WP 049907325036363614748750666 \$#\\dag{\partial parties \partial parties \ ELY83688.1 (JCM12890) (Natrinema altunense JCM12890)) MGYP000597840497 WP_007110832.1 - WP_144240185.1 (Natrinema altunense (AJ2)) AntAceMinimDraft_18_1070375.scaffolds.fasta_scaffold06785_2 (TBA) WP_174701778.1 (Haloterrigena sp. SYSU A121-1) AntAceMinimDraft_4_1070372.scaffolds.fasta_scaffold17623_2 (TBA) LKMP01000007_1 (Nanohaloarchaea archaeon B1-Br10_U2g21) AntAceMinimDraft_18_1070375.scaffolds.fasta_scaffold72169_1 (1 ORF scaffold) mc1328965482|Representative=gi|998585003|gb|LFCJ01001378.1|_8/1661 (TBA) 3300014208-Ga0172379-10000243-871512-158 (Methanomicrobia?) $MGYP000598426430-1548 \ (Archaea; Euryarchaeota; Halobacteria; Halobacteriales)$ NJD53946.1 (Candidatus Methanoperedens sp.) RLI53188.1 (Candidatus Thorarchaeota archaeon) 3300014613-Ga0180008-1001212-875221-12 GYP000179698888-RLG58774.1 (Candidatus Geothermarchaeota B85_G16) 3300014206-Ga0172377-10000119-870930-129 (Methanomicr HHR27186.1 (Candidatus Bathyarchaeota archaeon) 3300014208-Ga0172379-10001592-871560-40 (Methano MGYP000681001372 SAMEA2619974_10776_4 nDraft_18_1070375.scaffolds.fasta_scaffold31711_5 (1 ORF scaffold) RLG94066.1 (Candidatus Bathyarchaeota archaeor inimDraft_18_1070375.scaffolds.fasta_scaffold00713_2 (TBA) MGYP000565992219 Draft_18_1070375.scaffolds.fasta_scaffold50355_1 (1 ORF scaffold) 3300018015-Ga0187866-1000629-915963-9 imDraft_10_1070366.scaffolds.fasta_scaffold00069_38 (TBA) Ace Lake-Antarctica -1402-Ga0222697_1002427_6 (TBA) HGF63239.1 (Candidatus Micrarchaeota archaeon) HDO79935.1 or HEX32987.1 (Candidatus Aenigmarchaeota 003719155.1 HDD44259.1 (Candidatus Desulfofervidus auxili 000189635.1 MGYP000347782300 000190715.1 RKX41251.1 (**Thermotogae bacterium) HAP2s 000313135.1 HDI72891.1 (**Candidatus Altiarchaeales archae 001179505.1 MGYP000335988725 HID09282.1 (Candidatus Micrarchaeota archae WP_179268568.1 (Halobonum sp. NJ-3-1) WP_163487151.1 (Halogeometricum borinquense strain vis AntAceMinimDraft_18_1070375.scaffolds.fasta_scaffold06 AntAceMinimDraft_18_1070375.scaffolds.fasta_scaffold15402_2 (7 ntAceMinimDraft_4_1070372.scaffolds.fasta_scaffold07707_5 (TBA) AntAceMinimDraft_14_1070370.scaffolds.fasta_scaffold61386_1 raft_18_1070375.scaffolds.fasta_scaffold82302_1 (1 ORF scaffold) AntAceMinimDraft_10_1070366.scaffolds.fasta_scaffold03294 imDraft_4_1070372.scaffolds.fasta_scaffold00557_40 (TBA) AntAceMinimDraft_10_1070366.scaffolds.fasta_scaffold1267 imDraft_4_1070372.scaffolds.fasta_scaffold18285_3 (TBA) AntAceMinimDraft_18_1070375.scaffolds.fasta_scaffold20 imDraft_18_1070375.scaffolds.fasta_scaffold11370_2 (TBA) AntAceMinimDraft_4_1070372.scaffolds.fasta_scaffold10520_11 mDraft_18_1070375.scaffolds.fasta_scaffold02562_2 (TBA) AntAceMinimDraft_14_1070370.scaffolds.fasta_scaffold00694 highO2_12_1023870.scaffolds.fasta_scaffold04391_3 (TBA) MGYP000692789748 Draft_4_1070372.scaffolds.fasta_scaffold34113_4 (1 ORF scaffold) AntAceMinimDraft_10_1070366.scaffolds.fasta_scaffold19757_2 nDraft_18_1070375.scaffolds.fasta_scaffold55385_1 (1 ORF scaffold) AntAceMinimDraft_18_1070375.scaffolds.fasta_scaffold02754_4 imDraft_18_1070375.scaffolds.fasta_scaffold36732_2 (1 ORF scaffold) MGYP000559433360 AntAceMinimDraft_18_1070375.scaffolds.fasta_scaffold01114_5 (TBA) AntAceMinimDraft_18_1070375.scaffolds.fasta_scaffold13455_2 (TBA) AntAceMinimDraft_10_1070366.scaffolds.fasta_scaffold24857_2 (TBA) AntAceMinimDraft_4_1070372.scaffolds.fasta_scaffold00827_26 (TBA) AntAceMinimDraft_4_1070372.scaffolds.fasta_scaffold70733_1 (1 ORF scaffold) AntAceMinimDraft_10_1070366.scaffolds.fasta_scaffold06212_4 (TBA) AntAceMinimDraft_18_1070375.scaffolds.fasta_scaffold26691_1 (1 ORF scaffold) mc1330463358|Representative=gi|984778642|gb|LFUF01005182.1|_1/1417 (TBA) AntAceMinimDraft_4_1070372.scaffolds.fasta_scaffold83478_1 (1 ORF scaffold) AntAceMinimDraft_4_1070372.scaffolds.fasta_scaffold28048_2 (TBA) AntAceMinimDraft_10_1070366.scaffolds.fasta_scaffold00511_6 (TBA) PlalllAssembly_1097288.scaffolds.fasta_scaffold00281_9 (TBA) NOZ47386.1 (Chlorobi bacterium)

MGYP000359581082

MGYP000672152970

RKZ11204.1 (Candidatus Fermentibacteria bacterium)

RKZ11204.1 (Candidatus Fermentibacteria bacterium) MGYP000359581082