

misassemblies

5
4
3
2
1
0

• final.contigs

Alcanivorax pacificus W11 5
Aurantimonas manganoxydans S185 9A1
Desulfatibacter_alkalitolerans DSM_16504
Desulfatibacter_Desulfuromonas thiothophila
Flavobacterium sp._SCGC_AAA160_P02
Fluviicola taffensis DSM_16823
Geobacter daltonii FRC 32
Gracilibacteria bacterium JGI_0000069_P22
Lysiniphilus syllensis NOR5 1B
Magnetococcus marinus MC 1
Magnetospira sp._QH 2
Marinomonas communis
Methylobacter sp._XLMV4
Mollicutes bacterium HR1
Mycoplasma_arthritis 1581.3 1
Nitrospina_sp._SCGC_AAA288_L16
Oceanicola granulatus HTCC2516
Owenweeksia_hongkongensis DSM_17368
Paracoccus_aminophilus
Phaeocystidibacter sp._G18
Porticoccus_hydrocarbonoclasticus MCTG13d
Pseudoruegeria_aquimaris
Shewanella_amazonensis SB2B
Sphingobium_tuliginis
Sulfitobacter sp._NAS_14_1
Synechococcus sp._WH_8102
Syntrophus_aciditrophicus SB
Syntrophus aciditrophicus SB
Tenacibaculum skagerakense
Thioalkalivibrio sp._AL17
Thioalkalivibrio sulfidiphilus HL_EbGr7
Thiohalobacter thiocyanaticus
Xanthomonas citri subsp. malvacearum
alpha proteobacterium BAL199
endosymbiont of Ridgeia piscesae
gamma proteobacterium SCGC_AB_629_P17
marine_gamma_proteobacterium HTCC2080
mine_drainage_metagenome