

Genome sequences published outside of *Standards in Genomic Sciences*, January-March 2012

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The purpose of this table is to provide the community with a citable record of publications of ongoing genome sequencing projects that have led to a publication in the scientific literature. While our goal is to make the list complete, there is no guarantee that we may have omitted one or more publications appearing in this time frame. Readers and authors who wish to have publications added to subsequent versions of this list are invited to provide the bibliographic data for such references to the SIGS editorial office.

Phylum *Crenarchaeota*

Pyrobaculum strain 1860, sequence accession CP003098 [1]

Phylum *Deinococcus-Thermus*

"*Thermus sp.*" Strain CCB_US3_UF1, sequence accession CP003126 (chromosome), CP003127 (plasmid) [2]

Phylum *Proteobacteria*

"*Achromobacter arsenitoxydans*" SY8, sequence accession AGUF00000000 [3]

Acidovorax sp. Strain NO1, sequence accession AGTS00000000 [4]

Acinetobacter baumannii AB4857, sequence accession AHAG00000000 [5]

Acinetobacter baumannii AB5075, sequence accession AHAH00000000 [5]

Acinetobacter baumannii AB5256, sequence accession AHAI00000000 [5]

Acinetobacter baumannii AB5711, sequence accession AHAJ00000000 [5]

Aeromonas salmonicida, sequence accession AGVO00000000 [6]

Aggregatibacter actinomycetemcomitans RHAA1, sequence accession AHGR00000000 [7]

Agrobacterium tumefaciens 5A, sequence accession AGVZ00000000 [8]

Azoarcus sp. Strain KH32C, sequence accession AP012304, AP012305 [9]

Burkholderia sp. Strain YI23, sequence accession CP003087 (Chromosome 1), CP003088 (Chromosome 2), CP003089 (Chromosome 3), CP003090 (plasmid BYI23_D), CP003091 (plasmid BYI23_E) CP003092 (plasmid BYI23_F) [10]

Brucella suis VBI22, sequence accession CP003128, CP003129 [11]

"*Comamonas testosteroni*" ATCC 11996, sequence accession AHIL00000000 [12]

Commensalibacter intestini A911^T, sequence accession AGFR00000000 [13]

Edwardsiella ictaluri, sequence accession CP001600.1 [14]

Enterobacter cloacae subsp. *dissolvens* SDM, sequence accession AGSY00000000 [15]

"*Gluconobacter morbifer*" G707^T, sequence accession AGQV00000000 [16]

Legionella dumoffii TEX-KL, sequence accession AGVT00000000 [17]

Legionella dumoffii NY-23, sequence accession AGVU00000000 [17]

Legionella pneumophila serogroup 12 Strain 570-CO-H, sequence accession CP003192 [18]

Marinobacterium stanieri S30, sequence accession AFPL00000000 [19]

"*Marinobacter manganoxydans*" MnI7-9, sequence accession CP001978 to CP001980 [20]

Mesorhizobium alhagi CCNWXJ12-2T, sequence accession AHAM00000000 [21]

Mesorhizobium amorphae, sequence accession AGSN00000000 [22]

Methylomicrobium alcaliphilum 20Z, sequence accession FO082060 and FO082061 [23]

Mitsuaria sp. Strain H24L5A, sequence accession CAFG01000001 to CAFG01000607 [24]

Novosphingobium pentaromativorans US6-1, sequence accession AGFM00000000 [25]

Pantoea ananatis B1-9, sequence accession CAEI01000001 to CAEI01000169 [26]

Pantoea ananatis LMG 5342, sequence accession HE617160 (chromosome), HE617161 (pPANA10) [27]

Pantoea ananatis Strain PA13, sequence accession CP003085 and CP003086 [28]

Pseudomonas aeruginosa, sequence accession AFXI00000000 [29]

Pseudomonas aeruginosa, sequence accession AFXJ00000000 [29]

Pseudomonas aeruginosa, sequence accession AFXK00000000 [29]

Pseudomonas chlororaphis GP72, sequence accession AHAY01000000 [30]

Pseudomonas fluorescens F113, sequence accession CP003150 [31]

Pseudomonas fluorescens Wayne 1R, sequence accession CADX01000001 to CADX01000090 [32]

Pseudomonas fluorescens Wood1R, sequence accession CAFF01000001 to CAFF01001437 [32]

Pseudomonas psychrotolerans L19, sequence accession AHBD00000000 [33]

Pseudoalteromonas rubra ATCC 29570^T, sequence accession AHCD00000000 [34]

Pseudomonas stutzeri SDM-LAC, sequence accession AGSX00000000 [35]

Pseudoxanthomonas spadix BD-a59, sequence accession CP003093 [36]

Rickettsia slovacica, sequence accession CP002428 [37]

Salmonella enterica serovar Pullorum RKS5078, sequence accession CP003047 [38]

Sinorhizobium meliloti CCNWSX0020, sequence accession AGVV00000000 [39]

Sphingobium sp. Strain SYK-6, sequence accession AP012222 and AP012223 [40]

Sphingomonas sp. Strain PAMC 26605, sequence accession AHIS00000000 [41]

Stenotrophomonas maltophilia RR-10, sequence accession AGRB00000000 [42]

Strain HIMB30, sequence accession AGIG00000000 [43]

Taylorella equigenitalis, sequence accession CP003059 [44]

Vibrio campbellii DS40M4, sequence accession AGIE00000000 [45]

Vibrio fischeri SR5, sequence accession AHIH00000000 [46]

Yersinia enterocolitica, sequence accession AGQO00000000 [47]

Phylum *Tenericutes*

Candidatus Mycoplasma haemominutum, sequence accession HE613254 [48]

Mycoplasma haemocanis strain Illinois, sequence accession CP003199 [49]

Mycoplasma iowae, sequence accession AGFP00000000 [50]

Mycoplasma pneumoniae Type 2a Strain 309, sequence accession AP012303 [51]

Phylum *Firmicutes*

Bacillus cereus F837/76, sequence accession CP003187 (chromosome) CP003188 (pF837_55kb), CP003189 (pF837_10kb) [52]

Brevibacillus laterosporus Strain GI-9, sequence accession CAGD01000001 to CAGD01000061 [53]

Clostridium sporogenes PA 3679, sequence accession AGAH00000000 [54]

Enterococcus mundtii CRL1656, sequence accession AFWZ00000000.1 [55]

Geobacillus thermoleovorans CCB_US3_UF5, sequence accession CP003125 [56]

Lactobacillus curvatus Strain CRL705, sequence accession AGBU01000000 [57]

Lactobacillus rhamnosus ATCC 8530, sequence accession CP003094 [58]

Lactobacillus rhamnosus R0011, sequence accession AGKC00000000 [59]

Lactococcus garvieae TB25, sequence accession AGQX01000000 [60]

Lactococcus garvieae LG9, sequence accession AGQY01000000 [60]

Lactococcus lactis subsp. cremoris A76, sequence accession CP003132 (chromosome), CP003136

(pQA505), CP003135 (PQA518), CP003134 (pQA549), CP003133 (pQA554) [61]

Leuconostoc citreum LBAE C10, sequence accession CAGE00000000 [62]

Leuconostoc citreum LBAE C11, sequence accession CAGF00000000 [62]

Leuconostoc citreum LBAE E16, sequence accession CAGG00000000 [62]

Leuconostoc mesenteroides subsp. *mesenteroides* Strain J18, sequence accession CP003101 [63]

Paenibacillus peoriae Strain KCTC 3763^T, sequence accession AGFX00000000 [64]

Pediococcus acidilactici MA18/5M, sequence accession AGKB00000000 [65]

Pediococcus clausenii ATCC BAA-344^T, sequence accession CP003137 (chromosome), CP003138 (pPECL-1), CP003139 (pPECL-2), CP003140 (pPECL-3), CP003141 (pPECL-4), CP003142 (pPECL-5), CP003143 (pPECL-6), CP003144 (pPECL-7), CP003145 (pPECL-8) [66]

Staphylococcus aureus M013, sequence accession CP003166 [67]

Staphylococcus aureus subsp. *aureus* TW20, sequence accession FN433596 [68]

Weissella confusa LBAE C39-2, sequence accession CAGH00000000 [69]

Phylum Actinobacteria

Corynebacterium casei, sequence accession CAFW01000001 to CAFW01000106 [70]

Corynebacterium glutamicum, sequence accession AGQQ00000000 [71]

Leucobacter chromiirestiens, sequence accession AGCW00000000 [72]

Mycobacterium abscessus, sequence accession AGQU00000000 [73]

Propionibacterium acnes ST9, sequence accession CP003195 [74]

Propionibacterium acnes ST22, sequence accession CP003196 [74]

Propionibacterium acnes ST27, sequence accession CP003197 [74]

Saccharomonospora azurea SZMC 14600, sequence accession AHBX00000000 [75]

Streptomyces sp. Strain TOR3209, sequence accession AGNH00000000 [76]

Streptomyces sp. Strain W007, sequence accession AGSW00000000 [77]

Phylum Spirochaetes

Borrelia valaisiana VS116, sequence accession ABCY02000001 (chromosome), CP001439 (plasmid Ip17), CP001437 (Ip25), CP001440 (plasmid Ip 28-3), CP001442 (plasmid Ip28-8), CP001436 (Ip 36), CP001433 (plasmid Ip 54), CP001438 (plasmid cp9), CP001432 (plasmid cp26), CP001441 (plasmid cp32-5), CP001434 (plasmid cp32-7), CP001435 (plasmid cp32-10) [78]

"Borrelia bissetii" DN127, sequence accession CP002746 (chromosome), CP002756 (plasmid Ip12), CP002757 (plasmid Ip25), CP002758 (plasmid 28-3), CP002759 (plasmid Ip 28-4), CP002760 (Ip28-7), CP002761 (plasmid Ip54), CP002762 (plasmid Ip56), CP002755 (plasmid cp9), CP002747 (plasmid cp26), CP002749 (plasmid cp32-3), CP002750 (plasmid cp32-4), CP002751 (plasmid 32-5), CP002752 (plasmid cp32-6), CP002753 (plasmid cp32-7), CP0027554 (plasmid cp32-9), CP002748 (plasmid cp32-11) [78]

Borrelia spielmanii A14S, sequence accession ABKB02000001 (chromosome), CP001468 (plasmid Ip17), CP001471 (Ip28-3), CP001470 (plasmid Ip28-4), CP001465 (plasmid Ip28-2), CP001466 (plasmid Ip36), CP001464 (plasmid Ip38), CP001469 (plasmid Ip54), ABKB02000016, ABKB02000020 (plasmid cp9), CP001467 (plasmid cp26), ABKB02000026 (plasmid cp32-3), ABKB02000031 (plasmid 32-5), ABKB02000021 (plasmid cp32-12), ABKB02000014 (unidentified) [78]

Non-Bacterial genomes

Aspergillus flavus, sequence accession GSE32177 [79]

Bacteriophage SPN3UB, sequence accession JQ288021 [80]

Bamboo mitochondria, sequence accession JQ235166 to JQ235179 [81]

Boea hygrometrica chloroplast, sequence accession JN107811 [82]

Boea hygrometrica mitochondrial, sequence accession JN107812 [82]

Canine Picornavirus, sequence accession JN831356 [83]

Chandipura virus (CHPV) CIN0327, sequence accession GU212856.1 [84]

Chandipura virus (CHPV) CIN0451, sequence accession GU212857.1 [84]

Chandipura virus (CHPV) CIN0751, sequence accession GU212858.1 [84]
 Chandipura virus (CHPV) CIN0755, sequence accession GU190711.1 [84]
 Chinese Porcine Parvovirus Strain PPV2010, sequence accession JN872448 [85]
 Common midwife toad megavirus, sequence accession JQ231222 [86]
 Dengue Virus Serotype 4, sequence accession JN983813 [87]
 Duck Tembusu Virus, sequence accession JF270480 [88]
 Duck Tembusu Virus, sequence accession JQ314464 [88]
 Duck Tembusu Virus, sequence accession JQ314465 [88]
 Emiliana huxleyi Virus 202, sequence accession HQ634145 [89]
 Emiliana huxleyi Virus EhV-88, sequence accession JF974310 [89]
 Emiliana huxleyi EhV-201, sequence accession JF974311 [89]
 Emiliana huxleyi EhV-207, sequence accession JF974317 [89]
 Emiliana huxleyi EhV-208, sequence accession JF974318 [89]
 Glarea lozoyensis, sequence accession GUE0000000 [90]

Nannochloropsis gaditana, sequence accession AGNI00000000 [91]
Oryza sativa cv., sequence accession DRA000499 [92]
 Partetravirus, sequence accession JN990269 [93]
 Porcine Bocavirus PBoV5, sequence accession JN831651 [94]
 Porcine epidemic diarrhea virus, sequence accession JQ282909 [95]
 Pseudomonas aeruginosa lytic bacteriophage PA1Ø, sequence accession HM624080 [96]
 Pseudomonas fluorescens phage OBP, sequence accession JN627160 [97]
 RNA Virus from Avocado, sequence accession JN880414 [98]
 Salmonella enterica Serovar Typhimurium Bacteriophage SPN1S, sequence accession JN391180 [99]
Schistosoma haematobium, sequence accession PRJNA78265 [100]
Schistosoma mansoni, sequence accession ERP00038 [101]
Stenopirates sp., sequence accession JN100019 [102]
 T7-Like Virus, sequence accession JN651747 [103]
 Vibrio harveyi siphophage VHS1, sequence accession JF713456 [104]
 Tyrolean ice man, sequence accession ERP001144 [105]

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