

Total aligned length

Aligned length (kbp)

10
8
6
4
2
0

final.contigs

Illumatobacter coccineus YM16_304
 marine_gamma_proteobacterium_HTC2080
 Synechococcus sp_ KORDI 49
 endosymbiont_of_Ridgeia pisceae
 Ulvibacter litoralis
 Sulfatobacter sp. NAS 14.1
 Nitrospina gracilis
 Thiohalobacter thiocyanicus
 gamma_proteobacterium_SCGC AB 629 P17
 Pseudomonas aeruginosa
 Nitrospina sp. SCGC AAA288 L16
 Methylobacter sp. XLMV4
 Owenweeksia hongkongensis DSM 17368
 Oceanicola granulosus HTC2516
 Thioalkalibacter sp. AL17
 Thioalkalibacter HL-EbGr7
 Thiobacillus sulfidophilus
 Thiobacillus skagerakense
 Marinomonas communis
 Fluviicola taffensis DSM 16823
 alpha_proteobacterium BAL199
 Alcanivorax pacificus W115
 Nocardioides albus
 Simidula sp. J MY2
 Paracoccus aminophilus
 Magnetospira sp. OH 2
 Porticoccus hydrocarbonoclasticus MCTG13d
 Conexibacter woesei Iso977N
 Syntrophus aciditrophicus
 Desulfatibacter alkalitolans DSM 16504
 Aurantimonas manganoxydans SI85 9A1
 Desulfatimonas thiothiophila
 Sphingobium fuliginis
 Magnetococcus marinus MC 1
 Xanthomonas citri subsp. malvacearum
 Shewanella amazonensis SB2B
 marine metagenome
 Neisseria sp. CSL 7565
 Gracilibacteria_bacterium_JGI-0000069_P22
 Mollicutes bacterium HR1
 mine_drainage_metagenome