

Genome fraction (%)

5

4

3

2

1

0

final.contigs

gamma_proteobacterium_SCGC_AB_629_P17
 marine_gamma_proteobacterium_HTC2080
 Illuminobacter_coccineus_YM16_304
 endosymbiont_of_Ridgerea_piscisae
 Nitrospina_sp._SCGC_AAA288_L16
 Sulfatobacter_sp._NAS_14_1
 Nitrospina_gracilis
 Pseudoruegeria_aquamaris
 Thiohalobacter_thiocyanaticus
 Ulvibacter_litoralis
 Oceanicola_granulosus_HTC2516
 Thioalkalibrio_sp._AB17
 Thioalkalibrio_sulfidiphilus_HL_EbGr7
 Tenacibaculum_skagerakense
 Owenweeksia_hongkongensis_DSM_17368
 Methyllobacter_sp._XLMV4
 Marinomonas_communis
 Alcanivorax_pacificus_W11_5
 Fluviicola_taffensis_DSM_16823
 Porticoccus_hydrocarbonoclasticus_MCTG13d
 Paracoccus_aminophilus
 Simidua_sp._J_MY2
 Magnetospira_sp._QH_2
 Nocardioides_albus
 alpha_proteobacterium_BAL199
 Desulfuromonas_thiophila
 Syntrophus_aciditrophicus
 Aurantimonas_manganoxydans_SI85_9A1
 Conexibacter_woesei_Iso977N
 Desulfatibacter_alkalitolerans_DSM_16504
 Magnetococcus_marinus_MC_1
 Shewanella_amazonensis_SB2B
 Xanthomonas_citri_subsp._malvacearum
 Neisseria_sp._CSL_7565
 marine_metagenome
 Gracilibacteria_bacterium_GJ_0000069_P22
 Mollicutes_bacterium_HR1
 mine_drainage_metagenome