837afbb0ec189 |
| Species | *Bacteroides thetaiotaomicron* | 86.77% | cc7ab79329a59043e3aed5db8202bdb1 |
| Species | *Bacteroides thetaiotaomicron* | 88.67% | 0821e7745e37471884595d5ac44f6a90 |
| Species | *Bacteroides thetaiotaomicron* | 88.63% | 30188e081e77769a3484aeded92bfc47 |
| Species | *Bacteroides thetaiotaomicron* | 88.63% | 268c68ccbb2c00037682711f21520534 |
| Species | *Bacteroides thetaiotaomicron* | 84.93% | 6907f1d90712abb06b18fcbb7b81497e |
| Species | *Bacteroides uniformis* | 97.90% | 3de114d89f5ef3586cb45c0613b6f782 |
| Species | *Bacteroides uniformis* | 99.49% | 7a270269524e86429f7f0cdb2a8af19c |
| Species | *Bacteroides uniformis* | 97.90% | 0bfc73e3c5110eb7e2536c8bbffdf4b4 |
| Species | *Bacteroides uniformis* | 99.48% | 7231d6d332b3dcf6547bf28bcb6bb089 |
| Species | *Bacteroides vulgatus* | 80.83% | 59361bb35a047b3d011b44e31235aaea |
| Species | *Bacteroides vulgatus* | 80.97% | 916005967616bfb93e9d94728c7d4d67 |
| Species | *Bacteroides vulgatus* | 73.13% | 898176c93314421708734269cfbae087 |
| Species | *Bacteroides vulgatus* | 77.75% | f5d642a427b449de21a21302802704d6 |
| Species | *Bacteroides vulgatus* | 70.79% | eb3f00ce00b491e7608c719f68204a8b |
| Species | *Bactoderma rosea* | 99.66% | 9369c8553e1cfe42d1839147e6551704 |
| Species | *Bergeyella sp.* | 78.78% | 08397ad00dba5d204c75b94aed9fa0fd |
| Species | *Bilophila wadsworthia* | 92.46% | fab196834ee96c75cfdd5893624e6454 |
| Species | *Brachyspira sp.* | 96.45% | 56341489588cb4ed0175406821b3a0ab |
| Species | *Brevibacterium permense* | 99.46% | 7806288847c366b64fe93472480afb41 |
| Species | *Brevundimonas naejangsanensis* | 81.07% | 4208f2da56bf2625b0937679de9358fb |
| Species | *Brevundimonas naejangsanensis* | 78.67% | 41f8b15a2cf5cbe2f79ef3a767962c9a |
| Species | *Brevundimonas naejangsanensis* | 79.34% | c8fb4526335b41514efeaa5772121ac7 |
| Species | *Breznakia pachnodae* | 83.99% | 1cf0ad1783ee2048a62bc59c8b985e57 |
| Species | *Burkholderia sp.* | 70.90% | 8d0c7a1ae73b693e333a84eb7d3df16b |
| Species | *Candidatus Fritschea* | 99.67% | 318926299badb10196e7a251afd63757 |
| Species | *Candidatus Rhabdochlamydia* | 71.63% | 49a9a52b925692e7aa40234c6db7cd86 |
| Species | *Cardiobacterium valvarum* | 93.88% | d0d8f356bff5768ac3b56ca8242f09f9 |
| Species | *Cardiobacterium valvarum* | 92.18% | ab23de616146157c1b93004d2567c05b |
| Species | *Cardiobacterium valvarum* | 88.40% | daa4096c2f378516f34a3aed8ef01075 |
| Species | *Cardiobacterium valvarum* | 86.68% | 7dee2db8c745def63ff6317ee9a9d1d0 |
| Species | *Cardiobacterium valvarum* | 97.73% | 0fa0c26e2c481fffa464b7500977761f |
| Species | *Cardiobacterium valvarum* | 96.55% | 92567f85302ca5eb29432343f002f997 |
| Species | *Cardiobacterium valvarum* | 98.53% | 807486d7d1ee0bf0f47c2e6d8595e06d |
| Species | *Cardiobacterium valvarum* | 97.26% | dacdc210831675f408612fd26a672bd6 |
| Species | *Carnimonas nigrificans* | 76.48% | d9fdd6f7cbb93dfd9936ba02e624023e |
| Species | *Cenchrus americanus* | 96.81% | b60252eb76e3ba7e39df03bc3744da9f |
| Species | *Cenchrus americanus* | 86.75% | 6818764be5110bb8859d4f719ee1e2fe |
| Species | *Chryseobacterium hispalense* | 76.41% | 45436d72e4cc68e25522cfffd20ea278 |
| Species | *Chryseobacterium hispalense* | 79.63% | 07f4df513dd4b55ca037b0d20b2dfae3 |
| Species | *Chryseobacterium hispalense* | 79.41% | b140eb1ec2b873b2449bcb92022c70be |
| Species | *Chryseobacterium piscium* | 93.58% | a6290c26e6a6f044578fbebfc9fdf17c |
| Species | *Clostridioides difficile* | 99.98% | 17309c08d5300e4cb88a5e2b74b8a302 |
| Species | *Colobanthus quitensis* | 99.95% | 85c2213172a0bb333c4072a4e6640578 |
| Species | *Colobanthus quitensis* | 99.94% | e5a1ff68b192557b713757b2bdbd89e2 |
| Species | *Coptotermes gestroi* | 82.61% | 7e31d2fb30ca079608fd95762ababf65 |
| Species | *Coptotermes gestroi* | 99.70% | 8514064bce5f4a5429ca1dcb3af696f6 |
| Species | *Coptotermes gestroi* | 99.71% | b75dad7538fc3ed45baeed07344ac314 |
| Species | *Corynebacterium callunae* | 80.89% | dc7af4373d9eb88abf0febcb2075ea86 |
| Species | *Corynebacterium tuberculostearicum* | 89.14% | 1d0f1b6267509f545f4ef98cb541e744 |
| Species | *Daucus carota* | 99.67% | 515bb92406e66eac10c9293fef495ca8 |
| Species | *Daucus carota* | 99.94% | 5b7d35eaf6b871fcc5732dd80d559238 |
| Species | *Daucus carota* | 99.67% | b18ccc34b36511d065fca9053b306dee |
| Species | *Daucus carota* | 99.65% | 307a9f756a72ef19df4920e74b130417 |
| Species | *Daucus carota* | 99.67% | 0e11e8ee3090848f13d95442eeba812b |
| Species | *Daucus carota* | 99.99% | 5a2d242a5599dd5bfb2d57825e0b0fe6 |
| Species | *Daucus carota* | 99.90% | 5a31106ecc8656748927098449c0a74f |
| Species | *Daucus carota* | 99.99% | c128ef79d5fd78af2f3bcc841bda3946 |
| Species | *Daucus carota* | 99.99% | 6d90c25a35824cb9bf8547cdb9bf411d |
| Species | *Daucus carota* | 99.99% | 9797cca18013069ad09ad0d52b51bf24 |
| Species | *Deinococcus radiomollis* | 79.36% | 34f93cb0ad3c09e4bdcdf7e632e53dec |
| Species | *Dolosigranulum pigrum* | 92.91% | 79ec83414a5dd668295b09e332b12843 |
| Species | *Dysgonomonas alginatilytica* | 88.51% | a1f77493f91d561e132ff4ae4d73f87d |
| Species | *Dysgonomonas alginatilytica* | 99.95% | 15c22030c37b8bc5a45d844f45c015aa |
| Species | *Dysgonomonas macrotermitis* | 80.85% | 01bd02069cedce9d19940e1ac880938b |
| Species | *Dysgonomonas macrotermitis* | 81.95% | e5c77bf7fb94cc24b85bac93d6b2f5ca |
| Species | *Dysgonomonas mossii* | 71.54% | 7f183e518f7b1082334610c045c2e7d8 |
| Species | *Dysgonomonas mossii* | 81.57% | 91aa23d445d5689f924e527a7429dc0b |
| Species | *Dysgonomonas mossii* | 83.34% | 201f61c6e20c96515613863531f97380 |
| Species | *Dysgonomonas mossii* | 99.76% | 6a10ddd08426a6560e8f7d06c3bb1e8a |
| Species | *Dysgonomonas mossii* | 92.73% | b47aa613250a0a96f95c1a6fc71fe84e |
| Species | *Edaphochlorella mirabilis* | 73.70% | 8c0a4e6509f38df362bbb6eeec050022 |
| Species | *Edaphochlorella mirabilis* | 73.71% | 148b20b04def44b97bbb3185425a8751 |
| Species | *Edaphochlorella mirabilis* | 73.02% | 66ed1044c1e0da663587381934a9306e |
| Species | *Enterococcus flavescens* | 71.61% | de658bed4810d2bf6fd8e584ba25a775 |
| Species | *Enterococcus flavescens* | 96.24% | a2e46f566ff73361079ba719d5954051 |
| Species | *Enterococcus flavescens* | 96.24% | c35084b2ef74f14e59b44146b8402123 |
| Species | *Enterococcus flavescens* | 71.22% | 3157b3131e434adefcd6753acacae5ac |
| Species | *Enterococcus flavescens* | 70.97% | af35fef875a0953eac28e0e6c4d75ae4 |
| Species | *Enterococcus flavescens* | 71.30% | e5e70e59164317ff9c35feeb73956d4b |
| Species | *Enterococcus flavescens* | 71.69% | 029427fe0d93a4c1458a293861ffd731 |
| Species | *Enterococcus flavescens* | 71.59% | d4dda34b25bb40ea08e98c0c1c2bfba8 |
| Species | *Enterococcus flavescens* | 96.23% | 2839059f8683e52e99b059fa50f27c2c |
| Species | *Enterococcus flavescens* | 71.27% | 55c3624f234d38428d7d878b016153ca |
| Species | *Enterococcus flavescens* | 71.28% | accf1f851a9ba775ef70f70343ac4576 |
| Species | *Enterococcus flavescens* | 96.34% | b0ee421bab4925e4d294ed16dbb4860c |
| Species | *Enterococcus flavescens* | 70.20% | 9f2a7d3000b2bcc33e2ccba09100c899 |
| Species | *Enterococcus flavescens* | 73.28% | 3363c35d4e375908b235ae2bf17faaa3 |
| Species | *Enterococcus flavescens* | 70.92% | 3c80b0e44195b9bbd3f6ddd69a2da1ef |
| Species | *Enterococcus flavescens* | 71.28% | ef84f30a783f47b7ff6e9519f148072e |
| Species | *Enterococcus flavescens* | 70.68% | 62d3cf43194780bf3949eb13cdedc2b9 |
| Species | *Enterococcus flavescens* | 71.69% | 7129aa96be7fd020fb3618f539b9652c |
| Species | *Erysipelatoclostridium ramosum* | 81.73% | e86da07f37dfef91c130030d71c4e743 |
| Species | *Erysipelothrix inopinata* | 88.89% | 10eeb5bcdfbfb79c70e1cd3ddcd23f53 |
| Species | *Ferruginibacter lapsinanis* | 97.66% | 39a500ed383046987e3da65e144b59c7 |
| Species | *Finegoldia magna* | 86.55% | 5a5730e7206aad6f93f22bdf81ef4b18 |
| Species | *Flavobacteriaceae bacterium* | 96.47% | aca488f4cc434d53ac6217c9d3edbc4b |
| Species | *Fusobacterium equinum* | 85.87% | c537d1ea07c153f6e8b6b2808a213fc8 |
| Species | *Fusobacterium ulcerans* | 90.66% | 63146d2fb9df0878ed6a125151b2cea1 |
| Species | *Fusobacterium varium* | 92.12% | 3c70d457cdf65a18142fa2385f2fa91f |
| Species | *Gemella sp.* | 77.19% | 5a42f4ab2967b8bb9c6e2f1c4ff31bd7 |
| Species | *Gryllus firmus* | 99.70% | 40c8a38cf0f61fed23b7e9313de966db |
| Species | *Gryllus firmus* | 99.62% | db008f7a3f15698ac6b686da0e1a5006 |
| Species | *Gryllus firmus* | 99.55% | 5c8e8a98b571b54232dc6c175b8dff2a |
| Species | *Halomonas xiaochaidanensis* | 82.08% | 06afdf77bab6622389036b6457bdeeaf |
| Species | *Heliconius cydno* | 99.97% | b91d910a06ca5d497f3bada6029ea983 |
| Species | *Heliconius cydno* | 99.98% | 3ec83fbe38b7b1740f9473c63994c86c |
| Species | *Heliconius numata* | 99.50% | 7f61aa69ee04f2735186b382b9904257 |
| Species | *Heliconius numata* | 99.46% | b2f6ff6d4dcfffe3d0cf331774b979a8 |
| Species | *Heliconius numata* | 99.56% | 455fc931d03c066e23b9683599edfc51 |
| Species | *Heliconius numata* | 99.51% | 4a3658c4df1423a6d2a90ab6b81afa8b |
| Species | *Heliconius numata* | 99.51% | 42d5596d0deba80553f8456391200ccd |
| Species | *Heliconius timareta* | 99.96% | d59e59438f92dbbbcfe9664db17fdde2 |
| Species | *Huanghella arctica* | 99.29% | 8641251cbce6f4ee96a9bdc5c9240151 |
| Species | *Hymenobacter rivuli* | 98.44% | 96e278925eba70aaba34458d500dd34e |
| Species | *Hymenobacter rivuli* | 95.64% | cf9a33e877dcf53fc1a6f0a8227957d2 |
| Species | *Hymenobacter rivuli* | 95.59% | c30e2aafd1902823b0bef3ca3bf39132 |
| Species | *Hymenobacter rivuli* | 83.01% | c04675ccf6ec58d36a00ecb1905cc1c6 |
| Species | *Hymenobacter rivuli* | 74.51% | 51bd2eb430a10bd9240889c69c4d0178 |
| Species | *Lachnoclostridium sp.* | 84.60% | 9e69606bf8685a23ada2bd9ca6503149 |
| Species | *Lactobacillus spicheri* | 94.77% | c88a4ebaf6d4f005dbc465e0a6487fdd |
| Species | *Lawsonella clevelandensis* | 94.49% | b2712cc646bcaebb8eae5345ece18f67 |
| Species | *Leptotrichia goodfellowii* | 99.98% | 7405563b23d5b8a48ace18a211c3c823 |
| Species | *Leuconostocaceae bacterium* | 70.60% | c3101c872b4204675ad788e718d5fb7f |
| Species | *Loriellopsis cavernicola* | 97.74% | f62fd48227ce646ba1a8b70cbf56e5d3 |
| Species | *Luteibacter rhizovicinus* | 85.45% | 454c7e509db3d49cf75d6d469d9832ff |
| Species | *Mannheimia granulomatis* | 70.73% | 1e8f89ce32e320a143c54916442e5b54 |
| Species | *Mannheimia granulomatis* | 73.67% | 7fee6864a1c5c21193a504e87a1eff7d |
| Species | *Mannheimia granulomatis* | 75.34% | df199ef2d2d9ed1b1f55b43a63cc7617 |
| Species | *Mannheimia granulomatis* | 72.20% | 4de93055b8d899d10e4a2bc2c33b72ae |
| Species | *Mannheimia granulomatis* | 72.87% | 8d976cd1b75ecbd0b55e98de3c1a04e8 |
| Species | *Mannheimia granulomatis* | 90.41% | 789af18968faecfc33f373b54a54a2d6 |
| Species | *Mannheimia granulomatis* | 90.72% | 5918c8052194b5e7de281a91d7ad2dc6 |
| Species | *Methylobacterium adhaesivum* | 99.28% | 8019285b0bc60327e30dd97a6151c0fd |
| Species | *Methylobacterium adhaesivum* | 99.24% | 3e1e7b76162d81595b3af625a90eb8c0 |
| Species | *Methylobacterium adhaesivum* | 79.67% | 403483a6695a2fb2bd1b52fbe724b6c1 |
| Species | *Methylobacterium aerolatum* | 99.34% | 1c0d36231f6a2fca4a975142af302a9e |
| Species | *Methylobacterium aerolatum* | 99.33% | 8fd55105ad4294e826b48b24769423d7 |
| Species | *Methylobacterium cerastii* | 83.83% | bd121066d2393e3a2677f44ffc5303e8 |
| Species | *Methylobacterium sp.* | 74.11% | 208bf8422d69807df28e10ce691862b4 |
| Species | *Mycoplasma iguanae* | 87.48% | 157bd650e654e8e01b53c70ed70fb65a |
| Species | *Mycoplasma iguanae* | 87.48% | bdb25c65282256c5418c4fff28f0c194 |
| Species | *Mycoplasma iguanae* | 87.37% | 3d32523b2a9adfeb255c807fcdde7f1c |
| Species | *Mycoplasma iguanae* | 87.48% | fdd0560020a12d8b47ed357f5c88cb33 |
| Species | *Mycoplasma iguanae* | 89.43% | cebf0d550968a49b4ddb393303797740 |
| Species | *Mycoplasma iguanae* | 87.47% | e0f8324cab2e6284ca34020e0539883f |
| Species | *Mycoplasma iguanae* | 87.47% | 780938a29af5ac7ce7ef2db30ac2c1d9 |
| Species | *Mycoplasma iguanae* | 87.36% | 5b5440eb3e03649837fb4c9f092e89a0 |
| Species | *Mycoplasma iguanae* | 87.47% | 70b42feeccd33a350c76b3eb94356de8 |
| Species | *Mycoplasma iguanae* | 87.26% | e848cf16579d76be3d08196e3a721b1e |
| Species | *Mycoplasma iguanae* | 87.36% | f10ed559858e4266fa7d6f9bf4b91d10 |
| Species | *Mycoplasma iguanae* | 87.32% | 804959db9ed87a62bdcaa3de1abf7a46 |
| Species | *Mycoplasma iguanae* | 87.26% | 503b41cf2d46e85c8bfa4f30631b3192 |
| Species | *Mycoplasma iguanae* | 89.62% | 99e756333f8e4c86e7723e50773df230 |
| Species | *Mycoplasma iguanae* | 96.16% | 6db94a89445524ae7e6af22cdf0a10e6 |
| Species | *Mycoplasma iguanae* | 90.74% | d1469b5658f75f431fab9af19696dbee |
| Species | *Mycoplasma iguanae* | 93.81% | 7d4a01af35412fd1e8d366e97aef0b37 |
| Species | *Mycoplasma iguanae* | 92.53% | 9b588e523e44204e00140bc27af6ecd6 |
| Species | *Mycoplasma iguanae* | 93.81% | 8a75081e7eb86592f258882de87d7438 |
| Species | *Mycoplasma iguanae* | 84.98% | b3fdaf3462c6f27143823c68c8034b92 |
| Species | *Mycoplasma iguanae* | 82.38% | db21e5457a513343eb693581b769f832 |
| Species | *Mycoplasma iguanae* | 85.10% | 929dc17ebdd9524864c4b9d5db8083f3 |
| Species | *Mycoplasmopsis columboralis* | 99.82% | e3b25bbe677b2d35ba949473a68c1dfe |
| Species | *Mycoplasmopsis columboralis* | 99.87% | 01d96ba67c21d6eaa4d01381b141b12f |
| Species | *Mycoplasmopsis columboralis* | 99.89% | 5c1af6372edfb35b5e2d070a226be7dd |
| Species | *Mycoplasmopsis columboralis* | 99.87% | bea7fe025b41276519672d98096cf65f |
| Species | *Mycoplasmopsis columboralis* | 99.91% | 4f6611d6a4db92f18258afe3231de62d |
| Species | *Mycoplasmopsis columboralis* | 99.88% | 38cb02922fb43babf221789cbb4bc1a5 |
| Species | *Mycoplasmopsis columboralis* | 99.87% | eab668dddad29526a8733c99c1bfcc98 |
| Species | *Myroides odoratus* | 99.28% | efb896c7d0f319067a1648327586b292 |
| Species | *Nocardioides halotolerans* | 99.02% | 438aee17c9765c4a91ec8721540f80c3 |
| Species | *Novosphingobium nitrogenifigens* | 90.95% | 7bba9813b3332552cfe1fac7f0041bcd |
| Species | *Novosphingobium nitrogenifigens* | 90.81% | 7e5fc29b2551d52c703d27d99c53242e |
| Species | *Novosphingobium resinovorum* | 77.58% | bebc638b41571fd9ab752f6a2ac3accc |
| Species | *Orbus sasakiae* | 72.11% | b29c0d1f0e0a1f11f47f8b77a7e5a433 |
| Species | *Orbus sasakiae* | 74.48% | a1e5ecf56988e99596b8f7ef8c984190 |
| Species | *Orbus sasakiae* | 72.91% | 16b1f6d8fc9dff2f8fa41269797c5a1f |
| Species | *Orbus sasakiae* | 83.51% | 9b975647c565972b3a53ed9c6184e27a |
| Species | *Parabacteroides goldsteinii* | 98.92% | d0d9cc298f3f4cff6c8da85426e58ae7 |
| Species | *Parabacteroides goldsteinii* | 89.25% | fd986a851fc5a0ba21127ca57e5ad00d |
| Species | *Parabacteroides goldsteinii* | 98.95% | e8c197d2d747433e74fe05c0fbdaef6c |
| Species | *Parabacteroides goldsteinii* | 89.34% | 7ab4c67b09e084e38c025b58d92cc278 |
| Species | *Parabacteroides goldsteinii* | 99.49% | 6f459d5bccd5d1798f101ce8eb5b68c2 |
| Species | *Parabacteroides merdae* | 74.68% | b57409e5539402149cada796c4313a51 |
| Species | *Paracoccus marcusii* | 70.81% | 03c10400f997e45e60a516888c528af1 |
| Species | *Pararhodospirillum sulfurexigens* | 80.12% | a79b2b635d92f927551014b0b065937d |
| Species | *Pararhodospirillum sulfurexigens* | 80.63% | dac8f991cfba8e3394c7348ebe2a0030 |
| Species | *Peptoniphilaceae bacterium* | 93.81% | 80370902244ca8e86e3c6ff499af0f32 |
| Species | *Peptoniphilus urinimassiliensis* | 71.66% | 290bb6f69035003dcbd2b77ff7cc5e3f |
| Species | *Peptostreptococcus russellii* | 99.95% | 9ec73bdf1ba1b99dcd0c2f136ce19cea |
| Species | *Peptostreptococcus russellii* | 99.95% | 6cdddb95ec64633e1a416cddeabae69f |
| Species | *Petrimonas sp.* | 98.65% | e046054a2f0c2ed72e149281599460fa |
| Species | *Petrimonas sp.* | 98.65% | 38e09d5f0970941b69e5d46c94112678 |
| Species | *Petrimonas sp.* | 98.63% | 6b3a1bf55671372d6c829c58875fd45c |
| Species | *Petrimonas sp.* | 97.94% | ec9aa5ae80b7d0fd849578fb4f0e06a0 |
| Species | *Porphyromonadaceae bacterium* | 99.12% | cb05ce8f232424dc1342dba283003640 |
| Species | *Porphyromonadaceae bacterium* | 99.14% | 889651aa157ee090d4298ed86ae8efdb |
| Species | *proteobacterium symbiont* | 99.47% | 56340783778cfd33f9b4509cccbbcf21 |
| Species | *Pseudokineococcus lusitanus* | 84.34% | 740ce49698c87c8f4b77c6c348803cd0 |
| Species | *Pseudomonas fragi* | 96.38% | 800d29dc295b5f04cbbb3adeec2ba8b9 |
| Species | *Pseudomonas fragi* | 92.59% | 6e1b1078c83b4a501e5d1e0c15949acb |
| Species | *Pseudomonas fragi* | 95.54% | 883b9022f2538140bd34640d03cd8327 |
| Species | *Pseudomonas fragi* | 94.97% | aad74b763f37a9acd6c30c74ec5f36c5 |
| Species | *Pseudomonas fragi* | 94.03% | 5688266ef6648614063558a0ddd48bcf |
| Species | *Pseudomonas psychrotolerans* | 87.32% | bdec2d5b488593069b84a5cbb6a52e09 |
| Species | *Pseudomonas psychrotolerans* | 87.17% | 1b844b220769216e1abc8519d020fcaa |
| Species | *Pseudomonas psychrotolerans* | 87.85% | 05053227687ef8355f75c663c58822db |
| Species | *Pseudomonas psychrotolerans* | 87.97% | 0429ec3794902197f1f0a2a88503aa57 |
| Species | *Pseudomonas psychrotolerans* | 86.80% | 7e1baf28106ae70e3f904262f977d627 |
| Species | *Pygmaiobacter massiliensis* | 99.44% | 787b8322d2162de5f2e80cfefb9a3d2a |
| Species | *Ralstonia pickettii* | 94.92% | 30bfb6f99468f66abfbd40228576bb77 |
| Species | *Raoultibacter massiliensis* | 90.98% | 2b510fefc55e55432923030df11e1429 |
| Species | *Rickettsia asemboensis* | 90.06% | 426458fa5ae5c7ff10c3fc796a4b6ca8 |
| Species | *Rickettsia asemboensis* | 96.72% | 6400d1319716f9f9c0cb314b7b0a324c |
| Species | *Robinsoniella peoriensis* | 86.28% | 136a7eae571a0bc117f3ab348daf5fdc |
| Species | *Robinsoniella peoriensis* | 83.11% | dbda0ced23ed218a314b5be90eb25a98 |
| Species | *Sanguibacter gelidistatuariae* | 96.21% | 08c65b12b7de343ecd04d59240cf9233 |
| Species | *Siccationidurans occulans* | 82.08% | 9ea572c37796d4090c72a2fb2db94a1c |
| Species | *Siccationidurans soli* | 91.09% | df8846bf2603c944a7cac5f1dd3f0658 |
| Species | *Solirubrobacterales bacterium* | 99.45% | ff9693817969af8a3f631f50b056ae34 |
| Species | *Sphingobacterium kyonggiense* | 80.64% | 51ee7ae813f58d64577f3a256b525ad5 |
| Species | *Sphingobacterium rhinocerotis* | 75.41% | 7c4eb996be03294d331ccb0f6d1ebb87 |
| Species | *Sphingomonas crusticola* | 98.99% | e8a242b0975f90535cd666cc0bbe7e78 |
| Species | *Sphingomonas cynarae* | 70.30% | 7ea8900eb5062ca2ee1690ee6ac15a82 |
| Species | *Sphingomonas faeni* | 99.67% | 04600fe81990d48f673efbc393ea135f |
| Species | *Sphingomonas faeni* | 99.67% | 449091f976fbf8f679e67afd43174f50 |
| Species | *Sphingomonas faeni* | 99.66% | 67a6371d31630b03d9c5c3ad420e4841 |
| Species | *Sphingomonas phyllosphaerae* | 97.87% | 3abb34a99979df470633462fc38694d8 |
| Species | *Sphingomonas phyllosphaerae* | 97.72% | cd1875f207326ed83ca759234dde89b3 |
| Species | *Sphingomonas phyllosphaerae* | 98.43% | 00c6cde5df4907ca46225655f2e1075f |
| Species | *Sphingomonas roseiflava* | 75.12% | f52e9a18bdc8987e249168639480af3f |
| Species | *Sphingomonas roseiflava* | 75.12% | ce7c2acacca5bbc4a4a2245660fc973d |
| Species | *Sphingomonas roseiflava* | 74.93% | 624aa4ef52bef2f2c7ea5f09f902ad8a |
| Species | *Sphingomonas roseiflava* | 73.45% | 4aa0a28ac99e10d30a478b66b31a9134 |
| Species | *Spirosoma oryzae* | 99.95% | 4389ddcec9797073c6c1423cba7fc274 |
| Species | *Spirosoma oryzae* | 99.99% | 1079a501a5d220f30fef25cf5763e548 |
| Species | *Spirosoma oryzae* | 99.95% | 61ac6c4a4af9c8db519b130a270cc2a8 |
| Species | *Spirosoma pomorum* | 95.57% | 0c02277564f701af9642b5a7c407bc07 |
| Species | *Spirosoma pomorum* | 99.91% | a231650d6b41c07fa9f06f37c0a9087a |
| Species | *Staphylococcus carnosus* | 99.99% | baf51b24d5f98baa3d7494e8e8ddd6e8 |
| Species | *Staphylococcus carnosus* | 100.00% | 9442d373256297a1ea403679d7343740 |
| Species | *Staphylococcus carnosus* | 99.99% | 900789544f80d18d0dd8bbd265d18e38 |
| Species | *Staphylococcus carnosus* | 100.00% | 2089418b0fab403c8c51fb991816dc2e |
| Species | *Staphylococcus carnosus* | 99.86% | 590d2a40eb1d8e5bed490c2dfe58fbd5 |
| Species | *Staphylococcus carnosus* | 99.89% | e0a6a6968342f743c34cb757358ce72e |
| Species | *Staphylococcus carnosus* | 100.00% | ed1536b39bfbe8f360e2be051ba75b46 |
| Species | *Staphylococcus carnosus* | 100.00% | 475da4e9d4013ffa6bc00aa0aa8ca003 |
| Species | *Staphylococcus carnosus* | 99.99% | 4c9a4701836f20d6a206a5d468001cfe |
| Species | *Streptococcus pneumoniae* | 100.00% | de01fa220eb44aa99898ddd3316631a6 |
| Species | *Streptococcus pneumoniae* | 100.00% | 9ccbb9bd79a98dea8392197f937b2337 |
| Species | *Streptococcus pneumoniae* | 100.00% | db395ccd0114ec25a2aafe6c05296ded |
| Species | *Streptomyces sp.* | 88.00% | 7a7e8519e6a1789d5d9bda760e81b4c5 |
| Species | *Streptomyces sp.* | 91.16% | b4b71e4b5d6c6f219adc364cd47968ff |
| Species | *Streptomyces sp.* | 72.37% | 52904d58b891dfcb8f2de8478cae239f |
| Species | *Streptomyces sp.* | 82.20% | ffd76d0625f3d3a07ab3f00a31b3427c |
| Species | *Streptomyces sp.* | 79.66% | 2745c98f3a68d28f2c7c03d6c6672834 |
| Species | *Streptomyces sp.* | 99.99% | b5a008dd5d0b336538d6b5283401694f |
| Species | *Streptomyces sp.* | 99.72% | a70fc8072077bf6a008747c2755344b2 |
| Species | *Streptomyces sp.* | 97.97% | d24f531f6b59cf8c1fb69115929d237c |
| Species | *Streptomyces sp.* | 70.34% | 5a2c4763f3487ab0344bb817e805838f |
| Species | *Streptomyces sp.* | 77.26% | c6359f6d8ccedd53bd79998ec06c9da8 |
| Species | *Streptomyces sp.* | 82.36% | ae185fc37080ced41f2a6e8ac78f5e29 |
| Species | *Streptomyces sp.* | 99.97% | 7e21e000b1e6699290b61d34ca912ea0 |
| Species | *Streptomyces sp.* | 99.80% | 0d41ba775d0e31aa686de3c804f84144 |
| Species | *Streptomyces sp.* | 98.01% | 398e3c2d6c892ea5998dde7c71b33578 |
| Species | *Streptomyces sp.* | 99.98% | b075302b85c69a2db8c812f5fc058766 |
| Species | *Streptomyces sp.* | 99.97% | 9ed18a499418e8df75a3d6d2f37085e2 |
| Species | *Streptomyces sp.* | 100.00% | 6189bd389031938493319ef6a0c39fd1 |
| Species | *Streptomyces sp.* | 99.94% | 4d572f5295efefc34e73a3d42c7c9879 |
| Species | *Streptomyces sp.* | 100.00% | 6104ba50eaa7a7ea770e94e6555e2038 |
| Species | *Streptomyces sp.* | 97.36% | 302005b7b652f9536d12a02666ca28f7 |
| Species | *Streptomyces sp.* | 98.11% | 1093e21c662b010634d45f4d5dbe90da |
| Species | *Streptomyces sp.* | 100.00% | 01ee2b460f22ad6c4ea8448f377caac0 |
| Species | *Streptomyces sp.* | 100.00% | ddc1971d97f78401e99b4ffebb8f2da1 |
| Species | *Streptomyces sp.* | 99.99% | 5ee8a99d3fa4ef8da82d1212de8e9aa1 |
| Species | *Streptomyces sp.* | 99.99% | 623aefac464bf0c6730b6e7daaeac8fb |
| Species | *Streptomyces sp.* | 98.66% | 3581b7e06f1b69953de42323915958e8 |
| Species | *Streptomyces sp.* | 100.00% | 3519b70afb36075fd5cd3a2f0bc926dc |
| Species | *Streptomyces sp.* | 100.00% | 69c9afbb4ce81fb05e0c8c9a8f0e700e |
| Species | *Streptomyces sp.* | 95.11% | 3be50c401e27f8ebabb1c69f942118b2 |
| Species | *Streptomyces sp.* | 100.00% | 3b4fb9f1fbd01a808b2f450668e9afe4 |
| Species | *Streptomyces sp.* | 82.13% | 92df125c9c5a0b5bdd7d82b3de554907 |
| Species | *Streptomyces sp.* | 100.00% | 448c80da073c195e6960c6ed9ef0a31d |
| Species | *Streptomyces sp.* | 88.85% | 6a5ce6969434213b1cd8a01145add9ee |
| Species | *Streptomyces sp.* | 95.94% | 76045d140c7152ba0614f117321e8f79 |
| Species | *Terriglobus aquaticus* | 99.75% | 532ba12d310477c4390fd23ff7a7b77c |
| Species | *Vagococcus fluvialis* | 74.94% | 9e30822c297122ea922d039d2f180801 |
| Species | *Vagococcus fluvialis* | 74.33% | ab411eed4d386ec6e941e9a0cdba49ef |
| Species | *Vagococcus fluvialis* | 75.79% | 5c7f9e8c96e69c06c948656063d05ad8 |
| Species | *Vagococcus fluvialis* | 75.79% | 4f6ac13400d5347e5796106f6eecaa6f |
| Species | *Vagococcus teuberi* | 98.57% | 99d5da78d681dd40043e21a5beec83ba |
| Species | *Vagococcus teuberi* | 98.57% | 2682ab4e9b11c65c84d48138645daa2d |
| Species | *Vagococcus teuberi* | 98.57% | f354e872116988230db53261cfc7bb0d |
| Species | *Vagococcus teuberi* | 95.66% | 774c5fa442ff8b5a66255e3c352cccbf |
| Species | *Vagococcus teuberi* | 98.60% | b92d24ae252f3ffa4add8bf1141d1898 |
| Species | *Vibrio penaeicida* | 86.16% | 0a13430bfb746ac289152603436bdfb1 |
| Species | *Wohlfahrtiimonas chitiniclastica* | 92.71% | 38ce0e996edb8432921a6dedaaf627c4 |
| Species | *Xanthomonas albilineans* | 93.92% | 89c46ac3879cf76f91312624e6917e9b |