

feature_Taxon

| | | | | | | | | | |
|--|------|------|------|------|--------|------|------|------|------|
| ASV_3033__csb3; CRISPR-associated protein Csb3 | 591 | 452 | 882 | 1740 | 562 | 24 | 236 | 1081 | 800 |
| ASV_1492__"UNG, UDG; uracil-DNA glycosylase [EC:3.2.2.27]" | 1452 | 14 | 927 | 17 | 1030 | 172 | 295 | 70 | 2588 |
| ASV_2152__mutS2; DNA mismatch repair protein MutS2 | 334 | 64 | 54 | 18 | 42 | 135 | 34 | 2541 | 617 |
| ASV_3097__sum_of_decays_for_non-surrounding/distant_teeth | 3 | 1 | 3 | 1 | 3 | 3 | 3 | 3 | 3 |
| ASV_2929__"pglJ; N-acetylgalactosamine-N,N'-diacetylbaucillosaminyl-diphospho-undecaprenol 4-alpha-N-acetylgalactosaminyltransferase [EC:2.4.1.291]" | 574 | 1255 | 932 | 767 | 27 | 1241 | 335 | 2490 | 1050 |
| ASV_2708__wbpB; UDP-N-acetyl-2-amino-2-deoxyglucuronate dehydrogenase [EC:1.1.1.335] | 66 | 80 | 13 | 73 | 625 | 59 | 16 | 2886 | 461 |
| ASV_2971__vanW; vancomycin resistance protein VanW | 226 | 895 | 32 | 329 | 2895 | 136 | 225 | 31 | 84 |
| ASV_1322__"TC.MATE, SLC47A, norM, mdtK, dinF; multidrug resistance protein, MATE family" | 835 | 160 | 15 | 205 | 971 | 272 | 379 | 205 | 2026 |
| ASV_1268__ureI; acid-activated urea channel | 508 | 204 | 1866 | 83 | 15 | 986 | 142 | 102 | 423 |
| ASV_273__"purF, PPAT; amidophosphoribosyltransferase [EC:2.4.2.14]" | 1975 | 25 | 633 | 212 | 131 | 524 | 565 | 380 | 906 |
| ASV_2001__K07052; uncharacterized protein | 584 | 82 | 1831 | 189 | 941 | 205 | 685 | 26 | 58 |
| ASV_3096__sum_of_dmfs_for_all_other_teeth | 1 | 2 | 2 | 2 | 2 | 1 | 2 | 1 | 2 |
| ASV_253__E2.4.1.5; dextransucrase [EC:2.4.1.5] | 17 | 88 | 5 | 35 | 45 | 76 | 2611 | 1004 | 523 |
| ASV_1204__"RP-S12, MRPS12, rpsL; small subunit ribosomal protein S12" | 501 | 477 | 150 | 6 | 187 | 47 | 358 | 453 | 237 |
| ASV_2954__kce; 3-keto-5-aminohexanoate cleavage enzyme [EC:2.3.1.247] | 1383 | 179 | 106 | 940 | 784 | 405 | 28 | 98 | 124 |
| ASV_1814__dexA; dextranase [EC:3.2.1.11] | 6 | 172 | 318 | 520 | 1085.5 | 67 | 290 | 25 | 107 |
| ASV_2831__nikA; nickel transport system substrate-binding protein | 452 | 147 | 1196 | 36 | 615 | 26 | 187 | 237 | 1458 |
| ASV_3094__spatial-distance-weighted_mean_of_dmfs_of_all_remaining_teeth | 2 | 3 | 1 | 3 | 1 | 2 | 1 | 2 | 1 |
| ASV_2655__pchF; pyochelin synthetase | 25 | 28 | 12 | 222 | 5 | 74 | 6 | 6 | 22 |
| ASV_2705__rgpE; glucosyltransferase [EC:2.4.1.-] | 5 | 5 | 6 | 5 | 17 | 5 | 5 | 5 | 5 |
| ASV_1506__hslJ; heat shock protein HslJ | 20 | 307 | 70 | 88 | 754 | 1103 | 389 | 748 | 12 |
| ASV_2210__"lytS; two-component system, LytTR family, sensor histidine kinase LytS [EC:2.7.13.3]" | 1320 | 422 | 972 | 10 | 550 | 7 | 353 | 20 | 19 |
| ASV_2338__rutE; 3-hydroxypropanoate dehydrogenase [EC:1.1.1.-] | 2718 | 334 | 22 | 1193 | 79 | 516 | 146 | 657 | 524 |
| ASV_2432__ABC.MN.A; manganese/iron transport system ATP-binding protein | 890 | 34 | 261 | 893 | 22 | 2471 | 24 | 324 | 263 |
| ASV_1756__"mnhF, mrpF; multicomponent Na+:H+ antiporter subunit F" | 147 | 645 | 319 | 49 | 1193 | 425 | 37 | 17 | 272 |
| ASV_2834__nikD; nickel transport system ATP-binding protein [EC:3.6.3.24] | 2621 | 54 | 1245 | 45 | 797 | 29 | 115 | 155 | 1053 |
| ASV_2758__aroKB; shikimate kinase / 3-dehydroquinase synthase [EC:2.7.1.71 4.2.3.4] | 1378 | 289 | 71 | 90 | 48 | 10 | 35 | 326 | 8 |
| ASV_2765__pfbA; plasmin and fibronectin-binding protein A | 811 | 677 | 53 | 2797 | 47 | 1190 | 43 | 711 | 9 |
| ASV_922__"cobC, phpB; alpha-ribazole phosphatase [EC:3.1.3.73]" | 232 | 2624 | 83 | 22 | 281 | 18 | 7 | 21 | 252 |
| ASV_905__bioH; pimeloyl-[acyl-carrier protein] methyl ester esterase [EC:3.1.1.85] | 80 | 17 | 367 | 217 | 2601 | 717 | 1280 | 882 | 1100 |
| ASV_3098__sum_of_decays_for_surrounding_teeth | | 4 | 4 | 4 | 4 | 4 | 4 | 4 | 4 |
| ASV_737__"WARS, trpS; tryptophanyl-tRNA synthetase [EC:6.1.1.2]" | 1380 | 647 | 25 | 12 | 90 | 373 | 205 | 2408 | 323 |

