

Cliff Han

List of Publications by Year in descending order

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#	ARTICLE	IF	CITATIONS
1	High-Quality Draft Genome Sequences of Four Lignocellulose-Degrading Bacteria Isolated from Puerto Rican Forest Soil: <i>Gordonia</i> sp., <i>Paenibacillus</i> sp., <i>Variovorax</i> sp., and <i>Vogesella</i> sp. <i>Genome Announcements</i> , 2017, 5, .	0.8	18
2	Complete genome sequence of the haloalkaliphilic, obligately chemolithoautotrophic thiosulfate and sulfide-oxidizing β -proteobacterium <i>Thioalkalimicrobium cyclicum</i> type strain ALM 1 (DSM 14477T). <i>Standards in Genomic Sciences</i> , 2016, 11, 38.	1.5	6
3	A specific hygiene hypothesis. <i>Medical Hypotheses</i> , 2016, 93, 146-149.	0.8	2
4	Complete genome sequence of the phenanthrene-degrading soil bacterium <i>Delftia acidovorans</i> Cs1-4. <i>Standards in Genomic Sciences</i> , 2015, 10, 55.	1.5	43
5	Draft Genome Sequence of <i>Thauera</i> sp. Strain SWB20, Isolated from a Singapore Wastewater Treatment Facility Using Gel Microdroplets. <i>Genome Announcements</i> , 2015, 3, .	0.8	5
6	Genome sequence of <i>Bradyrhizobium</i> sp. WSM1253; a microsymbiont of <i>Ornithopus compressus</i> from the Greek Island of Sifnos. <i>Standards in Genomic Sciences</i> , 2015, 10, 113.	1.5	3
7	Complete Genome Sequence of <i>Candidatus Liberibacter africanus</i> , a Bacterium Associated with Citrus Huanglongbing. <i>Genome Announcements</i> , 2015, 3, .	0.8	37
8	High-Quality draft genome sequence of the Lotus spp. microsymbiont <i>Mesorhizobium loti</i> strain CJ3Sym. <i>Standards in Genomic Sciences</i> , 2015, 10, 54.	1.5	2
9	Genome sequence of the Lotus spp. microsymbiont <i>Mesorhizobium loti</i> strain R7A. <i>Standards in Genomic Sciences</i> , 2014, 9, 6.	1.5	22
10	Genome sequence of the <i>Thermotoga thermarum</i> type strain (LA3T) from an African solfataric spring. <i>Standards in Genomic Sciences</i> , 2014, 9, 1105-1117.	1.5	7
11	Genome sequence of the mud-dwelling archaeon <i>Methanoplanus limicola</i> type strain (DSM 2279T), reclassification of <i>Methanoplanus petrolearius</i> as <i>Methanolacinia petrolearia</i> and emended descriptions of the genera <i>Methanoplanus</i> and <i>Methanolacinia</i> . <i>Standards in Genomic Sciences</i> , 2014, 9, 1076-1088.	1.5	22
12	Complete genome sequence of <i>Anabaena variabilis</i> ATCC 29413. <i>Standards in Genomic Sciences</i> , 2014, 9, 562-573.	1.5	43
13	Genome sequence of <i>Microvirga lupini</i> strain LUT6T, a novel <i>Lupinus</i> alphaproteobacterial microsymbiont from Texas. <i>Standards in Genomic Sciences</i> , 2014, 9, 1159-1167.	1.5	6
14	Genome Sequence of <i>Streptomyces viridosporus</i> Strain T7A ATCC 39115, a Lignin-Degrading Actinomycete. <i>Genome Announcements</i> , 2013, 1, .	0.8	20
15	Complete genome sequence of <i>Arthrobacter</i> sp. strain FB24. <i>Standards in Genomic Sciences</i> , 2013, 9, 106-116.	1.5	17
16	Genome sequence of the <i>Trifolium rueppellianum</i> -nodulating <i>Rhizobium leguminosarum</i> bv. <i>trifolii</i> strain WSM2012.. <i>Standards in Genomic Sciences</i> , 2013, 9, 283-293.	1.5	3
17	Genome sequence of the <i>Listia angolensis</i> microsymbiont <i>Microvirga lotononidis</i> strain WSM3557T. <i>Standards in Genomic Sciences</i> , 2013, 9, 540-550.	1.5	7
18	High-quality-draft genome sequence of the yellow-pigmented flavobacterium <i>Joostella marina</i> type strain (En5T). <i>Standards in Genomic Sciences</i> , 2013, 8, 37-46.	1.5	11

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19	Thermus oshimai JL-2 and T. thermophilus JL-18 genome analysis illuminates pathways for carbon, nitrogen, and sulfur cycling. Standards in Genomic Sciences, 2013, 7, 449-468.	1.5	31
20	Complete genome sequence of Dehalobacter restrictus PER-K23T. Standards in Genomic Sciences, 2013, 8, 375-388.	1.5	44
21	Permanent draft genome sequences of the symbiotic nitrogen fixing Ensifer meliloti strains BO21CC and AK58. Standards in Genomic Sciences, 2013, 9, 352-333.	1.5	7
22	Non-contiguous finished genome sequence of plant-growth promoting Serratia proteamaculans S4. Standards in Genomic Sciences, 2013, 8, 441-449.	1.5	26
23	Genome sequence of the chemoheterotrophic soil bacterium Saccharomonospora cyanea type strain (NA-134T). Standards in Genomic Sciences, 2013, 9, 28-41.	1.5	4
24	Genome sequence of the moderately thermophilic sulfur-reducing bacterium Thermanaerovibrio velox type strain (Z-9701T) and emended description of the genus Thermanaerovibrio. Standards in Genomic Sciences, 2013, 9, 57-70.	1.5	8
25	Genome sequence of Frateuria aurantia type strain (KondÅ´ 67T), a xanthomonade isolated from Lilium auratum Lindl.. Standards in Genomic Sciences, 2013, 9, 83-92.	1.5	3
26	Genome sequence of the clover-nodulating Rhizobium leguminosarum bv. trifolii strain TA1. Standards in Genomic Sciences, 2013, 9, 243-253.	1.5	10
27	Genome sequence of the Ornithopus/Lupinus-nodulating Bradyrhizobium sp. strain WSM471. Standards in Genomic Sciences, 2013, 9, 254-263.	1.5	0
28	Genome sequence of the South American clover-nodulating Rhizobium leguminosarum bv. trifolii strain WSM597. Standards in Genomic Sciences, 2013, 9, 264-272.	1.5	4
29	Genome sequence of the lupin-nodulating Bradyrhizobium sp. strain WSM1417. Standards in Genomic Sciences, 2013, 9, 273-282.	1.5	3
30	Genome sequence of Ensifer arboris strain LMG 14919T; a microsymbiont of the legume Prosopis chilensis growing in Kosti, Sudan. Standards in Genomic Sciences, 2013, 9, 473-483.	1.5	6
31	Genome sequence of Rhizobium leguminosarum bv trifolii strain WSM1689, the microsymbiont of the one flowered clover Trifolium uniflorum. Standards in Genomic Sciences, 2013, 9, 527-539.	1.5	27
32	Complete genome sequence of Mesorhizobium australicum type strain (WSM2073T). Standards in Genomic Sciences, 2013, 9, 410-419.	1.5	11
33	Genome sequence of the thermophilic fresh-water bacterium Spirochaeta caldaria type strain (H11), reclassification of Spirochaeta caldaria, Spirochaeta stenostrepta, and Spirochaeta zuelzeriae in the genus Treponema as Treponema caldaria comb. nov., Treponema stenostrepta comb. nov., and Treponema zuelzeriae comb. nov., and emendation of the genus Treponema. Standards in Genomic Sciences, 2013, 8, 88-105.	1.5	44
34	Complete genome sequence of Enterobacter sp. IIT-BT 08: A potential microbial strain for high rate hydrogen production. Standards in Genomic Sciences, 2013, 9, 359-369.	1.5	16
35	Genome sequence of the Lebeckia ambigua-nodulating ÆBurkholderia sprentiaeÆ-strain WSM5005T. Standards in Genomic Sciences, 2013, 9, 385-394.	1.5	9
36	Complete genome sequence of Mesorhizobium australicum type strain (WSM2073T). Standards in Genomic Sciences, 2013, 9, 1-15.	1.5	1

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37	Genome sequence of the moderately thermophilic, amino-acid-degrading and sulfur-reducing bacterium <i>Thermovirga lienii</i> type strain (Cas60314T). Standards in Genomic Sciences, 2012, 6, 230-239.	1.5	26
38	Genome sequence of the orange-pigmented seawater bacterium <i>Owenweeksia hongkongensis</i> type strain (UST20020801T). Standards in Genomic Sciences, 2012, 7, 120-130.	1.5	11
39	Complete genome sequence of the moderately thermophilic mineral-sulfide-oxidizing firmicute <i>Sulfobacillus acidophilus</i> type strain (NALT). Standards in Genomic Sciences, 2012, 6, 293-303.	1.5	23
40	Complete genome sequence of the termite hindgut bacterium <i>Spirochaeta coccoides</i> type strain (SPN1T), reclassification in the genus <i>Sphaerochaeta</i> as <i>Sphaerochaeta coccoides</i> comb. nov. and emendations of the family Spirochaetaceae and the genus <i>Sphaerochaeta</i> . Standards in Genomic Sciences, 2012, 6, 194-209.	1.5	58
41	Permanent draft genome sequence of the gliding predator <i>Saprospira grandis</i> strain Sa g1 (= HR1). Standards in Genomic Sciences, 2012, 6, 210-219.	1.5	3
42	Genome sequence of the soil bacterium <i>Saccharomonospora azurea</i> type strain (NA-128T). Standards in Genomic Sciences, 2012, 6, 220-229.	1.5	13
43	Genome sequence of the ocean sediment bacterium <i>Saccharomonospora marina</i> type strain (XMU15T). Standards in Genomic Sciences, 2012, 6, 265-275.	1.5	5
44	Complete genome sequence of the thermophilic sulfate-reducing ocean bacterium <i>Thermodesulfator indicus</i> type strain (CIR29812T). Standards in Genomic Sciences, 2012, 6, 155-164.	1.5	14
45	Complete genome sequence of <i>Thauera aminoaromatica</i> strain MZ1T. Standards in Genomic Sciences, 2012, 6, 325-335.	1.5	53
46	Complete genome sequence of <i>Serratia plymuthica</i> strain AS12. Standards in Genomic Sciences, 2012, 6, 165-173.	1.5	19
47	Genome sequence of the homoacetogenic bacterium <i>Holophaga foetida</i> type strain (TMBS4T). Standards in Genomic Sciences, 2012, 6, 174-184.	1.5	26
48	Complete genome sequence of the orange-red pigmented, radioresistant <i>Deinococcus proteolyticus</i> type strain (MRPT). Standards in Genomic Sciences, 2012, 6, 240-250.	1.5	14
49	Complete genome sequence of the plant-associated <i>Serratia plymuthica</i> strain AS13. Standards in Genomic Sciences, 2012, 7, 22-30.	1.5	22
50	Genome sequence of the flexirubin-pigmented soil bacterium <i>Niabella soli</i> type strain (JS13-8T). Standards in Genomic Sciences, 2012, 7, 210-220.	1.5	2
51	Genome sequence of the Antarctic rhodopsins-containing flavobacterium <i>Gillisia limnaea</i> type strain (R-8282T). Standards in Genomic Sciences, 2012, 7, 107-119.	1.5	16
52	Complete genome sequence of the sulfate-reducing firmicute <i>Desulfotomaculum ruminis</i> type strain (DLT). Standards in Genomic Sciences, 2012, 7, 304-319.	1.5	22
53	Complete genome sequence of <i>Polynucleobacter necessarius</i> subsp. <i>asymbioticus</i> type strain (QLW-P1DMWA-1T). Standards in Genomic Sciences, 2012, 6, 1-14.	1.5	30
54	Complete genome sequence of <i>Halopiger xanaduensis</i> type strain (SH-6T). Standards in Genomic Sciences, 2012, 6, 31-42.	1.5	9

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55	Complete genome sequence of the facultatively chemolithoautotrophic and methylotrophic alpha Proteobacterium <i>Starkeya novella</i> type strain (ATCC 8093T). <i>Standards in Genomic Sciences</i> , 2012, 7, 44-58.	1.5	16
56	Complete genome sequence of the sulfur compounds oxidizing chemolithoautotroph <i>Sulfuricurvum kujiense</i> type strain (YK-1T). <i>Standards in Genomic Sciences</i> , 2012, 6, 94-103.	1.5	37
57	Complete genome sequence of the rapeseed plant-growth promoting <i>Serratia plymuthica</i> strain AS9. <i>Standards in Genomic Sciences</i> , 2012, 6, 54-62.	1.5	27
58	Complete genome sequence of the facultatively chemolithoautotrophic and methylotrophic alpha Proteobacterium <i>Starkeya novella</i> type strain (ATCC 8093T). <i>Standards in Genomic Sciences</i> , 2012, 7, 44-58.	1.5	19
59	Comparative genome sequence analysis underscores mycoparasitism as the ancestral life style of <i>Trichoderma</i> . <i>Genome Biology</i> , 2011, 12, R40.	3.8	594
60	Complete genome sequence of the filamentous gliding predatory bacterium <i>Herpetosiphon aurantiacus</i> type strain (114-95T). <i>Standards in Genomic Sciences</i> , 2011, 5, 356-370.	1.5	47
61	Complete genome sequence of <i>Leadbetterella byssophila</i> type strain (4M15T). <i>Standards in Genomic Sciences</i> , 2011, 4, 2-12.	1.5	22
62	Complete genome sequence of <i>Hydrogenobacter thermophilus</i> type strain (TK-6T). <i>Standards in Genomic Sciences</i> , 2011, 4, 131-143.	1.5	12
63	Complete genome sequence of <i>Paludibacter propionigenes</i> type strain (WB4T). <i>Standards in Genomic Sciences</i> , 2011, 4, 36-44.	1.5	30
64	Complete genome sequence of <i>Bacteroides helcogenes</i> type strain (P 36-108T). <i>Standards in Genomic Sciences</i> , 2011, 4, 45-53.	1.5	14
65	Complete genome sequence of <i>Weeksella virosa</i> type strain (9751T). <i>Standards in Genomic Sciences</i> , 2011, 4, 81-90.	1.5	15
66	Complete genome sequence of <i>Desulfobulbus propionicus</i> type strain (1pr3T). <i>Standards in Genomic Sciences</i> , 2011, 4, 100-110.	1.5	51
67	Complete genome sequence of <i>Marivirga tractuosa</i> type strain (H-43T). <i>Standards in Genomic Sciences</i> , 2011, 4, 154-162.	1.5	18
68	Complete genome sequence of <i>Desulfurococcus mucosus</i> type strain (O7/1T). <i>Standards in Genomic Sciences</i> , 2011, 4, 173-182.	1.5	10
69	Complete genome sequence of <i>Cellulophaga lytica</i> type strain (LIM-21T). <i>Standards in Genomic Sciences</i> , 2011, 4, 221-232.	1.5	33
70	Non-contiguous finished genome sequence of <i>Bacteroides coprosuis</i> type strain (PC139T). <i>Standards in Genomic Sciences</i> , 2011, 4, 233-243.	1.5	9
71	Complete genome sequence of <i>Rhodospirillum rubrum</i> type strain (S1T). <i>Standards in Genomic Sciences</i> , 2011, 4, 293-302.	1.5	44
72	Complete genome sequence of the extremely halophilic <i>Halanaerobium praevalens</i> type strain (GSLT). <i>Standards in Genomic Sciences</i> , 2011, 4, 312-321.	1.5	36

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73	Complete genome sequence of <i>Nitratifactor salsuginis</i> type strain (E9I37-1T). <i>Standards in Genomic Sciences</i> , 2011, 4, 322-330.	1.5	13
74	Complete genome sequence of <i>Mahella australiensis</i> type strain (50-1 BONT). <i>Standards in Genomic Sciences</i> , 2011, 4, 331-341.	1.5	7
75	Complete genome sequence of <i>Treponema succinifaciens</i> type strain (6091T). <i>Standards in Genomic Sciences</i> , 2011, 4, 361-370.	1.5	41
76	Complete genome sequence of <i>Syntrophobotulus glycolicus</i> type strain (FIGlyRT). <i>Standards in Genomic Sciences</i> , 2011, 4, 371-380.	1.5	11
77	Complete genome sequence of the hyperthermophilic chemolithoautotroph <i>Pyrolobus fumarii</i> type strain (1AT). <i>Standards in Genomic Sciences</i> , 2011, 4, 381-392.	1.5	13
78	Complete genome sequence of <i>Staphylothermus hellenicus</i> P8T. <i>Standards in Genomic Sciences</i> , 2011, 5, 12-20.	1.5	7
79	Complete genome sequence of the acetate-degrading sulfate reducer <i>Desulfobacca acetoxidans</i> type strain (ASRB2T). <i>Standards in Genomic Sciences</i> , 2011, 4, 393-401.	1.5	25
80	Complete genome sequence of the thermophilic, hydrogen-oxidizing <i>Bacillus tusciae</i> type strain (T2T) and reclassification in the new genus, <i>Kyrpidia</i> gen. nov. as <i>Kyrpidia tusciae</i> comb. nov. and emendation of the family Alicyclobacillaceae da Costa and Rainey, 2010.. <i>Standards in Genomic Sciences</i> , 2011, 5, 121-134.	1.5	51
81	Complete genome sequence of the gliding, heparinolytic <i>Pedobacter saltans</i> type strain (113T). <i>Standards in Genomic Sciences</i> , 2011, 5, 30-40.	1.5	16
82	Non-contiguous finished genome sequence of the opportunistic oral pathogen <i>Prevotella multisaccharivorax</i> type strain (PPPA20T). <i>Standards in Genomic Sciences</i> , 2011, 5, 41-49.	1.5	7
83	Complete genome sequence of <i>Tolomonas auensis</i> type strain (TA 4T). <i>Standards in Genomic Sciences</i> , 2011, 5, 112-120.	1.5	6
84	Complete genome sequence of <i>Ferroplasma placidus</i> AEDII12DO. <i>Standards in Genomic Sciences</i> , 2011, 5, 50-60.	1.5	32
85	Complete genome sequence of the halophilic and highly halotolerant <i>Chromohalobacter salexigens</i> type strain (1H11T). <i>Standards in Genomic Sciences</i> , 2011, 5, 379-388.	1.5	35
86	Complete genome sequence of <i>Allochromatium vinosum</i> DSM 180T. <i>Standards in Genomic Sciences</i> , 2011, 5, 311-330.	1.5	65
87	Genome sequence of the filamentous, gliding <i>Thiothrix nivea</i> neotype strain (JP2T). <i>Standards in Genomic Sciences</i> , 2011, 5, 398-406.	1.5	22
88	Complete genome sequence of the thermophilic sulfur-reducer <i>Desulfurobacterium thermolithotrophum</i> type strain (BSAT) from a deep-sea hydrothermal vent. <i>Standards in Genomic Sciences</i> , 2011, 5, 407-415.	1.5	11
89	Complete genome sequence of <i>Calditerrivibrio nitroreducens</i> type strain (Yu37-1T). <i>Standards in Genomic Sciences</i> , 2011, 4, 54-62.	1.5	10
90	Complete genome sequence of <i>Truepera radiovictrix</i> type strain (RQ-24T). <i>Standards in Genomic Sciences</i> , 2011, 4, 91-99.	1.5	46

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91	Complete genome sequence of <i>Bacteroides salanitronis</i> type strain (BL78T). Standards in Genomic Sciences, 2011, 4, 191-199.	1.5	11
92	Complete genome sequence of <i>Odoribacter splanchnicus</i> type strain (1651/6T). Standards in Genomic Sciences, 2011, 4, 200-209.	1.5	96
93	Complete genome sequence of <i>Oceanithermus profundus</i> type strain (506T). Standards in Genomic Sciences, 2011, 4, 210-220.	1.5	4
94	Complete genome sequence of <i>Tsukamurella paurometabola</i> type strain (no. 33T). Standards in Genomic Sciences, 2011, 4, 342-351.	1.5	10
95	Non-contiguous finished genome sequence and contextual data of the filamentous soil bacterium <i>Ktedonobacter racemifer</i> type strain (SOSP1-21T). Standards in Genomic Sciences, 2011, 5, 97-111.	1.5	115
96	Complete genome sequence of <i>Isosphaera pallida</i> type strain (IS1BT). Standards in Genomic Sciences, 2011, 4, 63-71.	1.5	46
97	Complete genome sequence of <i>Cellulophaga algicola</i> type strain (IC166T). Standards in Genomic Sciences, 2011, 4, 72-80.	1.5	28
98	Complete genome sequence of <i>Riemerella anatipestifer</i> type strain (ATCC 11845T). Standards in Genomic Sciences, 2011, 4, 144-153.	1.5	33
99	Complete genome sequence of the thermophilic sulfur-reducer <i>Hippea maritima</i> type strain (MH2T). Standards in Genomic Sciences, 2011, 4, 303-311.	1.5	8
100	Complete genome sequence of <i>Haliscomenobacter hydrossis</i> type strain (OT). Standards in Genomic Sciences, 2011, 4, 352-360.	1.5	26
101	Complete genome sequence of the gliding freshwater bacterium <i>Fluviicola taffensis</i> type strain (RW262T). Standards in Genomic Sciences, 2011, 5, 21-29.	1.5	23
102	Genome sequence of the moderately thermophilic halophile <i>Flexistipes sinusarabici</i> strain (MAS10T). Standards in Genomic Sciences, 2011, 5, 86-96.	1.5	12
103	Complete genome sequence of <i>Deinococcus maricopensis</i> type strain (LB-34T). Standards in Genomic Sciences, 2011, 4, 163-172.	1.5	14
104	Complete genome sequence of <i>Thermomonospora curvata</i> type strain (B9T). Standards in Genomic Sciences, 2011, 4, 13-22.	1.5	35
105	Complete genome sequence of <i>Riemerella anatipestifer</i> type strain (ATCC 11845). Standards in Genomic Sciences, 2011, 4, 144-53.	1.5	20
106	Complete genome sequence of <i>Syntrophobotulus glycolicus</i> type strain (FIGlyR). Standards in Genomic Sciences, 2011, 4, 371-80.	1.5	4
107	Complete genome sequence of <i>Ferrimonas balearica</i> type strain (PATT). Standards in Genomic Sciences, 2010, 3, 174-182.	1.5	14
108	Non-contiguous finished genome sequence of <i>Aminomonas paucivorans</i> type strain (GLU-3T). Standards in Genomic Sciences, 2010, 3, 285-293.	1.5	9

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109	Complete genome sequence of <i>Ilyobacter polytropus</i> type strain (CuHbu1T). Standards in Genomic Sciences, 2010, 3, 304-314.	1.5	10
110	Complete genome sequence of <i>Nocardioopsis dassonvillei</i> type strain (IMRU 509T). Standards in Genomic Sciences, 2010, 3, 325-336.	1.5	32
111	Complete genome sequence of <i>Veillonella parvula</i> type strain (Te3T). Standards in Genomic Sciences, 2010, 2, 57-65.	1.5	44
112	Complete genome sequence of <i>Thermocrinis albus</i> type strain (HI 11/12T). Standards in Genomic Sciences, 2010, 2, 194-202.	1.5	18
113	Complete genome sequence of <i>Meiothermus silvanus</i> type strain (VI-R2T). Standards in Genomic Sciences, 2010, 3, 37-46.	1.5	19
114	Complete genome sequence of <i>Olsenella uli</i> type strain (VPI D76D-27CT). Standards in Genomic Sciences, 2010, 3, 76-84.	1.5	43
115	Complete genome sequence of <i>Acidaminococcus fermentans</i> type strain (VR4T). Standards in Genomic Sciences, 2010, 3, 1-14.	1.5	31
116	Complete genome sequence of <i>Meiothermus ruber</i> type strain (21T). Standards in Genomic Sciences, 2010, 3, 26-36.	1.5	33
117	Complete genome sequence of <i>Acetohalobium arabaticum</i> type strain (Z-7288T). Standards in Genomic Sciences, 2010, 3, 57-65.	1.5	24
118	Complete genome sequence of <i>Ignisphaera aggregans</i> type strain (AQ1.S1T). Standards in Genomic Sciences, 2010, 3, 66-75.	1.5	15
119	Complete genome sequence of <i>Arcanobacterium haemolyticum</i> type strain (11018T). Standards in Genomic Sciences, 2010, 3, 126-135.	1.5	8
120	Complete genome sequence of <i>Thermosediminibacter oceani</i> type strain (JW/IW-1228PT). Standards in Genomic Sciences, 2010, 3, 108-116.	1.5	12
121	Complete genome sequence of <i>Spirochaeta smaragdinae</i> type strain (SEBR 4228T). Standards in Genomic Sciences, 2010, 3, 1-9.	1.5	18
122	Complete genome sequence of <i>Thermobaculum terrenum</i> ™ type strain (YNP1T). Standards in Genomic Sciences, 2010, 3, 153-162.	1.5	11
123	Complete genome sequence of <i>Syntrophothermus lipocalidus</i> type strain (TGB-C1T). Standards in Genomic Sciences, 2010, 3, 268-275.	1.5	13
124	Complete genome sequence of <i>Desulfarculus baarsii</i> type strain (2st14T). Standards in Genomic Sciences, 2010, 3, 276-284.	1.5	37
125	Complete genome sequence of <i>Intrasporangium calvum</i> type strain (7 KIPT). Standards in Genomic Sciences, 2010, 3, 294-303.	1.5	7
126	Complete genome sequence of <i>Methanothermus fervidus</i> type strain (V24ST). Standards in Genomic Sciences, 2010, 3, 315-324.	1.5	17

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127	Complete genome sequence of <i>Thermaerobacter marianensis</i> type strain (7p75aT). Standards in Genomic Sciences, 2010, 3, 337-345.	1.5	10
128	Complete genome sequence of <i>Xylanimonas cellulositytica</i> type strain (XIL07T). Standards in Genomic Sciences, 2010, 2, 1-8.	1.5	10
129	Complete genome sequence of <i>Alicyclobacillus acidocaldarius</i> type strain (104-IAT). Standards in Genomic Sciences, 2010, 2, 9-18.	1.5	24
130	Complete genome sequence of <i>Streptosporangium roseum</i> type strain (NI 9100T). Standards in Genomic Sciences, 2010, 2, 29-37.	1.5	27
131	Complete genome sequence of <i>Sulfurospirillum deleyianum</i> type strain (5175T). Standards in Genomic Sciences, 2010, 2, 149-157.	1.5	29
132	Complete genome sequence of <i>Haloterrigena turkmenica</i> type strain (4kT). Standards in Genomic Sciences, 2010, 2, 107-116.	1.5	32
133	Complete genome sequence of <i>Segniliparus rotundus</i> type strain (CDC 1076T). Standards in Genomic Sciences, 2010, 2, 203-211.	1.5	10
134	Complete genome sequence of <i>Sebaldella termitidis</i> type strain (NCTC 11300T). Standards in Genomic Sciences, 2010, 2, 220-227.	1.5	34
135	Complete genome sequence of <i>Thermosphaera aggregans</i> type strain (M11TLT). Standards in Genomic Sciences, 2010, 2, 245-259.	1.5	14
136	Complete genome sequence of <i>Brachyspira murdochii</i> type strain (56-150T). Standards in Genomic Sciences, 2010, 2, 260-269.	1.5	20
137	Complete genome sequence of <i>Aminobacterium colombiense</i> type strain (ALA-1T). Standards in Genomic Sciences, 2010, 2, 280-289.	1.5	32
138	Complete genome sequence of <i>Arcobacter nitrofigilis</i> type strain (CIT). Standards in Genomic Sciences, 2010, 2, 300-308.	1.5	40
139	Complete genome sequence of <i>Coralimargarita akajimensis</i> type strain (04OKA010-24T). Standards in Genomic Sciences, 2010, 2, 290-299.	1.5	28
140	Complete genome sequence of <i>Thermobispora bispora</i> type strain (R51T). Standards in Genomic Sciences, 2010, 2, 318-326.	1.5	23
141	Complete genome sequence of <i>Desulfohalobium retbaense</i> type strain (HR100T). Standards in Genomic Sciences, 2010, 2, 38-48.	1.5	22
142	Complete genome sequence of <i>Archaeoglobus profundus</i> type strain (AV18T). Standards in Genomic Sciences, 2010, 2, 327-346.	1.5	26
143	Complete genome sequence of <i>Denitrovibrio acetiphilus</i> type strain (N2460T). Standards in Genomic Sciences, 2010, 2, 270-279.	1.5	16
144	Complete genome sequence of <i>Gordonia bronchialis</i> type strain (3410T). Standards in Genomic Sciences, 2010, 2, 19-28.	1.5	26

#	ARTICLE	IF	CITATIONS
145	The genome of <i>Syntrophomonas wolfei</i> : new insights into syntrophic metabolism and biohydrogen production. <i>Environmental Microbiology</i> , 2010, 12, 2289-2301.	1.8	158
146	Complete genome sequence of <i>Rhizobium leguminosarum</i> bv. <i>trifolii</i> strain WSM1325, an effective microsymbiont of annual Mediterranean clovers.. <i>Standards in Genomic Sciences</i> , 2010, 2, 347-356.	1.5	53
147	Complete genome sequence of <i>Sulfurimonas autotrophica</i> type strain (OK10T). <i>Standards in Genomic Sciences</i> , 2010, 3, 194-202.	1.5	37
148	Complete genome sequence of <i>Methanoplanus petrolearius</i> type strain (SEBR 4847T). <i>Standards in Genomic Sciences</i> , 2010, 3, 203-211.	1.5	14
149	Complete genome sequence of <i>Capnocytophaga ochracea</i> type strain (VPI 2845T). <i>Standards in Genomic Sciences</i> , 2009, 1, 101-109.	1.5	14
150	A phylogeny-driven genomic encyclopaedia of Bacteria and Archaea. <i>Nature</i> , 2009, 462, 1056-1060.	13.7	924
151	Complete genome sequence of <i>Halorhabdus utahensis</i> type strain (AX-2T). <i>Standards in Genomic Sciences</i> , 2009, 1, 218-225.	1.5	22
152	Complete genome sequence of <i>Methanoculleus marisnigri</i> Romesser et al. 1981 type strain JR1. <i>Standards in Genomic Sciences</i> , 2009, 1, 189-196.	1.5	34
153	Complete genome sequence of <i>Beutenbergia cavernae</i> type strain (HKI 0122T). <i>Standards in Genomic Sciences</i> , 2009, 1, 21-28.	1.5	12
154	Complete genome sequence of <i>Cryptobacterium curtum</i> type strain (12-3T). <i>Standards in Genomic Sciences</i> , 2009, 1, 93-100.	1.5	17
155	Complete genome sequence of <i>Desulfomicrobium baculatum</i> type strain (XT). <i>Standards in Genomic Sciences</i> , 2009, 1, 29-37.	1.5	36
156	Complete genome sequence of <i>Sanguibacter keddieii</i> type strain (ST-74T). <i>Standards in Genomic Sciences</i> , 2009, 1, 110-118.	1.5	16
157	Complete genome sequence of <i>Catenulispora acidiphila</i> type strain (ID 139908T). <i>Standards in Genomic Sciences</i> , 2009, 1, 119-125.	1.5	24
158	Complete genome sequence of <i>Leptotrichia buccalis</i> type strain (C-1013-bT). <i>Standards in Genomic Sciences</i> , 2009, 1, 126-132.	1.5	24
159	Complete genome sequence of <i>Saccharomonospora viridis</i> type strain (P101T). <i>Standards in Genomic Sciences</i> , 2009, 1, 141-149.	1.5	24
160	Complete genome sequence of <i>Actinosynnema mirum</i> type strain (101T). <i>Standards in Genomic Sciences</i> , 2009, 1, 46-53.	1.5	38
161	Complete genome sequence of <i>Pedobacter heparinus</i> type strain (HIM 762-3T). <i>Standards in Genomic Sciences</i> , 2009, 1, 54-62.	1.5	25
162	Complete genome sequence of <i>Anaerococcus prevotii</i> type strain (PC1T). <i>Standards in Genomic Sciences</i> , 2009, 1, 159-165.	1.5	25

#	ARTICLE	IF	CITATIONS
163	Complete genome sequence of <i>Atopobium parvulum</i> type strain (IPP 1246T). <i>Standards in Genomic Sciences</i> , 2009, 1, 166-173.	1.5	30
164	Complete genome sequence of <i>Eggerthella lenta</i> type strain (VPI 0255T). <i>Standards in Genomic Sciences</i> , 2009, 1, 174-182.	1.5	37
165	Complete genome sequence of <i>Kangiella koreensis</i> type strain (SW-125T). <i>Standards in Genomic Sciences</i> , 2009, 1, 226-233.	1.5	28
166	Complete genome sequence of <i>Jonesia denitrificans</i> type strain (Prevot 55134T). <i>Standards in Genomic Sciences</i> , 2009, 1, 262-269.	1.5	14
167	Complete genome sequence of <i>Halomicrobium mukohataei</i> type strain (arg-2T). <i>Standards in Genomic Sciences</i> , 2009, 1, 270-277.	1.5	31
168	Complete genome sequence of <i>Rhodothermus marinus</i> type strain (R-10T). <i>Standards in Genomic Sciences</i> , 2009, 1, 283-290.	1.5	45
169	Complete genome sequence of <i>Streptobacillus moniliformis</i> type strain (9901T). <i>Standards in Genomic Sciences</i> , 2009, 1, 300-307.	1.5	21
170	Complete genome sequence of <i>Pirellula staleyi</i> type strain (ATCC 27377T). <i>Standards in Genomic Sciences</i> , 2009, 1, 308-316.	1.5	34
171	Complete genome sequence of <i>Kytococcus sedentarius</i> type strain (541T). <i>Standards in Genomic Sciences</i> , 2009, 1, 12-20.	1.5	100
172	Complete genome sequence of <i>Dyadobacter fermentans</i> type strain (NS114T). <i>Standards in Genomic Sciences</i> , 2009, 1, 133-140.	1.5	25
173	Complete genome sequence of <i>Slackia heliotrinireducens</i> type strain (RHS 1T). <i>Standards in Genomic Sciences</i> , 2009, 1, 234-241.	1.5	20
174	Complete genome sequence of <i>Desulfotomaculum acetoxidans</i> type strain (5575T). <i>Standards in Genomic Sciences</i> , 2009, 1, 242-253.	1.5	35
175	Complete genome sequence of <i>Stackebrandtia nassauensis</i> type strain (LLR-40K-21T). <i>Standards in Genomic Sciences</i> , 2009, 1, 292-299.	1.5	23
176	Assembling the Marine Metagenome, One Cell at a Time. <i>PLoS ONE</i> , 2009, 4, e5299.	1.1	320
177	Genomic Characterization of Methanomicrobiales Reveals Three Classes of Methanogens. <i>PLoS ONE</i> , 2009, 4, e5797.	1.1	103
178	A 360-kb interchromosomal duplication of the human HYDIN locus. <i>Genomics</i> , 2006, 88, 762-771.	1.3	52
179	The sequence and analysis of duplication-rich human chromosome 16. <i>Nature</i> , 2004, 432, 988-994.	13.7	156