Michel Doumith

List of Publications by Year in descending order

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47006 30087 11,262 109 47 103 citations h-index g-index papers 111 111 111 11842 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Emergence of a new antibiotic resistance mechanism in India, Pakistan, and the UK: a molecular, biological, and epidemiological study. Lancet Infectious Diseases, The, 2010, 10, 597-602.	9.1	2,485
2	Differentiation of the Major Listeria monocytogenes Serovars by Multiplex PCR. Journal of Clinical Microbiology, 2004, 42, 3819-3822.	3.9	915
3	The role of whole genome sequencing in antimicrobial susceptibility testing of bacteria: report from the EUCAST Subcommittee. Clinical Microbiology and Infection, 2017, 23, 2-22.	6.0	428
4	Molecular mechanisms disrupting porin expression in ertapenem-resistant Klebsiella and Enterobacter spp. clinical isolates from the UK. Journal of Antimicrobial Chemotherapy, 2009, 63, 659-667.	3.0	390
5	Phosphoethanolamine Modification of Lipid A in Colistin-Resistant Variants of Acinetobacter baumannii Mediated by the pmrAB Two-Component Regulatory System. Antimicrobial Agents and Chemotherapy, 2011, 55, 3370-3379.	3.2	354
6	New Aspects Regarding Evolution and Virulence of Listeria monocytogenes Revealed by Comparative Genomics and DNA Arrays. Infection and Immunity, 2004, 72, 1072-1083.	2.2	307
7	Identification of bacterial pathogens and antimicrobial resistance directly from clinical urines by nanopore-based metagenomic sequencing. Journal of Antimicrobial Chemotherapy, 2017, 72, 104-114.	3.0	296
8	What remains against carbapenem-resistant Enterobacteriaceae? Evaluation of chloramphenicol, ciprofloxacin, colistin, fosfomycin, minocycline, nitrofurantoin, temocillin and tigecycline. International Journal of Antimicrobial Agents, 2011, 37, 415-419.	2.5	292
9	Activity of aminoglycosides, including ACHN-490, against carbapenem-resistant Enterobacteriaceae isolates. Journal of Antimicrobial Chemotherapy, 2011, 66, 48-53.	3.0	252
10	Detection of the plasmid-mediated <i>mcr-1</i> gene conferring colistin resistance in human and food isolates of <i>Salmonella enterica </i> and <i>Escherichia coli</i> in England and Wales. Journal of Antimicrobial Chemotherapy, 2016, 71, 2300-2305.	3.0	247
11	Activities of NXL104 Combinations with Ceftazidime and Aztreonam against Carbapenemase-Producing <i>Enterobacteriaceae</i> . Antimicrobial Agents and Chemotherapy, 2011, 55, 390-394.	3.2	240
12	A Molecular Marker for Evaluating the Pathogenic Potential of FoodborneListeria monocytogenes. Journal of Infectious Diseases, 2004, 189, 2094-2100.	4.0	217
13	Methicillin-resistant Staphylococcus aureus emerged long before the introduction of methicillin into clinical practice. Genome Biology, 2017, 18, 130.	8.8	193
14	Plasmid Classification in an Era of Whole-Genome Sequencing: Application in Studies of Antibiotic Resistance Epidemiology. Frontiers in Microbiology, 2017, 8, 182.	3.5	191
15	Phylogenetic diversity of Escherichia coli strains producing NDM-type carbapenemases. Journal of Antimicrobial Chemotherapy, 2011, 66, 2002-2005.	3.0	168
16	Extended-spectrum Î ² -lactamase-producing Escherichia coli in human-derived and foodchain-derived samples from England, Wales, and Scotland: an epidemiological surveillance and typing study. Lancet Infectious Diseases, The, 2019, 19, 1325-1335.	9.1	150
17	Improved Multiplex PCR Strategy for Rapid Assignment of the Four Major Escherichia coli Phylogenetic Groups. Journal of Clinical Microbiology, 2012, 50, 3108-3110.	3.9	145
18	Prediction of Phenotypic Antimicrobial Resistance Profiles From Whole Genome Sequences of Non-typhoidal Salmonella enterica. Frontiers in Microbiology, 2018, 9, 592.	3 . 5	139

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19	Efflux Pumps, OprD Porin, AmpC \hat{l}^2 -Lactamase, and Multiresistance in <i>Pseudomonas aeruginosa</i> lsolates from Cystic Fibrosis Patients. Antimicrobial Agents and Chemotherapy, 2010, 54, 2219-2224.	3.2	130
20	AdeABC-mediated efflux and tigecycline MICs for epidemic clones of Acinetobacter baumannii. Journal of Antimicrobial Chemotherapy, 2010, 65, 1589-1593.	3.0	129
21	Diversity, virulence, and antimicrobial resistance of the KPC-producing Klebsiella pneumoniae ST307 clone. Microbial Genomics, 2017, 3, e000110.	2.0	122
22	Rapid Identification of Major Escherichia coli Sequence Types Causing Urinary Tract and Bloodstream Infections. Journal of Clinical Microbiology, 2015, 53, 160-166.	3.9	121
23	Whole-genome comparison of two Acinetobacter baumannii isolates from a single patient, where resistance developed during tigecycline therapy. Journal of Antimicrobial Chemotherapy, 2011, 66, 1499-1503.	3.0	96
24	Virulence genes in isolates of Klebsiella pneumoniae from the UK during 2016, including among carbapenemase gene-positive hypervirulent K1-ST23 and †non-hypervirulent†types ST147, ST15 and ST383. Journal of Medical Microbiology, 2018, 67, 118-128.	1.8	94
25	OXA-1 \hat{l}^2 -lactamase and non-susceptibility to penicillin/ \hat{l}^2 -lactamase inhibitor combinations among ESBL-producing < i>Escherichia coli < /i>. Journal of Antimicrobial Chemotherapy, 2019, 74, 326-333.	3.0	91
26	Diversity of carbapenem resistance mechanisms in Acinetobacter baumannii from a Taiwan hospital: spread of plasmid-borne OXA-72 carbapenemase. Journal of Antimicrobial Chemotherapy, 2009, 63, 641-647.	3.0	90
27	Ordering the mob: Insights into replicon and MOB typing schemes from analysis of a curated dataset of publicly available plasmids. Plasmid, 2017, 91, 42-52.	1.4	89
28	Isolation of fluoroquinolone-resistant O25b:H4-ST131 Escherichia coli with CTX-M-14 extended-spectrum Â-lactamase from UK river water. Journal of Antimicrobial Chemotherapy, 2011, 66, 512-516.	3.0	88
29	Pathogenomics of Listeria spp International Journal of Medical Microbiology, 2007, 297, 541-557.	3.6	84
30	Cephalosporin resistance mechanisms in Escherichia coli isolated from raw chicken imported into the UK. Journal of Antimicrobial Chemotherapy, 2010, 65, 2534-2537.	3.0	78
31	Variation in the genetic environments of blaCTX-M-15 in Escherichia coli from the faeces of travellers returning to the United Kingdom. Journal of Antimicrobial Chemotherapy, 2011, 66, 1005-1012.	3.0	76
32	OXA-48-like carbapenemases in the UK: an analysis of isolates and cases from 2007 to 2014. Journal of Antimicrobial Chemotherapy, 2017, 72, 1340-1349.	3.0	76
33	Characterization of plasmids encoding extended-spectrum Â-lactamases and their addiction systems circulating among Escherichia coli clinical isolates in the UK. Journal of Antimicrobial Chemotherapy, 2012, 67, 878-885.	3.0	74
34	Multicenter Validation of a Multiplex PCR Assay for Differentiating the Major Listeria monocytogenes Serovars 1/2a, 1/2b, 1/2c, and 4b: Toward an International Standard. Journal of Food Protection, 2005, 68, 2648-2650.	1.7	73
35	Clonal expansion of Escherichia coli ST38 carrying a chromosomally integrated OXA-48 carbapenemase gene. Journal of Medical Microbiology, 2016, 65, 538-546.	1.8	62
36	Livestock-Associated Methicillin Resistant Staphylococcus aureus (LA-MRSA) Clonal Complex (CC) 398 Isolated from UK Animals belong to European Lineages. Frontiers in Microbiology, 2016, 7, 1741.	3.5	61

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37	Comparison of phenotypic and WGS-derived antimicrobial resistance profiles of Shigella sonnei isolated from cases of diarrhoeal disease in England and Wales, 2015. Journal of Antimicrobial Chemotherapy, 2017, 72, 2496-2502.	3.0	61
38	Accuracy of Different Bioinformatics Methods in Detecting Antibiotic Resistance and Virulence Factors from Staphylococcus aureus Whole-Genome Sequences. Journal of Clinical Microbiology, 2018, 56, .	3.9	61
39	NDM carbapenemases in the United Kingdom: an analysis of the first 250 cases. Journal of Antimicrobial Chemotherapy, 2014, 69, 1777-1784.	3.0	59
40	WGS for surveillance of antimicrobial resistance: a pilot study to detect the prevalence and mechanism of resistance to azithromycin in a UK population of non-typhoidal <i>Salmonella</i> Journal of Antimicrobial Chemotherapy, 2016, 71, 3400-3408.	3.0	58
41	A curated dataset of complete Enterobacteriaceae plasmids compiled from the NCBI nucleotide database. Data in Brief, 2017, 12, 423-426.	1.0	58
42	Activity of ceftazidime/avibactam against problem Enterobacteriaceae and Pseudomonas aeruginosa in the UK, 2015–16. Journal of Antimicrobial Chemotherapy, 2018, 73, 648-657.	3.0	56
43	Emergence and clonal spread of colistin resistance due to multiple mutational mechanisms in carbapenemase-producing Klebsiella pneumoniae in London. Scientific Reports, 2017, 7, 12711.	3.3	55
44	Linkage of acquired quinolone resistance (qnrS1) and metallo-Â-lactamase (blaVIM-1) genes in multiple species of Enterobacteriaceae from Bolzano, Italy. Journal of Antimicrobial Chemotherapy, 2008, 61, 515-523.	3.0	54
45	Molecular epidemiology of fluoroquinolone-resistant ST131 Escherichia coli producing CTX-M extended-spectrum Â-lactamases in nursing homes in Belfast, UK. Journal of Antimicrobial Chemotherapy, 2011, 66, 297-303.	3.0	54
46	Interspecies complementation in Saccharopolyspora erythraea: elucidation of the function of oleP1, oleG1 and oleG2 from the oleandomycin biosynthetic gene cluster of Streptomyces antibioticus and generation of new erythromycin derivatives. Molecular Microbiology, 1999, 34, 1039-1048.	2.5	53
47	Molecular and epidemiological characterization of carbapenemase-producing Enterobacteriaceae in Norway, 2007 to 2014. PLoS ONE, 2017, 12, e0187832.	2.5	53
48	Antimicrobial resistance in Shiga toxin-producing <i>Escherichia coli</i> serogroups O157 and O26 isolated from human cases of diarrhoeal disease in England, 2015. Journal of Antimicrobial Chemotherapy, 2017, 72, 145-152.	3.0	51
49	High-Resolution Analysis by Whole-Genome Sequencing of an International Lineage (Sequence Type 111) of Pseudomonas aeruginosa Associated with Metallo-Carbapenemases in the United Kingdom. Journal of Clinical Microbiology, 2015, 53, 2622-2631.	3.9	50
50	Comparative virulence of urinary and bloodstream isolates of extra-intestinal pathogenic <i>Escherichia coli</i> in a <i>Galleria mellonella</i> model. Virulence, 2015, 6, 145-151.	4.4	50
51	Comparison of phenotypic and WGS-derived antimicrobial resistance profiles of Salmonella enterica serovars Typhi and Paratyphi. Journal of Antimicrobial Chemotherapy, 2018, 73, 365-372.	3.0	50
52	Real-time PCR for detection of the O25b-ST131 clone of Escherichia coli and its CTX-M-15-like extended-spectrum β-lactamases. International Journal of Antimicrobial Agents, 2010, 36, 355-358.	2.5	49
53	Association of Novel Nonsynonymous Single Nucleotide Polymorphisms in <i>ampD</i> with Cephalosporin Resistance and Phylogenetic Variations in <i>ampC</i> , <i>ampR</i> , <i>ompF</i> , and <i>ompC</i> in Enterobacter cloacae Isolates That Are Highly Resistant to Carbapenems. Antimicrobial Agents and Chemotherapy, 2016, 60, 2383-2390.	3.2	47
54	Major role of pKpQIL-like plasmids in the early dissemination of KPC-type carbapenemases in the UK. Journal of Antimicrobial Chemotherapy, 2017, 72, 2241-2248.	3.0	47

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55	Carbapenem-resistant Enterobacteriaceae dispersal from sinks is linked to drain position and drainage ratesAin a laboratory model system. Journal of Hospital Infection, 2019, 102, 63-69.	2.9	46
56	Population structure of <i>Escherichia coli </i> causing bacteraemia in the UK and Ireland between 2001 and 2010. Journal of Antimicrobial Chemotherapy, 2016, 71, 2139-2142.	3.0	45
57	False extended-spectrum \hat{l}^2 -lactamase phenotype in clinical isolates of Escherichia coli associated with increased expression of OXA-1 or TEM-1 penicillinases and loss of porins. Journal of Antimicrobial Chemotherapy, 2011, 66, 2006-2010.	3.0	44
58	An Investigation of the Diversity of Strains of Enteroaggregative Escherichia coli Isolated from Cases Associated with a Large Multi-Pathogen Foodborne Outbreak in the UK. PLoS ONE, 2014, 9, e98103.	2.5	41
59	Rise of multidrug-resistant non-vaccine serotype 15A Streptococcus pneumoniae in the United Kingdom, 2001 to 2014. Eurosurveillance, 2016, 21, .	7.0	41
60	Early (2008â€"2010) hospital outbreak of Klebsiella pneumoniae producing OXA-48 carbapenemase in the UK. International Journal of Antimicrobial Agents, 2013, 42, 531-536.	2.5	38
61	Comparison of phenotypic and WGS-derived antimicrobial resistance profiles of enteroaggregative Escherichia coli isolated from cases of diarrhoeal disease in England, 2015–16. Journal of Antimicrobial Chemotherapy, 2017, 72, 3288-3297.	3.0	38
62	Genetic diversity of Listeria monocytogenes recovered from infected persons and pork, seafood and dairy products on retail sale in France during 2000 and 2001. International Journal of Food Microbiology, 2007, 114, 187-194.	4.7	37
63	Emergence of AcrAB-mediated tigecycline resistance in a clinical isolate of Enterobacter cloacae during ciprofloxacin treatment. International Journal of Antimicrobial Agents, 2010, 35, 478-481.	2.5	36
64	Use of nrdA gene sequence clustering to estimate the prevalence of different Achromobacter species among Cystic Fibrosis patients in the UK. Journal of Cystic Fibrosis, 2016, 15, 479-485.	0.7	36
65	Metallo- \hat{l}^2 -lactamases among Enterobacteriaceae from routine samples in an Italian tertiary-care hospital and long-term care facilities during 2008. Clinical Microbiology and Infection, 2011, 17, 181-189.	6.0	35
66	IMI-2 carbapenemase in a clinical Klebsiella variicola isolated in the UK. Journal of Antimicrobial Chemotherapy, 2017, 72, 2129-2131.	3.0	34
67	Conformational flexibility and polymerization of vesicular stomatitis virus matrix protein. Journal of Molecular Biology, 1997, 274, 816-825.	4.2	33
68	Detection and molecular characterization of Livestock-Associated MRSA in raw meat on retail sale in North West England. Letters in Applied Microbiology, 2017, 64, 239-245.	2.2	33
69	Tigecycline resistance in Serratia marcescens associated with up-regulation of the SdeXY-HasF efflux system also active against ciprofloxacin and cefpirome. Journal of Antimicrobial Chemotherapy, 2010, 65, 479-482.	3.0	32
70	In vitro activity of rifaximin against clinical isolates of Escherichia coli and other enteropathogenic bacteria isolated from travellers returning to the UK. International Journal of Antimicrobial Agents, 2014, 43, 431-437.	2.5	31
71	Activity of \hat{I}^2 -lactam/taniborbactam (VNRX-5133) combinations against carbapenem-resistant Gram-negative bacteria. Journal of Antimicrobial Chemotherapy, 2021, 76, 160-170.	3.0	29
72	Selection of mutants with resistance or diminished susceptibility to ceftazidime/avibactam from ESBL-and AmpC-producing Enterobacteriaceae. Journal of Antimicrobial Chemotherapy, 2018, 73, 3336-3345.	3.0	26

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73	Genomic Epidemiology of Complex, Multispecies, Plasmid-Borne <i>bla</i> _{KPC} Carbapenemase in <i>Enterobacterales</i> in the United Kingdom from 2009 to 2014. Antimicrobial Agents and Chemotherapy, 2020, 64, .	3.2	26
74	Comparison of phenotypic and WGS-derived antimicrobial resistance profiles of Campylobacter jejuni and Campylobacter coli isolated from cases of diarrhoeal disease in England and Wales, 2015–16. Journal of Antimicrobial Chemotherapy, 2020, 75, 883-889.	3.0	26
75	Epidemiology of extended-spectrum beta-lactamase-producing Enterobacteriaceae in a UK district hospital; an observational study. Journal of Hospital Infection, 2012, 81, 270-277.	2.9	25
76	Trends in ExPEC serogroups in the UK and their significance. European Journal of Clinical Microbiology and Infectious Diseases, 2016, 35, 1661-1666.	2.9	25
77	Changes in colonization of residents and staff of a long-term care facility and an adjacent acute-care hospital geriatric unit by multidrug-resistant bacteria over a four-year period. Scandinavian Journal of Infectious Diseases, 2014, 46, 114-122.	1.5	24
78	Emergence of diversity in carbapenemase-producing Escherichia coli ST131, England, January 2014 to June 2016. Eurosurveillance, 2019, 24, .	7.0	24
79	First report of lukM-positive livestock-associated methicillin-resistant Staphylococcus aureus CC30 from fattening pigs in Northern Ireland. Veterinary Microbiology, 2016, 182, 131-134.	1.9	23
80	FRI-2 carbapenemase-producing Enterobacter cloacae complex in the UK. Journal of Antimicrobial Chemotherapy, 2017, 72, 2478-2482.	3.0	23
81	ISEcp1-mediated transposition of linked blaCTX-M-3 and blaTEM-1b from the IncI1 plasmid pEK204 found in clinical isolates of Escherichia coli from Belfast, UK. Journal of Antimicrobial Chemotherapy, 2011, 66, 2263-2265.	3.0	22
82	New insights into the regulatory pathways associated with the activation of the stringent response in bacterial resistance to the PBP2-targeted antibiotics, mecillinam and OP0595/RG6080. Journal of Antimicrobial Chemotherapy, 2016, 71, 2810-2814.	3.0	22
83	KPC enzymes in the UK: an analysis of the first 160 cases outside the North-West region. Journal of Antimicrobial Chemotherapy, 2016, 71, 1199-1206.	3.0	21
84	Cysteamine, an Endogenous Aminothiol, and Cystamine, the Disulfide Product of Oxidation, Increase Pseudomonas aeruginosa Sensitivity to Reactive Oxygen and Nitrogen Species and Potentiate Therapeutic Antibiotics against Bacterial Infection. Infection and Immunity, 2018, 86, .	2.2	21
85	Characterization of the extra-intestinal pathogenic Escherichia coli ST131 clone among isolates recovered from urinary and bloodstream infections in the United Kingdom. Journal of Medical Microbiology, 2015, 64, 1496-1503.	1.8	21
86	Carbapenemase-producing Salmonella enterica isolates in the UK. Journal of Antimicrobial Chemotherapy, 2015, 70, 2165-7.	3.0	20
87	Successful treatment of infective endocarditis due to pandrug-resistant Klebsiella pneumoniae with ceftazidime-avibactam and aztreonam. Scientific Reports, 2021, 11, 9684.	3.3	20
88	Integration of Genomic and Other Epidemiologic Data to Investigate and Control a Cross-Institutional Outbreak of <i>Streptococcus pyogenes </i> Emerging Infectious Diseases, 2016, 22, 973-980.	4.3	18
89	Genomic analysis of the first KPC-producing Klebsiella pneumoniae isolated from a patient in Riyadh: A new public health concern in Saudi Arabia. Journal of Infection and Public Health, 2020, 13, 647-650.	4.1	17
90	Clonal expansion of community-associated meticillin-resistant Staphylococcus aureus (MRSA) in people who inject drugs (PWID): prevalence, risk factors and molecular epidemiology, Bristol, United Kingdom, 2012 to 2017. Eurosurveillance, 2019, 24, .	7.0	17

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91	Carbapenem resistance mediated byblaOXA-181inPseudomonas aeruginosa. Journal of Antimicrobial Chemotherapy, 2016, 71, 2056-2057.	3.0	16
92	The characterization of mobile colistin resistance (mcr) genes among 33 000 Salmonella enterica genomes from routine public health surveillance in England. Microbial Genomics, 2020, 6, .	2.0	16
93	The differential importance of mutations within AmpD in cephalosporin resistance of Enterobacter aerogenes and Enterobacter cloacae. International Journal of Antimicrobial Agents, 2016, 48, 555-558.	2.5	15
94	S. Enteritidis and S. Typhimurium Harboring SPI-1 and SPI-2 Are the Predominant Serotypes Associated With Human Salmonellosis in Saudi Arabia. Frontiers in Cellular and Infection Microbiology, 2019, 9, 187.	3.9	15
95	Genomic Characterization of Carbapenem-Non-susceptible Pseudomonas aeruginosa Clinical Isolates From Saudi Arabia Revealed a Global Dissemination of GES-5-Producing ST235 and VIM-2-Producing ST233 Sub-Lineages. Frontiers in Microbiology, 2021, 12, 765113.	3.5	13
96	Metallo-Î ² -lactamases among Enterobacteriaceae from routine samples in an Italian tertiary care hospital and long-term care facilities during 2008. Clinical Microbiology and Infection, 2010, 17, 181-9.	6.0	12
97	Use of DNA arrays for the analysis of outbreak-related strains of Listeria monocytogenes. International Journal of Medical Microbiology, 2006, 296, 559-562.	3.6	11
98	First outbreak of colonization by linezolid- and glycopeptide-resistant Enterococcus faecium harbouring the cfr gene in a UK nephrology unit. Journal of Hospital Infection, 2017, 97, 397-402.	2.9	11
99	Genomic sequences of Streptococcus agalactiae with high-level gentamicin resistance, collected in the BSAC bacteraemia surveillance. Journal of Antimicrobial Chemotherapy, 2017, 72, 2704-2707.	3.0	11
100	Investigation of a hospital Enterobacter cloacae NDM-1 outbreak using whole genome sequencing. Access Microbiology, 2019, 1, .	0.5	11
101	Diversity of carbapenemase-producing Enterobacterales in England as revealed by whole-genome sequencing of isolates referred to a national reference laboratory over a 30-month period. Journal of Medical Microbiology, 2022, 71, .	1.8	10
102	Complete Genome Sequence of a Colistin-Resistant Uropathogenic Escherichia coli Sequence Type 131 <i>fimH</i> 22 Strain Harboring <i>mcr-1</i> on an IncHI2 Plasmid, Isolated in Riyadh, Saudi Arabia. Microbiology Resource Announcements, 2019, 8, .	0.6	8
103	Acquired resistome and plasmid sequencing of $\langle i\rangle$ mcr $\langle i\rangle$ - $\langle i\rangle$ 1 $\langle i\rangle$ carrying MDR Enterobacteriaceae from poultry and their relationship to STs associated with humans. JAC-Antimicrobial Resistance, 2022, 4, dlab198.	2.1	6
104	Invasive fungal infection of the brain caused by Neoscytalidium dimidiatum in a post-renal transplant patient: A case report. Medical Mycology Case Reports, 2021, 34, 27-31.	1.3	5
105	Activity of RX-04 Pyrrolocytosine Protein Synthesis Inhibitors against Multidrug-Resistant Gram-Negative Bacteria. Antimicrobial Agents and Chemotherapy, 2018, 62, .	3.2	3
106	OXA-48 carbapenemase-producing Salmonella enterica serovar Kentucky ST198 isolated from Saudi Arabia. Journal of Antimicrobial Chemotherapy, 2020, 75, 2006-2008.	3.0	2
107	Integration of Genomic and Other Epidemiologic Data to Investigate and Control a Cross-Institutional Outbreak of <i>Streptococcus pyogenes </i> <ir> <ir> <ir> Integration of Genomic and Other Epidemiologic Data to Investigate and Control a Cross-Institutional Diseases, 2016, 22, 973-980.</ir></ir></ir>	4.3	2
108	Genotypes of Listeria monocytogenes strains isolated from 2000 to 2002 in Poland. Polish Journal of Microbiology, 2006, 55, 31-5.	1.7	2

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109	Molecular and Phenotypic Properties of Multidrug-Resistant Extraintestinal Pathogenic Escherichia coli (ExPEC) From Clinical Samples. Open Forum Infectious Diseases, 2015, 2, .	0.9	0