

Supplementary Table 6. Frequency that each bacterial sequence or OTU was retrieved by cultivation or by culture-independent methods.

| Phylum | Sequences | | | OTUs | | |
|-----------------------|-----------|------------|---------|----------|------------|-------|
| | Cultured | Uncultured | Total | Cultured | Uncultured | Total |
| Proteobacteria | 119742 | 293919 | 413661 | 4927 | 21726 | 26653 |
| Firmicutes | 84752 | 392921 | 477673 | 2575 | 19489 | 22064 |
| Bacteroidetes | 9744 | 155663 | 165407 | 1590 | 11840 | 13430 |
| Actinobacteria | 41122 | 178892 | 220014 | 1687 | 4023 | 5710 |
| Planctomycetes | 539 | 15008 | 15547 | 126 | 4236 | 4362 |
| Chloroflexi | 197 | 23202 | 23399 | 64 | 4095 | 4159 |
| Acidobacteria | 224 | 15683 | 15907 | 101 | 3714 | 3815 |
| Spirochaetae | 2105 | 9723 | 11828 | 107 | 1546 | 1653 |
| Cyanobacteria | 3541 | 14656 | 18197 | 415 | 1234 | 1649 |
| Verrucomicrobia | 177 | 8436 | 8613 | 81 | 1459 | 1540 |
| Gemmatimonadetes | 23 | 2890 | 2913 | 17 | 1003 | 1020 |
| Lentisphaerae | 18 | 3593 | 3611 | 9 | 898 | 907 |
| Parcubacteria | 2 | 1298 | 1300 | 2 | 900 | 902 |
| Saccharibacteria | 10 | 2567 | 2577 | 10 | 869 | 879 |
| Tenericutes | 2182 | 3564 | 5746 | 198 | 664 | 862 |
| Armatimonadetes | 10 | 1412 | 1422 | 5 | 746 | 751 |
| Nitrospirae | 112 | 2897 | 3009 | 24 | 596 | 620 |
| Microgenomates | 0 | 779 | 779 | 0 | 592 | 592 |
| TM6 | 5 | 797 | 802 | 3 | 541 | 544 |
| Hydrogenedentes | 5 | 950 | 955 | 2 | 507 | 509 |
| Chlorobi | 83 | 1360 | 1443 | 17 | 415 | 432 |
| Gracilibacteria | 3 | 1143 | 1146 | 3 | 423 | 426 |
| Latescibacteria | 1 | 997 | 998 | 1 | 416 | 417 |
| Marinimicrobia | 8 | 1307 | 1315 | 3 | 410 | 413 |
| OP3 | 0 | 575 | 575 | 0 | 379 | 379 |
| Fibrobacteres | 62 | 2480 | 2542 | 9 | 364 | 373 |
| Aminicenantes | 0 | 1734 | 1734 | 0 | 365 | 365 |
| TA06 | 0 | 574 | 574 | 0 | 325 | 325 |
| Synergistetes | 103 | 1724 | 1827 | 21 | 281 | 302 |
| Deinococcus-Thermus | 554 | 1599 | 2153 | 89 | 201 | 290 |
| Atribacteria | 4 | 2515 | 2519 | 4 | 256 | 260 |
| Deferribacteres | 44 | 1505 | 1549 | 17 | 212 | 229 |
| Elusimicrobia | 4 | 458 | 462 | 3 | 225 | 228 |
| Kazan-3B-09 | 0 | 1236 | 1236 | 0 | 206 | 206 |
| Fusobacteria | 534 | 8926 | 9460 | 27 | 173 | 200 |
| Omnitrophica | 0 | 633 | 633 | 0 | 188 | 188 |
| WS6 | 0 | 240 | 240 | 0 | 172 | 172 |
| Acetothermia | 0 | 299 | 299 | 0 | 163 | 163 |
| Thermotogae | 156 | 650 | 806 | 32 | 112 | 144 |
| SHA-109 | 0 | 175 | 175 | 0 | 120 | 120 |
| Aerophobetes | 1 | 276 | 277 | 1 | 118 | 119 |
| PAUC34f | 1 | 169 | 170 | 1 | 113 | 114 |
| Chlamydiae | 536 | 187 | 723 | 49 | 63 | 112 |
| Hyd24-12 | 0 | 242 | 242 | 0 | 111 | 111 |
| WD272 | 0 | 140 | 140 | 0 | 106 | 106 |
| SR1 | 0 | 197 | 197 | 0 | 93 | 93 |
| Cloacimonetes | 2 | 443 | 445 | 1 | 85 | 86 |
| SM2F11 | 0 | 107 | 107 | 0 | 75 | 75 |
| Aquificae | 211 | 702 | 913 | 34 | 27 | 61 |
| WCHB1-60 | 0 | 70 | 70 | 0 | 60 | 60 |
| Caldiserica | 4 | 249 | 253 | 2 | 54 | 56 |
| JL-ETNP-Z39 | 0 | 45 | 45 | 0 | 38 | 38 |
| GOUTA4 | 0 | 31 | 31 | 0 | 28 | 28 |
| LCP-89 | 0 | 32 | 32 | 0 | 25 | 25 |
| Thermodesulfobacteria | 21 | 97 | 118 | 8 | 15 | 23 |
| CKC4 | 2 | 40 | 42 | 2 | 19 | 21 |
| Dictyoglomi | 11 | 17 | 28 | 1 | 11 | 12 |
| GAL08 | 0 | 17 | 17 | 0 | 11 | 11 |
| LD1-PA38 | 0 | 9 | 9 | 0 | 9 | 9 |
| OC31 | 0 | 7 | 7 | 0 | 6 | 6 |
| SBYG-2791 | 0 | 16 | 16 | 0 | 6 | 6 |
| Chrysiogenetes | 12 | 1 | 13 | 5 | 0 | 5 |
| RsaHF231 | 0 | 7 | 7 | 0 | 5 | 5 |
| Caescamantes | 0 | 3 | 3 | 0 | 3 | 3 |
| S2R-29 | 0 | 2 | 2 | 0 | 2 | 2 |
| aquifer1 | 0 | 1 | 1 | 0 | 1 | 1 |
| Total | 266867 | 1162087 | 1428954 | 12273 | 87168 | 99441 |