

Miriam L Land

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

313
papers

18,349
citations

53
h-index

126
g-index

314
ext. papers

23,353
ext. citations

6.7
avg, IF

5.87
L-index

#	Paper	IF	Citations
313	Computationally Guided Discovery and Experimental Validation of Indole-3-acetic Acid Synthesis Pathways. <i>ACS Chemical Biology</i> , 2019 , 14, 2867-2875	4.9	5
312	Genome Characterization of Oleaginous <i>Aspergillus oryzae</i> BCC7051: A Potential Fungal-Based Platform for Lipid Production. <i>Current Microbiology</i> , 2018 , 75, 57-70	2.4	24
311	KBase: The United States Department of Energy Systems Biology Knowledgebase. <i>Nature Biotechnology</i> , 2018 , 36, 566-569	44.5	419
310	Exploration of the Biosynthetic Potential of the Microbiome. <i>MSystems</i> , 2018 , 3,	7.6	21
309	Enrichment of Root Endophytic Bacteria from <i>Populus deltoides</i> and Single-Cell-Genomics Analysis. <i>Applied and Environmental Microbiology</i> , 2016 , 82, 5698-708	4.8	40
308	Complete genome sequence of the Antarctic <i>Halorubrum lacusprofundi</i> type strain ACAM 34. <i>Standards in Genomic Sciences</i> , 2016 , 11, 70		15
307	Complete genome sequence of the haloalkaliphilic, obligately chemolithoautotrophic thiosulfate and sulfide-oxidizing β proteobacterium <i>Thioalkalimicrobium cyclicum</i> type strain ALM 1 (DSM 14477(T)). <i>Standards in Genomic Sciences</i> , 2016 , 11, 38		5
306	Complete genome sequence of <i>Methanospirillum hungatei</i> type strain JF1. <i>Standards in Genomic Sciences</i> , 2016 , 11, 2		23
305	Complete genome sequences of <i>Geobacillus</i> sp. WCH70, a thermophilic strain isolated from wood compost. <i>Standards in Genomic Sciences</i> , 2016 , 11, 33		8
304	Diversity of <i>Pseudomonas</i> Genomes, Including <i>Populus</i> -Associated Isolates, as Revealed by Comparative Genome Analysis. <i>Applied and Environmental Microbiology</i> , 2016 , 82, 375-83	4.8	50
303	Application of Long Sequence Reads To Improve Genomes for <i>Clostridium thermocellum</i> AD2, <i>Clostridium thermocellum</i> LQRI, and <i>Pelosinus fermentans</i> R7. <i>Genome Announcements</i> , 2016 , 4,		2
302	Insights from 20 years of bacterial genome sequencing. <i>Functional and Integrative Genomics</i> , 2015 , 15, 141-61	3.8	391
301	Complete Genome Sequence of the Oritutaceae Bacterium Strain TAV5, a Potential Facultative Methylophile of the Wood-Feeding Termite <i>Reticulitermes flavipes</i> . <i>Genome Announcements</i> , 2015 , 3,		15
300	<i>Trichodesmium</i> genome maintains abundant, widespread noncoding DNA in situ, despite oligotrophic lifestyle. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015 , 112, 4251-6	11.5	31
299	Genome-scale resources for <i>Thermoanaerobacterium saccharolyticum</i> . <i>BMC Systems Biology</i> , 2015 , 9, 30	3.5	14
298	Ebolavirus comparative genomics. <i>FEMS Microbiology Reviews</i> , 2015 , 39, 764-78	15.1	34
297	Complete Genome Sequence of <i>Anaeromyxobacter</i> sp. Fw109-5, an Anaerobic, Metal-Reducing Bacterium Isolated from a Contaminated Subsurface Environment. <i>Genome Announcements</i> , 2015 , 3,		10

296	Complete genome sequence of <i>Geobacillus thermoglucosidasius</i> C56-YS93, a novel biomass degrader isolated from obsidian hot spring in Yellowstone National Park. <i>Standards in Genomic Sciences</i> , 2015 , 10, 73		16
295	Complete genome sequences of <i>Geobacillus</i> sp. Y412MC52, a xylan-degrading strain isolated from obsidian hot spring in Yellowstone National Park. <i>Standards in Genomic Sciences</i> , 2015 , 10, 81		9
294	Complete genome sequence of the phenanthrene-degrading soil bacterium <i>Delftia acidovorans</i> Cs1-4. <i>Standards in Genomic Sciences</i> , 2015 , 10, 55		17
293	Genome and Transcriptome of <i>Clostridium phytofermentans</i> , Catalyst for the Direct Conversion of Plant Feedstocks to Fuels. <i>PLoS ONE</i> , 2015 , 10, e0118285	3.7	16
292	High-Quality Draft Genome Sequence of <i>Desulfovibrio carbinoliphilus</i> FW-101-2B, an Organic Acid-Oxidizing Sulfate-Reducing Bacterium Isolated from Uranium(VI)-Contaminated Groundwater. <i>Genome Announcements</i> , 2015 , 3,		1
291	Complete Genome Sequence of <i>Geobacillus</i> strain Y4.1MC1, a Novel CO-Utilizing <i>Geobacillus thermoglucosidasius</i> Strain Isolated from Bath Hot Spring in Yellowstone National Park. <i>Bioenergy Research</i> , 2015 , 8, 1039-1045	3.1	13
290	Complete genome sequence of <i>Planctomyces brasiliensis</i> type strain (DSM 5305(T)), phylogenomic analysis and reclassification of Planctomycetes including the descriptions of <i>Gimesia</i> gen. nov., <i>Planctopirus</i> gen. nov. and <i>Rubinisphaera</i> gen. nov. and emended descriptions of the order Planctomycetales and the family Planctomycetaceae. <i>Standards in Genomic Sciences</i> , 2014 , 9, 10		39
289	Quality scores for 32,000 genomes. <i>Standards in Genomic Sciences</i> , 2014 , 9, 20		33
288	Comparison of single-molecule sequencing and hybrid approaches for finishing the genome of <i>Clostridium autoethanogenum</i> and analysis of CRISPR systems in industrial relevant Clostridia. <i>Biotechnology for Biofuels</i> , 2014 , 7, 40	7.8	105
287	Genome sequence of the <i>Thermotoga thermarum</i> type strain (LA3(T)) from an African solfataric spring. <i>Standards in Genomic Sciences</i> , 2014 , 9, 1105-17		4
286	Genome sequence of the mud-dwelling archaeon <i>Methanoplanus limicola</i> type strain (DSM 2279(T)), 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, 1076-88		10
285	Complete genome sequence of <i>Anabaena variabilis</i> ATCC 29413. <i>Standards in Genomic Sciences</i> , 2014 , 9, 562-73		32
284	Complete Genome sequence of <i>Burkholderia phymatum</i> STM815(T), a broad host range and efficient nitrogen-fixing symbiont of <i>Mimosa</i> species. <i>Standards in Genomic Sciences</i> , 2014 , 9, 763-74		36
283	Evaluation and validation of de novo and hybrid assembly techniques to derive high-quality genome sequences. <i>Bioinformatics</i> , 2014 , 30, 2709-16	7.2	86
282	Complete genome sequence of <i>Granulicella tundricola</i> type strain MP5ACTX9(T), an Acidobacteria from tundra soil. <i>Standards in Genomic Sciences</i> , 2014 , 9, 449-61		12
281	Complete genome sequence of <i>Mesorhizobium ciceri</i> bv. <i>biserrulae</i> type strain (WSM1271(T)). <i>Standards in Genomic Sciences</i> , 2014 , 9, 462-72		16
280	Global transcriptome analysis of <i>Clostridium thermocellum</i> ATCC 27405 during growth on dilute acid pretreated <i>Populus</i> and switchgrass. <i>Biotechnology for Biofuels</i> , 2013 , 6, 179	7.8	53
279	Draft genome sequence of <i>Frankia</i> sp. strain QA3, a nitrogen-fixing actinobacterium isolated from the root nodule of <i>Alnus nitida</i> . <i>Genome Announcements</i> , 2013 , 1, e0010313		35

278	Genome Sequence of the Obligate Gammaproteobacterial Methanotroph <i>Methylobacterium album</i> Strain BG8. <i>Genome Announcements</i> , 2013 , 1, e0017013	16
277	Draft genome sequence of <i>Frankia</i> sp. strain CN3, an atypical, noninfective (Nod-) ineffective (Fix-) isolate from <i>Coriaria nepalensis</i> . <i>Genome Announcements</i> , 2013 , 1, e0008513	46
276	Complete Genome Sequence of the Hyperthermophilic Sulfate-Reducing Bacterium <i>Thermodesulfobacterium geofontis</i> OPF15T. <i>Genome Announcements</i> , 2013 , 1, e0016213	4
275	Genome Sequence of <i>Streptomyces viridosporus</i> Strain T7A ATCC 39115, a Lignin-Degrading Actinomycete. <i>Genome Announcements</i> , 2013 , 1,	15
274	Complete Genome Sequence of <i>Micromonospora</i> Strain L5, a Potential Plant-Growth-Regulating Actinomycete, Originally Isolated from <i>Casuarina equisetifolia</i> Root Nodules. <i>Genome Announcements</i> , 2013 , 1,	14
273	Complete genome sequence of <i>Granulicella mallensis</i> type strain MP5ACTX8(T), an acidobacterium from tundra soil. <i>Standards in Genomic Sciences</i> , 2013 , 9, 71-82	15
272	Complete genome sequence of <i>Coriobacterium glomerans</i> type strain (PW2(T)) from the midgut of <i>Pyrrhocoris apterus</i> L. (red soldier bug). <i>Standards in Genomic Sciences</i> , 2013 , 8, 15-25	5
271	Complete genome sequence of the bile-resistant pigment-producing anaerobe <i>Alistipes finegoldii</i> type strain (AHN2437(T)). <i>Standards in Genomic Sciences</i> , 2013 , 8, 26-36	6
270	Complete genome sequence of the moderate thermophile <i>Anaerobaculum mobile</i> type strain (NGA(T)). <i>Standards in Genomic Sciences</i> , 2013 , 8, 47-57	10
269	Genome sequence of the free-living aerobic spirochete <i>Turneriella parva</i> type strain (H(T)), and emendation of the species <i>Turneriella parva</i> . <i>Standards in Genomic Sciences</i> , 2013 , 8, 228-38	8
268	Complete genome sequence of <i>Halorhodospira halophila</i> SL1. <i>Standards in Genomic Sciences</i> , 2013 , 8, 206-14	13
267	Genome sequence of the chemoheterotrophic soil bacterium <i>Saccharomonospora cyanea</i> type strain (NA-134(T)). <i>Standards in Genomic Sciences</i> , 2013 , 9, 28-41	1
266	Genome sequence of <i>Frateuria aurantia</i> type strain (Kond67(T)), a xanthomonade isolated from <i>Lilium auratum</i> Lindl. <i>Standards in Genomic Sciences</i> , 2013 , 9, 83-92	1
265	Complete genome sequence of <i>Mesorhizobium opportunistum</i> type strain WSM2075(T). <i>Standards in Genomic Sciences</i> , 2013 , 9, 294-303	10
264	Permanent draft genome sequence of <i>Comamonas testosteroni</i> KF-1. <i>Standards in Genomic Sciences</i> , 2013 , 8, 239-54	9
263	Genome sequence of the thermophilic fresh-water bacterium <i>Spirochaeta caldaria</i> type strain (H1(T)), reclassification of <i>Spirochaeta caldaria</i> , <i>Spirochaeta stenostrepta</i> , and <i>Spirochaeta zuelzeri</i> in the genus <i>Treponema</i> as <i>Treponema caldaria</i> comb. nov., <i>Treponema stenostrepta</i> comb. nov., and <i>Treponema zuelzeri</i> comb. nov. <i>Standards in Genomic Sciences</i> , 2013 , 9, 103-11	24
262	Complete genome sequence of the halophilic bacterium <i>Spirochaeta africana</i> type strain (Z-7692(T)) from the alkaline Lake Magadi in the East African Rift. <i>Standards in Genomic Sciences</i> , 2013 , 8, 165-76	2
261	Genome sequence of the phylogenetically isolated spirochete <i>Leptonema illini</i> type strain (3055(T)). <i>Standards in Genomic Sciences</i> , 2013 , 8, 177-87	4

260	Industrial robustness: understanding the mechanism of tolerance for the Populus hydrolysate-tolerant mutant strain of Clostridium thermocellum. <i>PLoS ONE</i> , 2013 , 8, e78829	3.7	19
259	Genome analysis and physiological comparison of Alicyclophilus denitrificans strains BC and K601(T.). <i>PLoS ONE</i> , 2013 , 8, e66971	3.7	26
258	Complete genome sequence of Nitrosomonas sp. Is79, an ammonia oxidizing bacterium adapted to low ammonium concentrations. <i>Standards in Genomic Sciences</i> , 2013 , 7, 469-82		27
257	Draft genome sequences for Clostridium thermocellum wild-type strain YS and derived cellulose adhesion-defective mutant strain AD2. <i>Journal of Bacteriology</i> , 2012 , 194, 3290-1	3.5	22
256	Twenty-one genome sequences from Pseudomonas species and 19 genome sequences from diverse bacteria isolated from the rhizosphere and endosphere of Populus deltoides. <i>Journal of Bacteriology</i> , 2012 , 194, 5991-3	3.5	117
255	Caldicellulosiruptor core and pangenomes reveal determinants for noncellulosomal thermophilic deconstruction of plant biomass. <i>Journal of Bacteriology</i> , 2012 , 194, 4015-28	3.5	81
254	Draft genome sequence for Microbacterium laevaniformans strain OR221, a bacterium tolerant to metals, nitrate, and low pH. <i>Journal of Bacteriology</i> , 2012 , 194, 3279-80	3.5	19
253	The genome of Pelobacter carbinolicus reveals surprising metabolic capabilities and physiological features. <i>BMC Genomics</i> , 2012 , 13, 690	4.5	26
252	Complete genome sequence of the rapeseed plant-growth promoting Serratia plymuthica strain AS9. <i>Standards in Genomic Sciences</i> , 2012 , 6, 54-62		25
251	The fast changing landscape of sequencing technologies and their impact on microbial genome assemblies and annotation. <i>PLoS ONE</i> , 2012 , 7, e48837	3.7	112
250	Complete genome sequences of Desulfosporosinus orientis DSM765T, Desulfosporosinus youngiae DSM17734T, Desulfosporosinus meridiei DSM13257T, and Desulfosporosinus acidiphilus DSM22704T. <i>Journal of Bacteriology</i> , 2012 , 194, 6300-1	3.5	46
249	Complete genome sequence of the thermophilic, piezophilic, heterotrophic bacterium Marinitoga piezophila KA3. <i>Journal of Bacteriology</i> , 2012 , 194, 5974-5	3.5	18
248	High-quality draft genome sequence of the Opitutaceae bacterium strain TAV1, a symbiont of the wood-feeding termite Reticulitermes flavipes. <i>Journal of Bacteriology</i> , 2012 , 194, 2744-5	3.5	18
247	Draft genome sequence of Rhizobium sp. strain PDO1-076, a bacterium isolated from Populus deltoides. <i>Journal of Bacteriology</i> , 2012 , 194, 2383-4	3.5	15
246	Complete genome sequence of Rahnella sp. strain Y9602, a gammaproteobacterium isolate from metal- and radionuclide-contaminated soil. <i>Journal of Bacteriology</i> , 2012 , 194, 2113-4	3.5	9
245	Complete genome sequence of Desulfurococcus fermentans, a hyperthermophilic cellulolytic crenarchaeon isolated from a freshwater hot spring in Kamchatka, Russia. <i>Journal of Bacteriology</i> , 2012 , 194, 5703-4	3.5	11
244	Complete genome sequence of Rahnella aquatilis CIP 78.65. <i>Journal of Bacteriology</i> , 2012 , 194, 3020-1	3.5	16
243	Draft genome sequences for two metal-reducing Pelosinus fermentans strains isolated from a Cr(VI)-contaminated site and for type strain R7. <i>Journal of Bacteriology</i> , 2012 , 194, 5147-8	3.5	17

242	Genome sequence of the mesophilic Thermotogales bacterium <i>Mesotoga prima</i> MesG1.Ag.4.2 reveals the largest Thermotogales genome to date. <i>Genome Biology and Evolution</i> , 2012 , 4, 700-8	3.9	18
241	Complete genome sequences of six strains of the genus <i>Methylobacterium</i> . <i>Journal of Bacteriology</i> , 2012 , 194, 4746-8	3.5	70
240	The whole genome sequence of <i>Sphingobium chlorophenicum</i> L-1: insights into the evolution of the pentachlorophenol degradation pathway. <i>Genome Biology and Evolution</i> , 2012 , 4, 184-98	3.9	68
239	Genome sequence of the moderately thermophilic, amino-acid-degrading and sulfur-reducing bacterium <i>Thermovirga lienii</i> type strain (Cas60314(T)). <i>Standards in Genomic Sciences</i> , 2012 , 6, 230-9		21
238	Genome sequence of <i>Amycolatopsis</i> sp. strain ATCC 39116, a plant biomass-degrading actinomycete. <i>Journal of Bacteriology</i> , 2012 , 194, 2396-7	3.5	31
237	Complete genome sequence of the moderately thermophilic mineral-sulfide-oxidizing firmicute <i>Sulfobacillus acidophilus</i> type strain (NAL(T)). <i>Standards in Genomic Sciences</i> , 2012 , 6, 1-13		20
236	Complete genome sequence of the termite hindgut bacterium <i>Spirochaeta coccoides</i> type strain (SPN1(T)), 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> . <i>Standards in Genomic Sciences</i> , 2012 , 6, 194-209		46
235	Permanent draft genome sequence of the gliding predator <i>Saprospira grandis</i> strain Sa g1 (= HR1). <i>Standards in Genomic Sciences</i> , 2012 , 6, 210-9		1
234	Complete genome sequence of <i>Paenibacillus</i> sp. strain JDR-2. <i>Standards in Genomic Sciences</i> , 2012 , 6, 1-10		12
233	Complete genome sequence of the melanogenic marine bacterium <i>Marinomonas mediterranea</i> type strain (MMB-1(T)). <i>Standards in Genomic Sciences</i> , 2012 , 6, 63-73		18
232	Genome sequence of the soil bacterium <i>Saccharomonospora azurea</i> type strain (NA-128(T)). <i>Standards in Genomic Sciences</i> , 2012 , 6, 220-9		10
231	Complete genome sequence of the facultatively anaerobic, appendaged bacterium <i>Muricauda ruestringensis</i> type strain (B1(T)). <i>Standards in Genomic Sciences</i> , 2012 , 6, 185-93		8
230	Conceptualizing a Genomics Software Institute (GSI). <i>Standards in Genomic Sciences</i> , 2012 , 6, 136-44		1
229	Complete genome sequence of <i>Dehalogenimonas lykanthroporepellens</i> type strain (BL-DC-9(T)) and comparison to "Dehalococoides" strains. <i>Standards in Genomic Sciences</i> , 2012 , 6, 251-64		45
228	Complete genome sequence of <i>Polynucleobacter necessarius</i> subsp. <i>asymbioticus</i> type strain (QLW-P1DMWA-1(T)). <i>Standards in Genomic Sciences</i> , 2012 , 6, 74-83		27
227	Complete genome sequence of the aerobic, heterotroph <i>Marinithermus hydrothermalis</i> type strain (T1(T)) from a deep-sea hydrothermal vent chimney. <i>Standards in Genomic Sciences</i> , 2012 , 6, 21-30		6
226	Complete genome sequence of the aquatic bacterium <i>Runella slithyformis</i> type strain (LSU 4(T)). <i>Standards in Genomic Sciences</i> , 2012 , 6, 145-54		16
225	Complete Genome Sequence of <i>Paenibacillus</i> strain Y4.12MC10, a Novel <i>Paenibacillus lautus</i> strain Isolated from Obsidian Hot Spring in Yellowstone National Park. <i>Standards in Genomic Sciences</i> , 2012 , 6, 381-400		27

224	Genome sequence of the ocean sediment bacterium <i>Saccharomonospora marina</i> type strain (XMU15(T)). <i>Standards in Genomic Sciences</i> , 2012 , 6, 265-75	3
223	Complete genome sequence of the thermophilic sulfate-reducing ocean bacterium <i>Thermodesulfatator indicus</i> type strain (CIR29812(T)). <i>Standards in Genomic Sciences</i> , 2012 , 6, 155-64	9
222	Complete genome sequence of <i>Thauera aminoaromatica</i> strain MZ1T. <i>Standards in Genomic Sciences</i> , 2012 , 6, 325-35	47
221	Complete genome sequence of <i>Serratia plymuthica</i> strain AS12. <i>Standards in Genomic Sciences</i> , 2012 , 6, 165-73	15
220	Genome sequence of the homoacetogenic bacterium <i>Holophaga foetida</i> type strain (TMBS4(T)). <i>Standards in Genomic Sciences</i> , 2012 , 6, 174-84	23
219	Complete genome sequence of the orange-red pigmented, radioresistant <i>Deinococcus proteolyticus</i> type strain (MRP(T)). <i>Standards in Genomic Sciences</i> , 2012 , 6, 240-50	7
218	Complete genome sequence of <i>Thermovibrio ammonificans</i> HB-1(T), a thermophilic, chemolithoautotrophic bacterium isolated from a deep-sea hydrothermal vent. <i>Standards in Genomic Sciences</i> , 2012 , 7, 82-90	8
217	Complete genome sequence of the plant-associated <i>Serratia plymuthica</i> strain AS13. <i>Standards in Genomic Sciences</i> , 2012 , 7, 22-30	16
216	Complete genome sequence of <i>Marinomonas posidonica</i> type strain (IVIA-Po-181(T)). <i>Standards in Genomic Sciences</i> , 2012 , 7, 31-43	5
215	Genome sequence of the flexirubin-pigmented soil bacterium <i>Niabella soli</i> type strain (JS13-8(T)). <i>Standards in Genomic Sciences</i> , 2012 , 7, 210-20	0
214	Genome sequence of the Antarctic rhodopsins-containing flavobacterium <i>Gillisia limnaea</i> type strain (R-8282(T)). <i>Standards in Genomic Sciences</i> , 2012 , 7, 107-19	9
213	Complete genome sequence of the sulfate-reducing firmicute <i>Desulfotomaculum ruminis</i> type strain (DL(T)). <i>Standards in Genomic Sciences</i> , 2012 , 7, 304-19	18
212	Complete genome sequencing and analysis of <i>Saprospira grandis</i> str. Lewin, a predatory marine bacterium. <i>Standards in Genomic Sciences</i> , 2012 , 6, 84-93	19
211	Complete genome sequence of <i>Halopiger xanaduensis</i> type strain (SH-6(T)). <i>Standards in Genomic Sciences</i> , 2012 , 6, 31-42	7
210	Complete Genome Sequence of <i>Clostridium clariflavum</i> DSM 19732. <i>Standards in Genomic Sciences</i> , 2012 , 6, 104-15	40
209	Complete genome sequence of the facultatively chemolithoautotrophic and methylotrophic alpha Proteobacterium <i>Starkeya novella</i> type strain (ATCC 8093(T)). <i>Standards in Genomic Sciences</i> , 2012 , 7, 44-58	12
208	Complete genome sequence of <i>Terriglobus saanensis</i> type strain SP1PR4(T), an Acidobacteria from tundra soil. <i>Standards in Genomic Sciences</i> , 2012 , 7, 59-69	16
207	Complete genome sequence of the sulfur compounds oxidizing chemolithoautotroph <i>Sulfuricurvum kujiense</i> type strain (YK-1(T)). <i>Standards in Genomic Sciences</i> , 2012 , 6, 94-103	31

206	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	15
205	Complete genome sequence of the filamentous gliding predatory bacterium <i>Herpetosiphon aurantiacus</i> type strain (114-95(T)). <i>Standards in Genomic Sciences</i> , 2011 , 5, 356-70	35
204	Complete genome sequence of <i>Leadbetterella byssophila</i> type strain (4M15). <i>Standards in Genomic Sciences</i> , 2011 , 4, 2-12	15
203	Complete genome sequence of <i>Hydrogenobacter thermophilus</i> type strain (TK-6). <i>Standards in Genomic Sciences</i> , 2011 , 4, 131-43	8
202	Complete genome sequence of <i>Paludibacter propionigenes</i> type strain (WB4). <i>Standards in Genomic Sciences</i> , 2011 , 4, 36-44	24
201	Complete genome sequence of <i>Bacteroides helcogenes</i> type strain (P 36-108). <i>Standards in Genomic Sciences</i> , 2011 , 4, 45-53	7
200	Complete genome sequence of <i>Weeksella virosa</i> type strain (9751). <i>Standards in Genomic Sciences</i> , 2011 , 4, 81-90	8
199	Complete genome sequence of <i>Desulfobulbus propionicus</i> type strain (1pr3). <i>Standards in Genomic Sciences</i> , 2011 , 4, 100-10	39
198	Complete genome sequence of <i>Marivirga tractuosa</i> type strain (H-43). <i>Standards in Genomic Sciences</i> , 2011 , 4, 154-62	15
197	Complete genome sequence of <i>Desulfurococcus mucosus</i> type strain (O7/1). <i>Standards in Genomic Sciences</i> , 2011 , 4, 173-82	9
196	Complete genome sequence of <i>Cellulophaga lytica</i> type strain (LIM-21). <i>Standards in Genomic Sciences</i> , 2011 , 4, 221-32	23
195	Non-contiguous finished genome sequence of <i>Bacteroides coprosuis</i> type strain (PC139). <i>Standards in Genomic Sciences</i> , 2011 , 4, 233-43	4
194	Complete genome sequence of <i>Rhodospirillum rubrum</i> type strain (S1). <i>Standards in Genomic Sciences</i> , 2011 , 4, 293-302	31
193	Complete genome sequence of the extremely halophilic <i>Halanaerobium praevalens</i> type strain (GSL). <i>Standards in Genomic Sciences</i> , 2011 , 4, 312-21	24
192	Complete genome sequence of <i>Nitratifractor salsuginis</i> type strain (E9I37-1). <i>Standards in Genomic Sciences</i> , 2011 , 4, 322-30	10
191	Complete genome sequence of <i>Mahella australiensis</i> type strain (50-1 BON). <i>Standards in Genomic Sciences</i> , 2011 , 4, 331-41	6
190	Complete genome sequence of <i>Treponema succinifaciens</i> type strain (6091). <i>Standards in Genomic Sciences</i> , 2011 , 4, 361-70	20
189	Complete genome sequence of <i>Syntrophobotulus glycolicus</i> type strain (FIGlyRT). <i>Standards in Genomic Sciences</i> , 2011 , 4, 371-380	9

188	Complete genome sequence of the hyperthermophilic chemolithoautotroph <i>Pyrolobus fumarii</i> type strain (1A). <i>Standards in Genomic Sciences</i> , 2011 , 4, 381-92	10
187	Complete genome sequence of <i>Staphylothermus hellenicus</i> P8. <i>Standards in Genomic Sciences</i> , 2011 , 5, 12-20	5
186	Complete genome sequence of the acetate-degrading sulfate reducer <i>Desulfobacca acetoxidans</i> type strain (ASRB2). <i>Standards in Genomic Sciences</i> , 2011 , 4, 393-401	19
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42	Complete genome sequence of <i>Eggerthella lenta</i> type strain (IPP VPI 0255). <i>Standards in Genomic Sciences</i> , 2009 , 1, 174-82		31
41	Complete genome sequence of <i>Methanocorpusculum labreanum</i> type strain Z. <i>Standards in Genomic Sciences</i> , 2009 , 1, 197-203		26
40	Complete genome sequence of <i>Kangiella koreensis</i> type strain (SW-125). <i>Standards in Genomic Sciences</i> , 2009 , 1, 226-33		10
39	Complete genome sequence of <i>Jonesia denitrificans</i> type strain (Prevot 55134). <i>Standards in Genomic Sciences</i> , 2009 , 1, 262-9		10
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37	Complete genome sequence of <i>Rhodothermus marinus</i> type strain (R-10). <i>Standards in Genomic Sciences</i> , 2009 , 1, 283-90		20
36	Complete genome sequence of <i>Streptobacillus moniliformis</i> type strain (9901). <i>Standards in Genomic Sciences</i> , 2009 , 1, 300-7		16
35	Complete genome sequence of <i>Pirellula staleyi</i> type strain (ATCC 27377). <i>Standards in Genomic Sciences</i> , 2009 , 1, 308-16		26
34	Complete genome sequence of <i>Dyadobacter fermentans</i> type strain (NS114). <i>Standards in Genomic Sciences</i> , 2009 , 1, 133-40		19
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32	Complete genome sequence of <i>Desulfotomaculum acetoxidans</i> type strain (5575). <i>Standards in Genomic Sciences</i> , 2009 , 1, 242-53		33
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