

# Loren Hauser

## List of Publications by Citations

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7,128  
ext. citations

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#	Paper	IF	Citations
134	Genome divergence in two <i>Prochlorococcus</i> ecotypes reflects oceanic niche differentiation. <i>Nature</i> , <b>2003</b> , 424, 1042-7	50.4	904
133	Complete genome sequence of the metabolically versatile photosynthetic bacterium <i>Rhodospseudomonas palustris</i> . <i>Nature Biotechnology</i> , <b>2004</b> , 22, 55-61	44.5	553
132	Complete genome sequence of the ammonia-oxidizing bacterium and obligate chemolithoautotroph <i>Nitrosomonas europaea</i> . <i>Journal of Bacteriology</i> , <b>2003</b> , 185, 2759-73	3.5	436
131	<i>Burkholderia xenovorans</i> LB400 harbors a multi-replicon, 9.73-Mbp genome shaped for versatility. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2006</b> , 103, 15280-7	11.5	280
130	Mechanisms of thermal adaptation revealed from the genomes of the Antarctic Archaea <i>Methanogenium frigidum</i> and <i>Methanococcoides burtonii</i> . <i>Genome Research</i> , <b>2003</b> , 13, 1580-8	9.7	213
129	Community genomic and proteomic analyses of chemoautotrophic iron-oxidizing " <i>Leptospirillum rubrum</i> " (Group II) and " <i>Leptospirillum ferrodiazotrophum</i> " (Group III) bacteria in acid mine drainage biofilms. <i>Applied and Environmental Microbiology</i> , <b>2009</b> , 75, 4599-615	4.8	149
128	Genome sequence of the chemolithoautotrophic nitrite-oxidizing bacterium <i>Nitrobacter winogradskyi</i> Nb-255. <i>Applied and Environmental Microbiology</i> , <b>2006</b> , 72, 2050-63	4.8	141
127	The genome sequence of <i>Psychrobacter arcticus</i> 273-4, a psychroactive Siberian permafrost bacterium, reveals mechanisms for adaptation to low-temperature growth. <i>Applied and Environmental Microbiology</i> , <b>2010</b> , 76, 2304-12	4.8	134
126	The genome sequence of the psychrophilic archaeon, <i>Methanococcoides burtonii</i> : the role of genome evolution in cold adaptation. <i>ISME Journal</i> , <b>2009</b> , 3, 1012-35	11.9	128
125	Whole-genome analysis of the ammonia-oxidizing bacterium, <i>Nitrosomonas eutropha</i> C91: implications for niche adaptation. <i>Environmental Microbiology</i> , <b>2007</b> , 9, 2993-3007	5.2	128
124	Integrating "top-down" and "bottom-up" mass spectrometric approaches for proteomic analysis of <i>Shewanella oneidensis</i> . <i>Journal of Proteome Research</i> , <b>2002</b> , 1, 239-52	5.6	123
123	Multiple genome sequences reveal adaptations of a phototrophic bacterium to sediment microenvironments. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2008</b> , 105, 18543-8	11.5	100
122	An oligopeptide transport gene from <i>Candida albicans</i> . <i>Microbiology (United Kingdom)</i> , <b>1997</b> , 143 (Pt 2), 387-396	2.9	88
121	Complete genome sequence of the <i>Medicago</i> microsymbiont <i>Ensifer</i> ( <i>Sinorhizobium</i> ) <i>medicae</i> strain WSM419. <i>Standards in Genomic Sciences</i> , <b>2010</b> , 2, 77-86		80
120	Genome sequence of the obligate methanotroph <i>Methylosinus trichosporium</i> strain OB3b. <i>Journal of Bacteriology</i> , <b>2010</b> , 192, 6497-8	3.5	76
119	Non-contiguous finished genome sequence and contextual data of the filamentous soil bacterium <i>Ktedonobacter racemifer</i> type strain (SOSP1-21). <i>Standards in Genomic Sciences</i> , <b>2011</b> , 5, 97-111		72
118	Determination and comparison of the baseline proteomes of the versatile microbe <i>Rhodospseudomonas palustris</i> under its major metabolic states. <i>Journal of Proteome Research</i> , <b>2006</b> , 5, 287-98	5.6	64

117	Complete genome sequence of <i>Odoribacter splanchnicus</i> type strain (1651/6). <i>Standards in Genomic Sciences</i> , <b>2011</b> , 4, 200-9		62
116	Complete genome sequence of the aerobic marine methanotroph <i>Methylomonas methanica</i> MC09. <i>Journal of Bacteriology</i> , <b>2011</b> , 193, 7001-2	3.5	57
115	A novel ATPase on mouse chromosome 7 is a candidate gene for increased body fat. <i>Physiological Genomics</i> , <b>2000</b> , 4, 93-100	3.6	52
114	Genome sequence of the Arctic methanotroph <i>Methylobacter tundripaludum</i> SV96. <i>Journal of Bacteriology</i> , <b>2011</b> , 193, 6418-9	3.5	50
113	Complete genome sequence of <i>Geodermatophilus obscurus</i> type strain (G-20). <i>Standards in Genomic Sciences</i> , <b>2010</b> , 2, 158-67		49
112	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> , <b>2012</b> , 6, 181-200		46
111	Genome sequence of the methanotrophic alphaproteobacterium <i>Methylocystis</i> sp. strain Rockwell (ATCC 49242). <i>Journal of Bacteriology</i> , <b>2011</b> , 193, 2668-9	3.5	45
110	Biological consequences of ancient gene acquisition and duplication in the large genome of <i>Candidatus Solibacter usitatus</i> Ellin6076. <i>PLoS ONE</i> , <b>2011</b> , 6, e24882	3.7	41
109	Complete genome sequence of <i>Isosphaera pallida</i> type strain (IS1B). <i>Standards in Genomic Sciences</i> , <b>2011</b> , 4, 63-71		40
108	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> , <b>2014</b> , 9, 10		39
107	Complete genome sequence of <i>Truepera radiovictrix</i> type strain (RQ-24). <i>Standards in Genomic Sciences</i> , <b>2011</b> , 4, 91-9		37
106	Environment sensing and response mediated by ABC transporters. <i>BMC Genomics</i> , <b>2011</b> , 12 Suppl 1, S8	4.5	35
105	Complete genome sequence of the filamentous gliding predatory bacterium <i>Herpetosiphon aurantiacus</i> type strain (114-95(T)). <i>Standards in Genomic Sciences</i> , <b>2011</b> , 5, 356-70		35
104	Complete genome sequence of the thermophilic, hydrogen-oxidizing <i>Bacillus tusciae</i> type strain (T2) 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> , <b>2011</b> , 5, 121-34		35
103	Complete genome sequence of <i>Spirosoma linguale</i> type strain (1). <i>Standards in Genomic Sciences</i> , <b>2010</b> , 2, 176-85		35
102	Evaluation of "shotgun" proteomics for identification of biological threat agents in complex environmental matrixes: experimental simulations. <i>Analytical Chemistry</i> , <b>2005</b> , 77, 923-32	7.8	34
101	Complete genome sequence of <i>Desulfotomaculum acetoxidans</i> type strain (5575). <i>Standards in Genomic Sciences</i> , <b>2009</b> , 1, 242-53		33
100	Iron nutrition and physiological responses to iron stress in <i>Nitrosomonas europaea</i> . <i>Archives of Microbiology</i> , <b>2006</b> , 186, 107-18	3	33

99	Complete genome sequence of <i>Arcobacter nitrofigilis</i> type strain (CI). <i>Standards in Genomic Sciences</i> , <b>2010</b> , 2, 300-8		32
98	Complete genome sequence of <i>Sulfurimonas autotrophica</i> type strain (OK10). <i>Standards in Genomic Sciences</i> , <b>2010</b> , 3, 194-202		32
97	The genome sequence of <i>Methanohalophilus mahii</i> SLP(T) reveals differences in the energy metabolism among members of the Methanosarcinaceae inhabiting freshwater and saline environments. <i>Archaea</i> , <b>2010</b> , 2010, 690737	2	31
96	Complete genome sequence of <i>Cellulomonas flavigena</i> type strain (134). <i>Standards in Genomic Sciences</i> , <b>2010</b> , 3, 15-25		31
95	Complete genome sequence of the sulfur compounds oxidizing chemolithoautotroph <i>Sulfuricurvum kujiense</i> type strain (YK-1(T)). <i>Standards in Genomic Sciences</i> , <b>2012</b> , 6, 94-103		31
94	Complete genome sequence of <i>Eggerthella lenta</i> type strain (IPP VPI 0255). <i>Standards in Genomic Sciences</i> , <b>2009</b> , 1, 174-82		31
93	Complete genome sequence of <i>Desulfomicrobium baculatum</i> type strain (X). <i>Standards in Genomic Sciences</i> , <b>2009</b> , 1, 29-37		30
92	Complete genome sequence of <i>Halomicrobium mukohataei</i> type strain (arg-2). <i>Standards in Genomic Sciences</i> , <b>2009</b> , 1, 270-7		30
91	Complete genome sequence of <i>Veillonella parvula</i> type strain (Te3). <i>Standards in Genomic Sciences</i> , <b>2010</b> , 2, 57-65		29
90	Complete genome sequence of <i>Methanoculleus marisnigri</i> Romesser et al. 1981 type strain JR1. <i>Standards in Genomic Sciences</i> , <b>2009</b> , 1, 189-96		28
89	Complete genome sequence of <i>Planctomyces limnophilus</i> type strain (M290). <i>Standards in Genomic Sciences</i> , <b>2010</b> , 3, 47-56		28
88	Complete genome sequence of <i>Meiothermus ruber</i> type strain (21). <i>Standards in Genomic Sciences</i> , <b>2010</b> , 3, 26-36		28
87	Complete genome sequence of <i>Haloterrigena turkmenica</i> type strain (4k). <i>Standards in Genomic Sciences</i> , <b>2010</b> , 2, 107-16		28
86	Complete genome sequence of <i>Nakamurella multipartita</i> type strain (Y-104). <i>Standards in Genomic Sciences</i> , <b>2010</b> , 2, 168-75		28
85	Complete genome sequence of <i>Ferroglobus placidus</i> AEDII12DO. <i>Standards in Genomic Sciences</i> , <b>2011</b> , 5, 50-60		27
84	Complete genome sequence of <i>Olsenella uli</i> type strain (VPI D76D-27C). <i>Standards in Genomic Sciences</i> , <b>2010</b> , 3, 76-84		27
83	Complete genome sequence of <i>Desulfarculus baarsii</i> type strain (2st14). <i>Standards in Genomic Sciences</i> , <b>2010</b> , 3, 276-84		27
82	Complete genome sequence of <i>Aminobacterium colombiense</i> type strain (ALA-1). <i>Standards in Genomic Sciences</i> , <b>2010</b> , 2, 280-9		27

81	Complete genome sequence of the anaerobic, halophilic alkalithermophile <i>Natranaerobius thermophilus</i> JW/NM-WN-LF. <i>Journal of Bacteriology</i> , <b>2011</b> , 193, 4023-4	3.5	26
80	Complete genome sequence of <i>Methanocorpusculum labreanum</i> type strain Z. <i>Standards in Genomic Sciences</i> , <b>2009</b> , 1, 197-203		26
79	Complete genome sequence of <i>Pirellula staleyi</i> type strain (ATCC 27377). <i>Standards in Genomic Sciences</i> , <b>2009</b> , 1, 308-16		26
78	Complete genome sequence of the rapeseed plant-growth promoting <i>Serratia plymuthica</i> strain AS9. <i>Standards in Genomic Sciences</i> , <b>2012</b> , 6, 54-62		25
77	Complete genome sequence of <i>Streptosporangium roseum</i> type strain (NI 9100). <i>Standards in Genomic Sciences</i> , <b>2010</b> , 2, 29-37		25
76	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 zuelzeriae</i> in the genus <i>Treponema</i> as <i>Treponema caldaria</i> comb. nov., <i>Treponema stenostrepta</i> comb. nov., and <i>Treponema zuelzeriae</i> comb. nov. and nomenclature of the genus <i>Treponema</i>		24
75	Complete genome sequence of <i>Paludibacter propionigenes</i> type strain (WB4). <i>Standards in Genomic Sciences</i> , <b>2011</b> , 4, 36-44		24
74	Complete genome sequence of <i>Thermomonospora curvata</i> type strain (B9). <i>Standards in Genomic Sciences</i> , <b>2011</b> , 4, 13-22		24
73	Complete genome sequence of <i>Sulfurospirillum deleyianum</i> type strain (5175). <i>Standards in Genomic Sciences</i> , <b>2010</b> , 2, 149-57		24
72	Complete genome sequence of <i>Cellulophaga lytica</i> type strain (LIM-21). <i>Standards in Genomic Sciences</i> , <b>2011</b> , 4, 221-32		23
71	Complete genome sequence of <i>Cellulophaga algicola</i> type strain (IC166). <i>Standards in Genomic Sciences</i> , <b>2011</b> , 4, 72-80		22
70	Complete genome sequence of <i>Halorhabdus utahensis</i> type strain (AX-2). <i>Standards in Genomic Sciences</i> , <b>2009</b> , 1, 218-25		21
69	Complete genome sequence of <i>Treponema succinifaciens</i> type strain (6091). <i>Standards in Genomic Sciences</i> , <b>2011</b> , 4, 361-70		20
68	Complete genome sequence of <i>Alicyclobacillus acidocaldarius</i> type strain (104-IA). <i>Standards in Genomic Sciences</i> , <b>2010</b> , 2, 9-18		20
67	Complete genome sequence of <i>Sphaerobacter thermophilus</i> type strain (S 6022). <i>Standards in Genomic Sciences</i> , <b>2010</b> , 2, 49-56		20
66	Complete genome sequence of <i>Archaeoglobus profundus</i> type strain (AV18). <i>Standards in Genomic Sciences</i> , <b>2010</b> , 2, 327-46		20
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61	Complete genome sequence of <i>Acidaminococcus fermentans</i> type strain (VR4). <i>Standards in Genomic Sciences</i> , <b>2010</b> , 3, 1-14	19
60	Complete genome sequence of <i>Gordonia bronchialis</i> type strain (3410). <i>Standards in Genomic Sciences</i> , <b>2010</b> , 2, 19-28	19
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54	Complete genome sequence of <i>Desulfohalobium retbaense</i> type strain (HR(100)). <i>Standards in Genomic Sciences</i> , <b>2010</b> , 2, 38-48	17
53	An aminophospholipid translocase associated with body fat and type 2 diabetes phenotypes. <i>Obesity</i> , <b>2002</b> , 10, 695-702	17
52	Genome and Transcriptome of <i>Clostridium phytofermentans</i> , Catalyst for the Direct Conversion of Plant Feedstocks to Fuels. <i>PLoS ONE</i> , <b>2015</b> , 10, e0118285	3-7 16
51	Genome Sequence of the Obligate Gammaproteobacterial Methanotroph <i>Methylomicrobium album</i> Strain BG8. <i>Genome Announcements</i> , <b>2013</b> , 1, e0017013	16
50	Complete genome sequence of the aquatic bacterium <i>Runella slithyformis</i> type strain (LSU 4(T)). <i>Standards in Genomic Sciences</i> , <b>2012</b> , 6, 145-54	16
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48	Complete genome sequence of <i>Streptobacillus moniliformis</i> type strain (9901). <i>Standards in Genomic Sciences</i> , <b>2009</b> , 1, 300-7	16
47	Complete genome sequence of <i>Meiothermus silvanus</i> type strain (VI-R2). <i>Standards in Genomic Sciences</i> , <b>2010</b> , 3, 37-46	16
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42	Complete genome sequence of <i>Spirochaeta smaragdinae</i> type strain (SEBR 4228). <i>Standards in Genomic Sciences</i> , <b>2010</b> , 3, 136-44	15
41	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> , <b>2012</b> , 7, 44-58	15
40	Complete genome sequence of <i>Thermocrinis albus</i> type strain (HI 11/12). <i>Standards in Genomic Sciences</i> , <b>2010</b> , 2, 194-202	14
39	Complete genome sequence of <i>Ignisphaera aggregans</i> type strain (AQ1.S1). <i>Standards in Genomic Sciences</i> , <b>2010</b> , 3, 66-75	13
38	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> , <b>2012</b> , 7, 44-58	12
37	Complete genome sequence of <i>Thermosphaera aggregans</i> type strain (M11TL). <i>Standards in Genomic Sciences</i> , <b>2010</b> , 2, 245-59	12
36	Complete genome sequence of <i>Methanoplanus petrolearius</i> type strain (SEBR 4847). <i>Standards in Genomic Sciences</i> , <b>2010</b> , 3, 203-11	12
35	Complete genome sequence of the gliding, heparinolytic <i>Pedobacter saltans</i> type strain (113). <i>Standards in Genomic Sciences</i> , <b>2011</b> , 5, 30-40	11
34	Complete genome sequence of <i>Deinococcus maricopensis</i> type strain (LB-34). <i>Standards in Genomic Sciences</i> , <b>2011</b> , 4, 163-72	11
33	Complete genome sequence of <i>Beutenbergia cavernae</i> type strain (HKI 0122). <i>Standards in Genomic Sciences</i> , <b>2009</b> , 1, 21-8	11
32	Complete genome sequence of <i>Denitrovibrio acetiphilus</i> type strain (N2460). <i>Standards in Genomic Sciences</i> , <b>2010</b> , 2, 270-9	11
31	Complete genome sequence of <i>Nitratifactor salsuginis</i> type strain (E9I37-1). <i>Standards in Genomic Sciences</i> , <b>2011</b> , 4, 322-30	10
30	Complete genome sequence of the hyperthermophilic chemolithoautotroph <i>Pyrolobus fumarii</i> type strain (1A). <i>Standards in Genomic Sciences</i> , <b>2011</b> , 4, 381-92	10
29	Complete genome sequence of the thermophilic sulfur-reducer <i>Desulfurobacterium thermolithotrophum</i> type strain (BSA(T)) from a deep-sea hydrothermal vent. <i>Standards in Genomic Sciences</i> , <b>2011</b> , 5, 407-15	10
28	Complete genome sequence of <i>Jonesia denitrificans</i> type strain (Prevot 55134). <i>Standards in Genomic Sciences</i> , <b>2009</b> , 1, 262-9	10

27	Complete genome sequence of <i>Thermosediminibacter oceani</i> type strain (JW/IW-1228P). <i>Standards in Genomic Sciences</i> , <b>2010</b> , 3, 108-16	10
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22	Complete genome sequence of the thermophilic sulfate-reducing ocean bacterium <i>Thermodesulfatator indicus</i> type strain (CIR29812(T)). <i>Standards in Genomic Sciences</i> , <b>2012</b> , 6, 155-64	9
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11	Complete genome sequence of the orange-red pigmented, radioresistant <i>Deinococcus proteolyticus</i> type strain (MRP(T)). <i>Standards in Genomic Sciences</i> , <b>2012</b> , 6, 240-50	7
10	Complete genome sequence of <i>Ilyobacter polytropus</i> type strain (CuHbu1). <i>Standards in Genomic Sciences</i> , <b>2010</b> , 3, 304-14	7



9	Complete genome sequence of <i>Mahella australiensis</i> type strain (50-1 BON). <i>Standards in Genomic Sciences</i> , <b>2011</b> , 4, 331-41	6
8	Complete genome sequence of the haloalkaliphilic, obligately chemolithoautotrophic thiosulfate and sulfide-oxidizing $\beta$ proteobacterium <i>Thioalkalimicrobium cyclicum</i> type strain ALM 1 (DSM 14477(T)). <i>Standards in Genomic Sciences</i> , <b>2016</b> , 11, 38	5
7	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> , <b>2013</b> , 8, 15-25	5
6	Genome sequence of the <i>Thermotoga thermarum</i> type strain (LA3(T)) from an African solfataric spring. <i>Standards in Genomic Sciences</i> , <b>2014</b> , 9, 1105-17	4
5	Complete Genome Sequence of the Hyperthermophilic Sulfate-Reducing Bacterium <i>Thermodesulfobacterium geofontis</i> OPF15T. <i>Genome Announcements</i> , <b>2013</b> , 1, e0016213	4
4	Non-contiguous finished genome sequence of <i>Bacteroides coprosuis</i> type strain (PC139). <i>Standards in Genomic Sciences</i> , <b>2011</b> , 4, 233-43	4
3	Analysis and annotation of microbial genome sequences. <i>Genetic Engineering</i> , <b>2004</b> , 26, 225-38	4
2	Complete genome sequence of <i>Oceanithermus profundus</i> type strain (506). <i>Standards in Genomic Sciences</i> , <b>2011</b> , 4, 210-20	3
1	Complete genome sequence of <i>Syntrophobotulus glycolicus</i> type strain (FIGlyR). <i>Standards in Genomic Sciences</i> , <b>2011</b> , 4, 371-80	3