

Christopher G. Dowson

List of Publications by Year in descending order

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163
papers

12,836
citations

20817

60
h-index

26613

107
g-index

172
all docs

172
docs citations

172
times ranked

10666
citing authors

#	ARTICLE	IF	CITATIONS
1	Intercontinental Spread of a Multiresistant Clone of Serotype 23F <i>Streptococcus pneumoniae</i> . <i>Journal of Infectious Diseases</i> , 1991, 164, 302-306.	4.0	463
2	Development of a Multilocus Sequence Typing Scheme for the Opportunistic Pathogen <i>Pseudomonas aeruginosa</i> . <i>Journal of Clinical Microbiology</i> , 2004, 42, 5644-5649.	3.9	450
3	Localized sex in bacteria. <i>Nature</i> , 1991, 349, 29-31.	27.8	436
4	Horizontal transfer of penicillin-binding protein genes in penicillin-resistant clinical isolates of <i>Streptococcus pneumoniae</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1989, 86, 8842-8846.	7.1	411
5	Horizontal transfer of multiple penicillin-binding protein genes, and capsular biosynthetic genes, in natural populations of <i>Streptococcus pneumoniae</i> . <i>Molecular Microbiology</i> , 1991, 5, 2255-2260.	2.5	344
6	<i>Burkholderia cepacia</i> complex bacteria: opportunistic pathogens with important natural biology. <i>Journal of Applied Microbiology</i> , 2008, 104, 1539-1551.	3.1	336
7	The Genome of <i>Burkholderia cenocepacia</i> J2315, an Epidemic Pathogen of Cystic Fibrosis Patients. <i>Journal of Bacteriology</i> , 2009, 191, 261-277.	2.2	329
8	Metal complexes as a promising source for new antibiotics. <i>Chemical Science</i> , 2020, 11, 2627-2639.	7.4	290
9	Taxon K, a complex within the <i>Burkholderia cepacia</i> complex, comprises at least two novel species, <i>Burkholderia contaminans</i> sp. nov. and <i>Burkholderia lata</i> sp. nov.. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2009, 59, 102-111.	1.7	280
10	Evolution of penicillin resistance in <i>Streptococcus pneumoniae</i> ; the role of <i>Streptococcus mitis</i> in the formation of a low affinity PBP2B in <i>S. pneumoniae</i> . <i>Molecular Microbiology</i> , 1993, 9, 635-643.	2.5	264
11	Genetic Relationships between Clinical Isolates of <i>Streptococcus pneumoniae</i> , <i>Streptococcus oralis</i> , and <i>Streptococcus mitis</i> : Characterization of Atypical Pneumococci and Organisms Allied to <i>S. mitis</i> Harboring <i>S. pneumoniae</i> Virulence Factor-Encoding Genes. <i>Infection and Immunity</i> , 2000, 68, 1374-1382.	2.2	259
12	Development of a Multilocus Sequence Typing Scheme for the Pig Pathogen <i>Streptococcus suis</i> : Identification of Virulent Clones and Potential Capsular Serotype Exchange. <i>Journal of Clinical Microbiology</i> , 2002, 40, 3671-3680.	3.9	236
13	Expression of resistance to tetracyclines in strains of methicillin-resistant <i>Staphylococcus aureus</i> . <i>Journal of Antimicrobial Chemotherapy</i> , 2000, 45, 763-770.	3.0	233
14	Bacterial cell wall assembly: still an attractive antibacterial target. <i>Trends in Biotechnology</i> , 2011, 29, 167-173.	9.3	230
15	<i>Burkholderia latens</i> sp. nov., <i>Burkholderia diffusa</i> sp. nov., <i>Burkholderia arboris</i> sp. nov., <i>Burkholderia seminalis</i> sp. nov. and <i>Burkholderia metallica</i> sp. nov., novel species within the <i>Burkholderia cepacia</i> complex. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2008, 58, 1580-1590.	1.7	218
16	Rapid Evolution of Virulence and Drug Resistance in the Emerging Zoonotic Pathogen <i>Streptococcus suis</i> . <i>PLoS ONE</i> , 2009, 4, e6072.	2.5	214
17	Rapid Covalent-Probe Discovery by Electrophile-Fragment Screening. <i>Journal of the American Chemical Society</i> , 2019, 141, 8951-8968.	13.7	213
18	Penicillin-resistant viridans streptococci have obtained altered penicillin-binding protein genes from penicillin-resistant strains of <i>Streptococcus pneumoniae</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1990, 87, 5858-5862.	7.1	205

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19	Recruitment of a penicillin-binding protein gene from <i>Neisseria flavescens</i> during the emergence of penicillin resistance in <i>Neisseria meningitidis</i> . Proceedings of the National Academy of Sciences of the United States of America, 1989, 86, 8988-8992.	7.1	198
20	Genetics of resistance to third-generation cephalosporins in clinical isolates of <i>Streptococcus pneumoniae</i> . Molecular Microbiology, 1992, 6, 2461-2465.	2.5	197
21	Barriers to Genetic Exchange between Bacterial Species: <i>Streptococcus pneumoniae</i> Transformation. Journal of Bacteriology, 2000, 182, 1016-1023.	2.2	194
22	Multilocus Sequence Typing Scheme That Provides Both Species and Strain Differentiation for the <i>Burkholderia cepacia</i> Complex. Journal of Clinical Microbiology, 2005, 43, 4665-4673.	3.9	193
23	Larval therapy for leg ulcers (VenUS II): randomised controlled trial. BMJ: British Medical Journal, 2009, 338, b773-b773.	2.3	193
24	Origin and molecular epidemiology of penicillin-binding-protein-mediated resistance to β -lactam antibiotics. Trends in Microbiology, 1994, 2, 361-366.	7.7	189
25	Extensive re-modelling of the transpeptidase domain of penicillin-binding protein 2B of a penicillin-resistant South African isolate of <i>Streptococcus pneumoniae</i> . Molecular Microbiology, 1989, 3, 95-102.	2.5	176
26	Genetic analysis of clinical isolates of <i>Streptococcus pneumoniae</i> with high-level resistance to expanded-spectrum cephalosporins. Antimicrobial Agents and Chemotherapy, 1995, 39, 1306-1313.	3.2	171
27	Multilocus sequence typing of <i>Cronobacter sakazakii</i> and <i>Cronobacter malonaticus</i> reveals stable clonal structures with clinical significance which do not correlate with biotypes. BMC Microbiology, 2009, 9, 223.	3.3	165
28	Multilocus Sequence Typing for Comparison of Veterinary and Human Isolates of <i>Campylobacter jejuni</i> . Applied and Environmental Microbiology, 2003, 69, 6370-6379.	3.1	158
29	Multilocus Sequence Typing of Intercontinental Bovine <i>Staphylococcus aureus</i> Isolates. Journal of Clinical Microbiology, 2005, 43, 4737-4743.	3.9	158
30	Expanded Multilocus Sequence Typing for <i>Burkholderia</i> Species. Journal of Clinical Microbiology, 2009, 47, 2607-2610.	3.9	158
31	Genetic Diversity of the Streptococcal Competence (<i>com</i>) Gene Locus. Journal of Bacteriology, 1999, 181, 3144-3154.	2.2	145
32	Infection With Transmissible Strains of <i>Pseudomonas aeruginosa</i> and Clinical Outcomes in Adults With Cystic Fibrosis. JAMA - Journal of the American Medical Association, 2010, 304, 2145.	7.4	142
33	A novel multilocus sequence typing scheme for the opportunistic pathogen <i>Propionibacterium acnes</i> and characterization of type I cell surface-associated antigens. Microbiology (United Kingdom), 2011, 157, 1990-2003.	1.8	131
34	Multilocus Sequence Typing Identifies Evidence for Recombination and Two Distinct Lineages of <i>Corynebacterium diphtheriae</i> . Journal of Clinical Microbiology, 2010, 48, 4177-4185.	3.9	124
35	Repeated Botulinum Toxin Type A Injections for Refractory Overactive Bladder: Medium-Term Outcomes, Safety Profile, and Discontinuation Rates. European Urology, 2012, 61, 834-839.	1.9	120
36	Genetic Diversity of the tet (M) Gene in Tetracycline-Resistant Clonal Lineages of <i>Streptococcus pneumoniae</i> . Antimicrobial Agents and Chemotherapy, 2000, 44, 2979-2984.	3.2	117

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37	Genetics of high level penicillin resistance in clinical isolates of <i>Streptococcus pneumoniae</i> . FEMS Microbiology Letters, 1995, 126, 299-303.	1.8	116
38	Environmental <i>Burkholderia cepacia</i> Complex Isolates from Human Infections. Emerging Infectious Diseases, 2007, 13, 458-461.	4.3	112
39	Homeologous recombination and mismatch repair during transformation in <i>Streptococcus pneumoniae</i> : saturation of the Hex mismatch repair system.. Proceedings of the National Academy of Sciences of the United States of America, 1995, 92, 9052-9056.	7.1	101
40	Evidence for niche adaptation in the genome of the bovine pathogen <i>Streptococcus uberis</i> . BMC Genomics, 2009, 10, 54.	2.8	101
41	Insertion of an extra amino acid is the main cause of the low affinity of penicillin-binding protein 2 in penicillin-resistant strains of <i>Neisseria gonorrhoeae</i> . Molecular Microbiology, 1990, 4, 913-919.	2.5	99
42	Combating AMR: photoactivatable ruthenium(II)-isoniazid complex exhibits rapid selective antimycobacterial activity. Chemical Science, 2017, 8, 395-404.	7.4	99
43	Genetics and Molecular Biology of β -Lactam-Resistant Pneumococci. Microbial Drug Resistance, 1995, 1, 29-34.	2.0	97
44	Spontaneous sequence duplication within an open reading frame of the pneumococcal type 3 capsule locus causes high-frequency phase variation. Molecular Microbiology, 2002, 42, 1223-1232.	2.5	92
45	Biocide susceptibility of the <i>Burkholderia cepacia</i> complex. Journal of Antimicrobial Chemotherapy, 2009, 63, 502-510.	3.0	85
46	Sequence Control as a Powerful Tool for Improving the Selectivity of Antimicrobial Polymers. ACS Applied Materials & Interfaces, 2017, 9, 40117-40126.	8.0	83
47	Deletion analysis of the essentiality of penicillin-binding proteins 1A, 2B and 2X of <i>Streptococcus pneumoniae</i> . FEMS Microbiology Letters, 1993, 106, 171-175.	1.8	81
48	Resource relationships of foraging mycelial systems of <i>Phanerochaete velutina</i> and <i>Hypholoma fasciculare</i> in soil. New Phytologist, 1989, 111, 501-509.	7.3	79
49	Biguanide Iridium(III) Complexes with Potent Antimicrobial Activity. Journal of Medicinal Chemistry, 2018, 61, 7330-7344.	6.4	79
50	The Tetracycline Resistance Genetet(M) Exhibits Mosaic Structure. Plasmid, 1996, 35, 156-163.	1.4	78
51	IFN- γ Enhances Production of Nitric Oxide from Macrophages via a Mechanism That Depends on Nucleotide Oligomerization Domain-2. Journal of Immunology, 2006, 176, 4804-4810.	0.8	72
52	Nod1 Signaling Overcomes Resistance of <i>S. pneumoniae</i> to Opsonophagocytic Killing. PLoS Pathogens, 2007, 3, e118.	4.7	72
53	Characterization of tRNA-dependent Peptide Bond Formation by MurM in the Synthesis of <i>Streptococcus pneumoniae</i> Peptidoglycan. Journal of Biological Chemistry, 2008, 283, 6402-6417.	3.4	70
54	Penicillin-binding protein 2 genes of non- β -lactamase-producing, penicillin-resistant strains of <i>Neisseria gonorrhoeae</i> . Molecular Microbiology, 1989, 3, 35-41.	2.5	69

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55	Repeated Injections of Botulinum Toxin-A for Idiopathic Detrusor Overactivity. <i>Urology</i> , 2010, 75, 552-558.	1.0	69
56	Horizontal spread of an altered penicillin-binding protein 2B gene between <i>Streptococcus pneumoniae</i> and <i>Streptococcus oralis</i> . <i>FEMS Microbiology Letters</i> , 1993, 110, 335-339.	1.8	68
57	Distribution and Genetic Diversity of Sulysin in <i>Streptococcus suis</i> Isolated from Different Diseases of Pigs and Characterization of the Genetic Basis of Sulysin Absence. <i>Infection and Immunity</i> , 2001, 69, 7572-7582.	2.2	68
58	Spontaneous sequence duplications within capsule genes <i>cap8E</i> and <i>tts</i> control phase variation in <i>Streptococcus pneumoniae</i> serotypes 8 and 37. <i>Microbiology (United Kingdom)</i> , 2003, 149, 497-504.	1.8	67
59	Novel <i>Corynebacterium diphtheriae</i> in Domestic Cats. <i>Emerging Infectious Diseases</i> , 2010, 16, 688-691.	4.3	66
60	Genetic Analysis of Diverse Disease-Causing Pneumococci Indicates High Levels of Diversity within Serotypes and Capsule Switching. <i>Journal of Clinical Microbiology</i> , 2004, 42, 5681-5688.	3.9	65
61	VenUS II: a randomised controlled trial of larval therapy in the management of leg ulcers. <i>Health Technology Assessment</i> , 2009, 13, 1-182, iii-iv.	2.8	64
62	Incremental Increase in Fitness Cost with Increased β -Lactam Resistance in Pneumococci Evaluated by Competition in an Infant Rat Nasal Colonization Model. <i>Journal of Infectious Diseases</i> , 2006, 193, 1296-1303.	4.0	63
63	The form and outcome of mycelial interactions involving cord-forming decomposer basidiomycetes in homogeneous and heterogeneous environments. <i>New Phytologist</i> , 1988, 109, 423-432.	7.3	61
64	Nucleotide sequence of the penicillin-binding protein 2B gene of <i>Streptococcus pneumoniae</i> strain R6. <i>Nucleic Acids Research</i> , 1989, 17, 7518-7518.	14.5	61
65	Multilocus Sequence Typing Breathes Life into a Microbial Metagenome. <i>PLoS ONE</i> , 2006, 1, e17.	2.5	61
66	Inoculation of mycelial cord-forming basidiomycetes into woodland soil and litter II. Resource capture and persistence. <i>New Phytologist</i> , 1988, 109, 343-349.	7.3	59
67	Spatial dynamics and interactions of the woodland fairy ring fungus, <i>Clitocybe nebularis</i> . <i>New Phytologist</i> , 1989, 111, 699-705.	7.3	59
68	Inhibition of D-Ala:D-Ala ligase through a phosphorylated form of the antibiotic D-cycloserine. <i>Nature Communications</i> , 2017, 8, 1939.	12.8	59
69	Horizontal gene transfer and the evolution of resistance and virulence determinants in <i>Streptococcus</i> . <i>Journal of Applied Microbiology</i> , 1997, 83, 42S-51S.	3.1	58
70	Key Role for Efflux in the Preservative Susceptibility and Adaptive Resistance of <i>Burkholderia cepacia</i> Complex Bacteria. <i>Antimicrobial Agents and Chemotherapy</i> , 2013, 57, 2972-2980.	3.2	58
71	Genetics of oxacillin resistance in clinical isolates of <i>Streptococcus pneumoniae</i> that are oxacillin resistant and penicillin susceptible. <i>Antimicrobial Agents and Chemotherapy</i> , 1994, 38, 49-53.	3.2	57
72	Association between Hypermutator Phenotype, Clinical Variables, Mucoïd Phenotype, and Antimicrobial Resistance in <i>Pseudomonas aeruginosa</i> . <i>Journal of Clinical Microbiology</i> , 2008, 46, 3491-3493.	3.9	57

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73	Elucidating Global Epidemiology of <i>Burkholderia multivorans</i> in Cases of Cystic Fibrosis by Multilocus Sequence Typing. <i>Journal of Clinical Microbiology</i> , 2008, 46, 290-295.	3.9	57
74	NanA, a Neuraminidase from <i>Streptococcus pneumoniae</i> , Shows High Levels of Sequence Diversity, at Least in Part through Recombination with <i>Streptococcus oralis</i> . <i>Journal of Bacteriology</i> , 2005, 187, 5376-5386.	2.2	55
75	Population biology of <i>Streptococcus pneumoniae</i> isolated from oropharyngeal carriage and invasive disease The GenBank accession numbers for the sequences of the trpA/B alleles determined in this study are AF157817 to AF157826.. <i>Microbiology (United Kingdom)</i> , 1999, 145, 3283-3293.	1.8	54
76	A combined approach for comparative exoproteome analysis of <i>Corynebacterium pseudotuberculosis</i> . <i>BMC Microbiology</i> , 2011, 11, 12.	3.3	52
77	Molecular Evolution of Rifampicin Resistance in <i>Streptococcus pneumoniae</i> . <i>Microbial Drug Resistance</i> , 1998, 4, 65-70.	2.0	50
78	Distribution and Genetic Diversity of the ABC Transporter Lipoproteins PiuA and PiaA within <i>Streptococcus pneumoniae</i> and Related Streptococci. <i>Journal of Bacteriology</i> , 2006, 188, 1031-1038.	2.2	47
79	Molecular Characterization of Equine Isolates of <i>Streptococcus pneumoniae</i> : Natural Disruption of Genes Encoding the Virulence Factors Pneumolysin and Autolysin. <i>Infection and Immunity</i> , 1999, 67, 2776-2782.	2.2	44
80	The Autolysin-Encoding Gene (<i>lytA</i>) of <i>Streptococcus pneumoniae</i> Displays Restricted Allelic Variation despite Localized Recombination Events with Genes of Pneumococcal Bacteriophage Encoding Cell Wall Lytic Enzymes. <i>Infection and Immunity</i> , 1999, 67, 4551-4556.	2.2	44
81	Sansamycin natural product analogues as potent and selective anti-mycobacterials that inhibit lipid I biosynthesis. <i>Nature Communications</i> , 2017, 8, 14414.	12.8	43
82	Penicillin-binding protein 2b of <i>Streptococcus pneumoniae</i> in piperacillin-resistant laboratory mutants. <i>Journal of Bacteriology</i> , 1994, 176, 5574-5577.	2.2	41
83	Multilocus Sequence Typing of <i>Staphylococcus aureus</i> Isolated from High-Somatic-Cell-Count Cows and the Environment of an Organic Dairy Farm in the United Kingdom. <i>Journal of Clinical Microbiology</i> , 2005, 43, 4731-4736.	3.9	39
84	Inoculation of mycelial cord-forming basidiomycetes into woodland soil and litter I. Initial establishment. <i>New Phytologist</i> , 1988, 109, 335-341.	7.3	37
85	Investigating <i>Burkholderia cepacia</i> complex populations recovered from Italian maize rhizosphere by multilocus sequence typing. <i>Environmental Microbiology</i> , 2007, 9, 1632-1639.	3.8	35
86	Mutational Analysis of the Substrate Specificity of <i>Escherichia coli</i> Penicillin Binding Protein 4. <i>Biochemistry</i> , 2009, 48, 2675-2683.	2.5	35
87	In vitro characterization of the antivirulence target of Gram-positive pathogens, peptidoglycan O-acetyltransferase A (OatA). <i>PLoS Pathogens</i> , 2017, 13, e1006667.	4.7	35
88	Carbohydrate scaffolds as glycosyltransferase inhibitors with in vivo antibacterial activity. <i>Nature Communications</i> , 2015, 6, 7719.	12.8	34
89	Profiling interactions of vaborbactam with metallo- β -lactamases. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2019, 29, 1981-1984.	2.2	34
90	Specificity Determinants for Lysine Incorporation in <i>Staphylococcus aureus</i> Peptidoglycan as Revealed by the Structure of a MurE Enzyme Ternary Complex. <i>Journal of Biological Chemistry</i> , 2013, 288, 33439-33448.	3.4	33

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91	Automated Pneumococcal MLST Using Liquid-Handling Robotics and a Capillary DNA Sequencer. <i>Molecular Biotechnology</i> , 2003, 24, 303-308.	2.4	32
92	Fluorescent reagents for in vitro studies of lipid-linked steps of bacterial peptidoglycan biosynthesis: derivatives of UDPMurNAc-pentapeptide containing d-cysteine at position 4 or 5. <i>Molecular BioSystems</i> , 2006, 2, 484.	2.9	32
93	The safety and efficacy of botulinum toxin-A in the management of bladder oversensitivity: a randomised double-blind placebo-controlled trial. <i>International Journal of Clinical Practice</i> , 2011, 65, 698-704.	1.7	32
94	Novel and Improved Crystal Structures of <i>H. influenzae</i> , <i>E. coli</i> and <i>P. aeruginosa</i> Penicillin-Binding Protein 3 (PBP3) and <i>N. gonorrhoeae</i> PBP2: Toward a Better Understanding of β -Lactam Target-Mediated Resistance. <i>Journal of Molecular Biology</i> , 2019, 431, 3501-3519.	4.2	31
95	Oxidative Stress of <i>Burkholderia cenocepacia</i> Induces Insertion Sequence-Mediated Genomic Rearrangements That Interfere with Macrorestriction-Based Genotyping. <i>Journal of Clinical Microbiology</i> , 2010, 48, 34-40.	3.9	30
96	To Push or To Pull? In a Post-COVID World, Supporting and Incentivizing Antimicrobial Drug Development Must Become a Governmental Priority. <i>ACS Infectious Diseases</i> , 2021, 7, 2029-2042.	3.8	30
97	Increased pathogenicity of pneumococcal serotype 1 is driven by rapid autolysis and release of pneumolysin. <i>Nature Communications</i> , 2020, 11, 1892.	12.8	28
98	DNA methylation in fibrosis. <i>European Journal of Cell Biology</i> , 2016, 95, 323-330.	3.6	27
99	Touching proteins with virtual bare hands. <i>Journal of Computer-Aided Molecular Design</i> , 2018, 32, 703-709.	2.9	27
100	There is no market for new antibiotics: This allows an open approach to research and development. <i>Wellcome Open Research</i> , 2021, 6, 146.	1.8	27
101	Bayesian modeling of recombination events in bacterial populations. <i>BMC Bioinformatics</i> , 2008, 9, 421.	2.6	26
102	Investigating Bacteriophages Targeting the Opportunistic Pathogen <i>Acinetobacter baumannii</i> . <i>Antibiotics</i> , 2020, 9, 200.	3.7	26
103	<i>Pseudomonas aeruginosa</i> MurE amide ligase: enzyme kinetics and peptide inhibitor. <i>Biochemical Journal</i> , 2009, 421, 263-272.	3.7	25
104	Pneumolysin Detection Identifies Atypical Isolates of <i>Streptococcus pneumoniae</i> . <i>Journal of Clinical Microbiology</i> , 2000, 38, 1309-1310.	3.9	25
105	PiuA and PiaA, iron uptake lipoproteins of <i>Streptococcus pneumoniae</i> , elicit serotype independent antibody responses following human pneumococcal septicaemia. <i>FEMS Immunology and Medical Microbiology</i> , 2005, 43, 73-80.	2.7	24
106	Repeat botulinum toxin-A injections for treatment of adult detrusor overactivity. <i>Nature Reviews Urology</i> , 2010, 7, 661-667.	3.8	24
107	What is a Pneumococcus?. , 0, , 1-14.		24
108	Structural studies suggest aggregation as one of the modes of action for teixobactin. <i>Chemical Science</i> , 2018, 9, 8850-8859.	7.4	24

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109	Multilocus sequence types of invasive <i>Corynebacterium diphtheriae</i> isolated in the Rio de Janeiro urban area, Brazil. <i>Epidemiology and Infection</i> , 2012, 140, 617-620.	2.1	23
110	Development and extension of mycelial cords in soil at different temperatures and moisture contents. <i>Mycological Research</i> , 1989, 92, 383-391.	2.5	22
111	Distribution, Genetic Diversity, and Variable Expression of the Gene Encoding Hyaluronate Lyase within the <i>Streptococcus suis</i> Population. <i>Journal of Bacteriology</i> , 2004, 186, 4740-4747.	2.2	22
112	Cross-Sectional and Longitudinal Multilocus Sequence Typing of <i>Pseudomonas aeruginosa</i> in Cystic Fibrosis Sputum Samples. <i>Journal of Clinical Microbiology</i> , 2009, 47, 3444-3448.	3.9	22
113	Meeting the discovery challenge of drug-resistant infections: progress and focusing resources. <i>Drug Discovery Today</i> , 2019, 24, 452-461.	6.4	22
114	Substrate and Stereochemical Control of Peptidoglycan Cross-Linking by Transpeptidation by <i>Escherichia coli</i> PBP1B. <i>Journal of the American Chemical Society</i> , 2020, 142, 5034-5048.	13.7	21
115	Patient experience and satisfaction with Onabotulinumtoxin A for refractory overactive bladder. <i>BJU International</i> , 2015, 116, 443-449.	2.5	20
116	Concordance in diabetic foot ulceration: a cross-sectional study of agreement between wound swabbing and tissue sampling in infected ulcers. <i>Health Technology Assessment</i> , 2016, 20, 1-176.	2.8	20
117	A Role for Sigma Factor σ^E in <i>Corynebacterium pseudotuberculosis</i> Resistance to Nitric Oxide/Peroxide Stress. <i>Frontiers in Microbiology</i> , 2012, 3, 126.	3.5	19
118	Horizontal gene transfer and the evolution of resistance and virulence determinants in <i>Streptococcus</i> . <i>Society for Applied Bacteriology Symposium Series</i> , 1997, 26, 42S-51S.	0.4	17
119	Data reduction in headspace analysis of blood and urine samples for robust bacterial identification. <i>Computer Methods and Programs in Biomedicine</i> , 2005, 79, 259-271.	4.7	15
120	Adenosine phosphonate inhibitors of lipid II: Alanyl tRNA ligase MurM from <i>Streptococcus pneumoniae</i> . <i>Bioorganic and Medicinal Chemistry Letters</i> , 2007, 17, 4654-4656.	2.2	15
121	Diversity of the <i>parB</i> and <i>repA</i> genes of the <i>Burkholderia cepacia</i> complex and their utility for rapid identification of <i>Burkholderia cenocepacia</i> . <i>BMC Microbiology</i> , 2008, 8, 44.	3.3	15
122	High-Throughput Crystallography Reveals Boron-Containing Inhibitors of a Penicillin-Binding Protein with Di- and Trivalent Binding Modes. <i>Journal of Medicinal Chemistry</i> , 2021, 64, 11379-11394.	6.4	15
123	Outgrowth Patterns of Mycelial Cord-forming Basidiomycetes from and between Woody Resource Units in Soil. <i>Microbiology (United Kingdom)</i> , 1986, 132, 203-211.	1.8	14
124	Kinetic Characterization of Lipid II-Ala:Alanyl-tRNA Ligase (MurN) from <i>Streptococcus pneumoniae</i> using Semisynthetic Aminoacyl-lipid II Substrates. <i>Journal of Biological Chemistry</i> , 2008, 283, 34571-34579.	3.4	14
125	Evaluation of a Library of FDA-Approved Drugs for Their Ability To Potentiate Antibiotics against Multidrug-Resistant Gram-Negative Pathogens. <i>Antimicrobial Agents and Chemotherapy</i> , 2019, 63, .	3.2	14
126	Commercial Mushrooms and Bean Sprouts Are a Source of <i>Pseudomonas aeruginosa</i> . <i>Journal of Clinical Microbiology</i> , 2005, 43, 5830-5831.	3.9	13

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127	Inhibition of tRNA-dependent ligase MurM from <i>Streptococcus pneumoniae</i> by phosphonate and sulfonamide inhibitors. <i>Bioorganic and Medicinal Chemistry</i> , 2009, 17, 3443-3455.	3.0	13
128	Diaryltriazenes as antibacterial agents against methicillin resistant <i>Staphylococcus aureus</i> (MRSA) and <i>Mycobacterium smegmatis</i> . <i>European Journal of Medicinal Chemistry</i> , 2017, 127, 223-234.	5.5	13
129	Oxidation of Tertiary Amine-Derivatized Surfaces To Control Protein Adhesion. <i>Langmuir</i> , 2013, 29, 2961-2970.	3.5	12
130	Reliability of multilocus sequence typing of the <i>Burkholderia cepacia</i> complex in cystic fibrosis. <i>Journal of Cystic Fibrosis</i> , 2007, 6, 215-219.	0.7	11
131	Demonstration of the utility of DOS-derived fragment libraries for rapid hit derivatisation in a multidirectional fashion. <i>Chemical Science</i> , 2020, 11, 10792-10801.	7.4	11
132	Whole-Genome Sequence of <i>Corynebacterium pseudotuberculosis</i> Strain Cp162, Isolated from Camel. <i>Journal of Bacteriology</i> , 2012, 194, 5718-5719.	2.2	10
133	Adenosine Tetraphosphoadenosine Drives a Continuous ATP-Release Assay for Aminoacyl-tRNA Synthetases and Other Adenylate-Forming Enzymes. <i>ACS Chemical Biology</i> , 2013, 8, 2157-2163.	3.4	10
134	Reconstruction of diaminopimelic acid biosynthesis allows characterisation of <i>Mycobacterium tuberculosis</i> N-succinyl-L,L-diaminopimelic acid desuccinylase. <i>Scientific Reports</i> , 2016, 6, 23191.	3.3	10
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