

Iain C Sutcliffe

List of Publications by Year in descending order

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Version: 2024-02-01

130
papers

4,959
citations

94433

37
h-index

110387

64
g-index

135
all docs

135
docs citations

135
times ranked

4945
citing authors

#	ARTICLE	IF	CITATIONS
1	ICSP response to "Science depends on nomenclature, but nomenclature is not science"™. <i>Nature Reviews Microbiology</i> , 2022, 20, 249-250.	28.6	14
2	It is time for a new type of type to facilitate naming the microbial world. <i>New Microbes and New Infections</i> , 2022, 47, 100991.	1.6	12
3	ICSP response to "Regulating access can restrict participation in reporting new species and taxa"™. <i>Nature Microbiology</i> , 2022, 7, 1711-1712.	13.3	5
4	Preparing a revision of the International Code of Nomenclature of Prokaryotes. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2021, 71, .	1.7	17
5	The <i>International Journal of Systematic and Evolutionary Microbiology</i> moves to "true continuous publication"™ at the beginning of 2021: Proposals to emend Rule 24b (2), Note 1 to Rule 27 and Note 2 to Rule 33b of the International Code of Nomenclature of Prokaryotes. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2021, 71, .	1.7	1
6	Emendation of Rules 5b, 8, 15 and 22 of the International Code of Nomenclature of Prokaryotes to include the rank of phylum. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2021, 71, .	1.7	27
7	Proposal of <i>Carbonactinosporaceae</i> fam. nov. within the class Actinomycetia. Reclassification of <i>Streptomyces thermoautotrophicus</i> as <i>Carbonactinospira thermoautotrophica</i> gen. nov., comb. nov. <i>Systematic and Applied Microbiology</i> , 2021, 44, 126223.	2.8	20
8	In memoriam "David Ernest Minnikin (1939" 2021). <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2021, 71, .	1.7	2
9	Out with the old and in with the new: time to rethink twentieth century chemotaxonomic practices in bacterial taxonomy. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2021, 71, .	1.7	17
10	Phylogenomic Reappraisal of Fatty Acid Biosynthesis, Mycolic Acid Biosynthesis and Clinical Relevance Among Members of the Genus <i>Corynebacterium</i> . <i>Frontiers in Microbiology</i> , 2021, 12, 802532.	3.5	12
11	Bringing the diversity of Planctomycetes into the light: Introduction to papers from the special issue on novel taxa of Planctomycetes. <i>Antonie Van Leeuwenhoek</i> , 2020, 113, 1715-1726.	1.7	4
12	Roadmap for naming uncultivated Archaea and Bacteria. <i>Nature Microbiology</i> , 2020, 5, 987-994.	13.3	115
13	Opinion: Response to concerns about the use of DNA sequences as types in the nomenclature of prokaryotes. <i>Systematic and Applied Microbiology</i> , 2020, 43, 126070.	2.8	8
14	Minutes of the International Committee on Systematics of Prokaryotes online discussion on the proposed use of gene sequences as type for naming of prokaryotes, and outcome of vote. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2020, 70, 4416-4417.	1.7	31
15	The van Niel International Prize for Studies in Bacterial Systematics, awarded in 2020 to Tanja Woyke. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2020, 70, 5594-5595.	1.7	1
16	Valediction: descriptions of novel prokaryotic taxa published in <i>Antonie van Leeuwenhoek</i> " change in editorial policy and a signpost to the future?. <i>Antonie Van Leeuwenhoek</i> , 2019, 112, 1281-1282.	1.7	4
17	Genomic analysis of a novel <i>Rhodococcus</i> (<i>Prescottella</i>) <i>equi</i> isolate from a bovine host. <i>Archives of Microbiology</i> , 2019, 201, 1317-1321.	2.2	4
18	Mechanism of Phosphatidylglycerol Activation Catalyzed by Prolipoprotein Diacylglyceryl Transferase. <i>Journal of Physical Chemistry B</i> , 2019, 123, 7092-7102.	2.6	11

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19	Reflections on the introduction of the Digital Protologue Database – A partial success?. <i>Systematic and Applied Microbiology</i> , 2019, 42, 1-2.	2.8	4
20	Reflections on the introduction of the Digital Protologue Database – a partial success?. <i>Antonie Van Leeuwenhoek</i> , 2019, 112, 141-143.	1.7	5
21	Refined Systematics of the Genus <i>Rhodococcus</i> Based on Whole Genome Analyses. <i>Microbiology Monographs</i> , 2019, , 1-21.	0.6	10
22	Proposal for changes in the International Code of Nomenclature of Prokaryotes: granting priority to Candidatus names. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2019, 69, 2174-2175.	1.7	27
23	Antidiabetic –gliptins– affect biofilm formation by <i>Streptococcus mutans</i> . <i>Microbiological Research</i> , 2018, 209, 79-85.	5.3	6
24	Deciphering the molecular basis of mycobacteria and lipoglycan recognition by the C-type lectin Dectin-2. <i>Scientific Reports</i> , 2018, 8, 16840.	3.3	34
25	Revisiting the Taxonomic Status of the Biomedically and Industrially Important Genus <i>Amycolatopsis</i> , Using a Phylogenomic Approach. <i>Frontiers in Microbiology</i> , 2018, 9, 2281.	3.5	23
26	Genomic analyses reveal two distinct lineages of <i>Corynebacterium ulcerans</i> strains. <i>New Microbes and New Infections</i> , 2018, 25, 7-13.	1.6	17
27	Proposal of the suffix –ota to denote phyla. Addendum to –Proposal to include the rank of phylum in the International Code of Nomenclature of Prokaryotes–™. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2018, 68, 967-969.	1.7	136
28	<i>Blastococcus atacamensis</i> sp. nov., a novel strain adapted to life in the Yungay core region of the Atacama Desert. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2018, 68, 2712-2721.	1.7	33
29	Introducing a Digital Protologue: A timely move towards a database-driven systematics of Archaea and Bacteria. <i>Systematic and Applied Microbiology</i> , 2017, 40, 121-122.	2.8	40
30	Introducing a digital protologue: a timely move towards a database-driven systematics of archaea and bacteria. <i>Antonie Van Leeuwenhoek</i> , 2017, 110, 455-456.	1.7	85
31	Genomic analysis of endemic clones of toxigenic and non-toxigenic <i>Corynebacterium diphtheriae</i> in Belarus during and after the major epidemic in 1990s. <i>BMC Genomics</i> , 2017, 18, 873.	2.8	41
32	<i>Mycobacterium eburneum</i> sp. nov., a non-chromogenic, fast-growing strain isolated from sputum. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2017, 67, 3174-3181.	1.7	13
33	Two novel species of rapidly growing mycobacteria: <i>Mycobacterium lehmannii</i> sp. nov. and <i>Mycobacterium neumannii</i> sp. nov.. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2017, 67, 4948-4955.	1.7	12
34	Next-generation systematics: An innovative approach to resolve the structure of complex prokaryotic taxa. <i>Scientific Reports</i> , 2016, 6, 38392.	3.3	114
35	Cosmid based mutagenesis causes genetic instability in <i>Streptomyces coelicolor</i> , as shown by targeting of the lipoprotein signal peptidase gene. <i>Scientific Reports</i> , 2016, 6, 29495.	3.3	4
36	Comment on Tocheva et al. –Sporulation, bacterial cell envelopes and the origin of life–. <i>Nature Reviews Microbiology</i> , 2016, 14, 600-600.	28.6	5

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37	Bacterial glycobiology: rhamnose-containing cell wall polysaccharides in Gram-positive bacteria. <i>FEMS Microbiology Reviews</i> , 2016, 40, 464-479.	8.6	146
38	Charting stormy waters: A commentary on the nomenclature of the equine pathogen variously named <i>Prescottella equi</i> and <i>Rhodococcus equi</i> and <i>Rhodococcus hoagii</i> . <i>Equine Veterinary Journal</i> , 2015, 47, 508-509.	1.7	13
39	Adherence and invasive properties of <i>Corynebacterium diphtheriae</i> strains correlates with the predicted membrane-associated and secreted proteome. <i>BMC Genomics</i> , 2015, 16, 765.	2.8	47
40	Challenging the anthropocentric emphasis on phenotypic testing in prokaryotic species descriptions: rip it up and start again. <i>Frontiers in Genetics</i> , 2015, 6, 218.	2.3	57
41	Characterisation of SEQ0694 (PrsA/PrtM) of <i>Streptococcus equi</i> as a functional peptidyl-prolyl isomerase affecting multiple secreted protein substrates. <i>Molecular BioSystems</i> , 2015, 11, 3279-3286.	2.9	8
42	Genomic analyses confirm close relatedness between <i>Rhodococcus defluvi</i> and <i>Rhodococcus equi</i> (<i>Rhodococcus hoagii</i>). <i>Archives of Microbiology</i> , 2015, 197, 113-116.	2.2	17
43	Proposal to include the rank of phylum in the International Code of Nomenclature of Prokaryotes. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2015, 65, 4284-4287.	1.7	84
44	Discovery of a cell wall porin in the mycolic acid-containing actinomycete <i>Dietzia Amaris</i> DSM 43672. <i>FEBS Journal</i> , 2014, 281, 2030-2041.	4.7	7
45	Comparative proteome analysis of <i>Acidaminococcus intestini</i> supports a relationship between outer membrane biogenesis in Negativicutes and Proteobacteria. <i>Archives of Microbiology</i> , 2014, 196, 307-310.	2.2	11
46	Comparative genomic analyses reveal a lack of a substantial signature of host adaptation in <i>Rhodococcus equi</i> (<i>Prescottella equi</i>). <i>Pathogens and Disease</i> , 2014, 71, 352-356.	2.0	11
47	Celebrating the 80th anniversary of Antonie van Leeuwenhoek: a special issue. <i>Antonie Van Leeuwenhoek</i> , 2014, 106, 1-2.	1.7	1
48	Proposal to replace the illegitimate genus name <i>Prescottia</i> Jones et al. 2013 with the genus name <i>Prescottella</i> gen. nov. and to replace the illegitimate combination <i>Prescottia equi</i> Jones et al. 2013 with <i>Prescottella equi</i> comb. nov. <i>Antonie Van Leeuwenhoek</i> , 2013, 103, 1405-1407.	1.7	35
49	Publication of descriptions of novel bacterial taxa in Antonie van Leeuwenhoek. <i>Antonie Van Leeuwenhoek</i> , 2013, 103, 1-2.	1.7	7
50	A call to action for the International Committee on Systematics of Prokaryotes. <i>Trends in Microbiology</i> , 2013, 21, 51-52.	7.7	31
51	<i>Prescottia equi</i> gen. nov., comb. nov.: a new home for an old pathogen. <i>Antonie Van Leeuwenhoek</i> , 2013, 103, 655-671.	1.7	35
52	Gram-Positive Bacterial Lipoglycans Based on a Glycosylated Diacylglycerol Lipid Anchor Are Microbe-Associated Molecular Patterns Recognized by TLR2. <i>PLoS ONE</i> , 2013, 8, e81593.	2.5	15
53	Polymicrobial challenges to Koch's postulates: Ecological lessons from the bacterial vaginosis and cystic fibrosis microbiomes. <i>Innate Immunity</i> , 2012, 18, 774-783.	2.4	35
54	Exposing a chink in the armor of methicillin-resistant <i>Staphylococcus aureus</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 18637-18638.	7.1	5

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55	A phylum level analysis reveals lipoprotein biosynthesis to be a fundamental property of bacteria. <i>Protein and Cell</i> , 2012, 3, 163-170.	11.0	42
56	The impact of pH and nutrient stress on the growth and survival of <i>Streptococcus agalactiae</i> . <i>Antonie Van Leeuwenhoek</i> , 2012, 102, 277-287.	1.7	34
57	A call to arms for systematists: revitalising the purpose and practises underpinning the description of novel microbial taxa. <i>Antonie Van Leeuwenhoek</i> , 2012, 101, 13-20.	1.7	97
58	The BISMIS 2011 special issue on prokaryotic systematics, a vital discipline entering a period of transition. <i>Antonie Van Leeuwenhoek</i> , 2012, 101, 1-2.	1.7	2
59	Cell envelope architecture in the <i>Chloroflexi</i> : a shifting frontline in a phylogenetic turf war. <i>Environmental Microbiology</i> , 2011, 13, 279-282.	3.8	105
60	Cell envelope architecture in the <i>Chloroflexi</i> : a shifting frontline in a phylogenetic turf war. <i>Environmental Microbiology</i> , 2011, 13, 2387-2387.	3.8	3
61	Priming and elongation: dissection of the lipoteichoic acid biosynthetic pathway in Gram-positive bacteria. <i>Molecular Microbiology</i> , 2011, 79, 553-556.	2.5	11
62	Dissecting the complete lipoprotein biogenesis pathway in <i>Streptomyces scabies</i> . <i>Molecular Microbiology</i> , 2011, 80, 1395-1412.	2.5	42
63	New insights into the distribution of WXG100 protein secretion systems. <i>Antonie Van Leeuwenhoek</i> , 2011, 99, 127-131.	1.7	30
64	A proteomic investigation of <i>Streptococcus agalactiae</i> reveals that human serum induces the C protein β^2 antigen and arginine deiminase. <i>Microbes and Infection</i> , 2011, 13, 757-760.	1.9	12
65	Investigating lipoprotein biogenesis and function in the model Gram-positive bacterium <i>Streptomyces coelicolor</i> . <i>Molecular Microbiology</i> , 2010, 77, 943-957.	2.5	56
66	A phylum level perspective on bacterial cell envelope architecture. <i>Trends in Microbiology</i> , 2010, 18, 464-470.	7.7	225
67	The Rhodococcal Cell Envelope: Composition, Organisation and Biosynthesis. <i>Microbiology Monographs</i> , 2010, , 29-71.	0.6	15
68	A proteomic investigation of <i>Streptococcus agalactiae</i> grown under conditions associated with neonatal exposure reveals the upregulation of the putative virulence factor C protein β^2 antigen. <i>International Journal of Medical Microbiology</i> , 2010, 300, 331-337.	3.6	14
69	Macroamphiphilic Components of Thermophilic Actinomycetes: Identification of Lipoteichoic Acid in <i>Thermobifida fusca</i> . <i>Journal of Bacteriology</i> , 2009, 191, 152-160.	2.2	24
70	The <i>Streptococcus equi</i> prophage-encoded protein SEQ2045 is a hyaluronan-specific hyaluronate lyase that is produced during equine infection. <i>Microbiology (United Kingdom)</i> , 2009, 155, 443-449.	1.8	20
71	Impact of <i>lgt</i> mutation on lipoprotein biosynthesis and in vitro phenotypes of <i>Streptococcus agalactiae</i> . <i>Microbiology (United Kingdom)</i> , 2009, 155, 1451-1458.	1.8	18
72	Expression of the MtsA lipoprotein of <i>Streptococcus agalactiae</i> A909 is regulated by manganese and iron. <i>Antonie Van Leeuwenhoek</i> , 2009, 95, 101-109.	1.7	28

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73	Phenotypic variation in <i>Streptomyces</i> sp. DSM 40537, a lipoteichoic acid producing actinomycete. <i>Letters in Applied Microbiology</i> , 2009, 48, 226-229.	2.2	10
74	Lipoprotein biogenesis in Gram-positive bacteria: knowing when to hold $\hat{e}m$, knowing when to fold $\hat{e}m$. <i>Trends in Microbiology</i> , 2009, 17, 13-21.	7.7	181
75	Lipoteichoic acid biosynthesis: two steps forwards, one step sideways?. <i>Trends in Microbiology</i> , 2009, 17, 219-225.	7.7	46
76	<i>Pyramidobacter piscolens</i> gen. nov., sp. nov., a member of the phylum 'Synergistetes' isolated from the human oral cavity. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2009, 59, 972-980.	1.7	108
77	Methods for the bioinformatic identification of bacterial lipoproteins encoded in the genomes of Gram-positive bacteria. <i>World Journal of Microbiology and Biotechnology</i> , 2008, 24, 2377-2382.	3.6	108
78	Actinomycetologists: a vibrant and strong scientific community. Papers from the 14th International Symposium on the Biology of Actinomycetes. <i>Antonie Van Leeuwenhoek</i> , 2008, 94, 1-2.	1.7	1
79	Mannan Chain Length Controls Lipoglycans Signaling via and Binding to TLR2. <i>Journal of Immunology</i> , 2008, 180, 6696-6702.	0.8	73
80	Bioinformatic insights into the biosynthesis of the Group B carbohydrate in <i>Streptococcus agalactiae</i> . <i>Microbiology (United Kingdom)</i> , 2008, 154, 1354-1363.	1.8	28
81	Making bioinformatics projects a meaningful experience in an undergraduate biotechnology or biomedical science programme. <i>Bioscience Education</i> , 2007, 10, 1-2.	0.4	1
82	<i>Prevotella maculosa</i> sp. nov., isolated from the human oral cavity. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2007, 57, 2936-2939.	1.7	28
83	Putative lipoproteins identified by bioinformatic genome analysis of <i>Leifsonia xyli</i> ssp. <i>xyli</i> , the causative agent of sugarcane ratoon stunting disease. <i>Molecular Plant Pathology</i> , 2007, 8, 121-128.	4.2	14
84	\hat{e} Antonie van Leeuwenhoek for the era of online academic publishing \hat{e} . <i>Antonie Van Leeuwenhoek</i> , 2007, 91, 97-98.	1.7	0
85	Identification of a lipoarabinomannan-like lipoglycan in the actinomycete <i>Gordonia bronchialis</i> . <i>Archives of Microbiology</i> , 2006, 184, 425-427.	2.2	5
86	The \hat{I} f E Cell Envelope Stress Response of <i>Streptomyces coelicolor</i> Is Influenced by a Novel Lipoprotein, CseA. <i>Journal of Bacteriology</i> , 2006, 188, 7222-7229.	2.2	57
87	<i>Dermacoccus abyssi</i> sp. nov., a piezotolerant actinomycete isolated from the Mariana Trench. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2006, 56, 1233-1237.	1.7	62
88	Mutation of the Maturase Lipoprotein Attenuates the Virulence of <i>Streptococcus equi</i> to a Greater Extent than Does Loss of General Lipoprotein Lipidation. <i>Infection and Immunity</i> , 2006, 74, 6907-6919.	2.2	55
89	<i>Prevotella bergensis</i> sp. nov., isolated from human infections. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2006, 56, 609-612.	1.7	25
90	<i>Williamsia marianensis</i> sp. nov., a novel actinomycete isolated from the Mariana Trench. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2006, 56, 1123-1126.	1.7	37

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91	Lipoarabinomannansâ€”structurally diverse and functionally enigmatic macroamphiphiles of mycobacteria and related actinomycetes. <i>Tuberculosis</i> , 2005, 85, 205-206.	1.9	12
92	<i>Prevotella marshii</i> sp. nov. and <i>Prevotella baroniae</i> sp. nov., isolated from the human oral cavity. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2005, 55, 1551-1555.	1.7	70
93	Characterization of a Truncated Lipoarabinomannan from the Actinomycete <i>Turicella otitidis</i> . <i>Journal of Bacteriology</i> , 2005, 187, 854-861.	2.2	35
94	Putative lipoproteins of <i>Streptococcus agalactiae</i> identified by bioinformatic genome analysis. <i>Antonie Van Leeuwenhoek</i> , 2004, 85, 305-315.	1.7	23
95	Purification and characterisation of lipoglycan macroamphiphiles from <i>Propionibacterium acnes</i> . <i>Antonie Van Leeuwenhoek</i> , 2004, 86, 77-85.	1.7	20
96	Lipoproteins of <i>Mycobacterium tuberculosis</i> : an abundant and functionally diverse class of cell envelope components. <i>FEMS Microbiology Reviews</i> , 2004, 28, 645-659.	8.6	160
97	Combined solvent and water activity stresses on turgor regulation and membrane adaptation in <i>Oceanimonas baumannii</i> ATCC 700832. <i>Antonie Van Leeuwenhoek</i> , 2003, 83, 275-283.	1.7	10
98	Lipoprotein signal peptidase of <i>Streptococcus suis</i> serotype 2. <i>Microbiology (United Kingdom)</i> , 2003, 149, 1399-1407.	1.8	32
99	The Cell Wall of the Pathogenic Bacterium <i>Rhodococcus equi</i> Contains Two Channel-Forming Proteins with Different Properties. <i>Journal of Bacteriology</i> , 2003, 185, 2952-2960.	2.2	22
100	A Novel Lipoarabinomannan from the Equine Pathogen <i>Rhodococcus equi</i> . <i>Journal of Biological Chemistry</i> , 2002, 277, 31722-31733.	3.4	85
101	The molecular basis of <i>Streptococcus equi</i> infection and disease. <i>Microbes and Infection</i> , 2002, 4, 501-510.	1.9	107
102	Pattern searches for the identification of putative lipoprotein genes in Gram-positive bacterial genomes. <i>Microbiology (United Kingdom)</i> , 2002, 148, 2065-2077.	1.8	150
103	Surface Immunolocalisation of HPr in the Equine Pathogen <i>Streptococcus equi</i> . <i>Systematic and Applied Microbiology</i> , 2001, 24, 486-489.	2.8	4
104	Reclassification of [<i>Pseudomonas</i>] <i>doudoroffii</i> (Baumann et al. 1983) into the genus <i>Oceanomonas</i> gen. nov. as <i>Oceanomonas doudoroffii</i> comb. nov., and description of a phenol-degrading bacterium from estuarine water as <i>Oceanomonas baumannii</i> sp. nov.. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2001, 51, 67-72.	1.7	49
105	The modification of the membrane of <i>Oceanomonas baumannii</i> when subjected to both osmotic and organic solvent stress. <i>FEMS Microbiology Letters</i> , 2000, 189, 149-154.	1.8	35
106	Characterization of Acid Phosphatase Activities in the Equine Pathogen <i>Streptococcus equi</i> . <i>Systematic and Applied Microbiology</i> , 2000, 23, 325-329.	2.8	22
107	Identification of Methionine-processed HPr in the Equine Pathogen <i>Streptococcus equi</i> . <i>Systematic and Applied Microbiology</i> , 2000, 23, 330-332.	2.8	1
108	Characterisation of a lipomannan lipoglycan from the mycolic acid containing actinomycete <i>Dietzia maris</i> . <i>Antonie Van Leeuwenhoek</i> , 2000, 78, 195-201.	1.7	47

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109	Identification of Lipoprotein Homologues of Pneumococcal PsaA in the Equine Pathogens <i>Streptococcus equi</i> and <i>Streptococcus zooepidemicus</i> . <i>Infection and Immunity</i> , 2000, 68, 6048-6051.	2.2	22
110	Identification of a Lipoarabinomannan-like Lipoglycan in <i>Gordonia rubropertincta</i> . <i>Systematic and Applied Microbiology</i> , 1999, 22, 530-533.	2.8	26
111	Cell envelope composition and organisation in the genus <i>Rhodococcus</i> . <i>Antonie Van Leeuwenhoek</i> , 1998, 74, 49-58.	1.7	78
112	Macroamphiphilic cell envelope components of <i>Rhodococcus equi</i> and closely related bacteria. <i>Veterinary Microbiology</i> , 1997, 56, 287-299.	1.9	44
113	Organization and nucleotide sequence of the <i>Streptococcus mutans</i> galactose operon. <i>Gene</i> , 1996, 180, 137-144.	2.2	34
114	A Chemotaxonomic Study of the Lipoglycans of <i>Rhodococcus rhodnii</i> N445 (NCIMB 11279). <i>Zentralblatt Fur Bakteriologie: International Journal of Medical Microbiology</i> , 1996, 285, 11-19.	0.5	21
115	Lipoproteins of gram-positive bacteria. <i>Journal of Bacteriology</i> , 1995, 177, 1123-1128.	2.2	381
116	A chemotaxonomic appraisal of the distribution of lipomannans within the genus <i>Micrococcus</i> . <i>FEMS Microbiology Letters</i> , 1995, 133, 233-237.	1.8	7
117	Identification of a lipoarabinomannan-like lipoglycan in <i>Corynebacterium matruchotii</i> . <i>Archives of Oral Biology</i> , 1995, 40, 1119-1124.	1.8	27
118	<i>Stomatococcus mucilaginosus</i> produces a mannose-containing lipoglycan rather than lipoteichoic acid. <i>Archives of Microbiology</i> , 1995, 163, 70-75.	2.2	5
119	The Lipoteichoic Acids and Lipoglycans of Gram-positive Bacteria: A Chemotaxonomic Perspective. <i>Systematic and Applied Microbiology</i> , 1995, 17, 467-480.	2.8	41
120	Identification of a Lipomannan from <i>Rothia dentocariosa</i> . <i>Systematic and Applied Microbiology</i> , 1994, 17, 321-326.	2.8	15
121	Identification of <i>Streptococcus mutans</i> antigen D as the HPr component of the sugar-phosphotransferase transport system. <i>FEMS Microbiology Letters</i> , 1993, 107, 67-70.	1.8	13
122	The Phospholipids of <i>Propionibacterium freudenreichii</i> : Absence of Phosphatidylinositol Mannosides. <i>Systematic and Applied Microbiology</i> , 1993, 16, 9-12.	2.8	6
123	Extraction of lipoteichoic acid from <i>Streptococcus mutans</i> with the non-ionic detergent Triton X-114. <i>Journal of Microbiological Methods</i> , 1993, 17, 215-225.	1.6	11
124	Transport of Sugars, Including Sucrose, by the msm Transport System of <i>Streptococcus mutans</i> . <i>Journal of Dental Research</i> , 1993, 72, 1386-1390.	5.2	57
125	MsmE, a lipoprotein involved in sugar transport in <i>Streptococcus mutans</i> . <i>Journal of Bacteriology</i> , 1993, 175, 1853-1855.	2.2	48
126	Cloning and expression of the multiple sugar metabolism (msm) operon of <i>Streptococcus mutans</i> in heterologous streptococcal hosts. <i>Infection and Immunity</i> , 1993, 61, 1121-1125.	2.2	26

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127	Abnormal activation of glycogen synthesis in fibroblasts from NIDDM subjects. Evidence for an abnormality specific to glucose metabolism. <i>Diabetes</i> , 1993, 42, 583-589.	0.6	16
128	Identification of <i>Streptococcus mutans</i> antigen D as the HPr component of the sugar-phosphotransferase transport system. <i>FEMS Microbiology Letters</i> , 1993, 107, 67-70.	1.8	4
129	Atypical lipoteichoic acids of gram-positive bacteria. <i>Journal of Bacteriology</i> , 1991, 173, 7065-7069.	2.2	81
130	An inositol containing lipomannan from <i>Propionibacterium freudenreichii</i> . <i>FEMS Microbiology Letters</i> , 1989, 59, 249-251.	1.8	11