

Genome fraction (%)

5
4
3
2
1
0

• final.contigs

gamma_proteobacterium_SCGC_AB_629_P17
 Synechococcus sp. KORDI_49
 marine_gamma_proteobacterium_HTC2080
 lummatobacter_coccineus_YM16_304
 endosymbiont_of_Ridgella_piscisae
 Nitrospina sp. SCGC_AA288_L16
 Roseobacter sp. SK209_2.6
 Nitrospina gracilis
 Sulfitobacter sp. NAS_14.1
 Thiohalobacter thiocyanaticus
 Pseudoruegeria aquimaris
 Ulvibacter litoralis
 Oceanicola granulosus_HTC2516
 Porticoccus_hydrocarbonoclasticus_MCTG13d
 Tenacibaculum skagerrakense
 Thioalkalivibrio sp. AL117
 Thioalkalivibrio sulfidophilus_HL_EbGr7
 Owenweeksia hongkongensis_DSM_17368
 Fluviicola taihensis_DSM_17369
 Alcanivorax pacificus_W11_5
 Paracoccus aminophilus
 Gaella occulta
 Simidula sp. MY2
 Sinobacterium calidoides
 alpha_proteobacterium_BAL199
 Streptomyces_cyaneogriseus_subsp._noncyanogenus
 Desulfuromonas thiophila
 Desulfitibacter_alkalitolerans_DSM_16504
 Nocardioides albus
 Syntrophus aciditrophicus
 Aurantimonas_manganoxydans_S185_9A1
 Neisseria sp. CSL_7565
 Shewanella amazonensis_SB2B
 Conexibacter woesei_Is0977N
 Sphingobium fuliginis
 Xanthomonas citri_subsp._malvacearum
 Mollicutes bacterium_HR1
 Rhodopirellula sp. SM50
 marine_metagenome
 Gracilibacteria_bacterium_JGI_0000069_P22
 mine_drainage_metagenome