## Emmanuelle Cambau

List of Publications by Year in descending order

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133 papers 9,860 citations

45 h-index 95 g-index

139 all docs 139 docs citations

139 times ranked 9382 citing authors

#	Article	lF	CITATIONS
1	A Diarylquinoline Drug Active on the ATP Synthase of Mycobacterium tuberculosis. Science, 2005, 307, 223-227.	12.6	1,907
2	Mycobacterium abscessus: a new antibiotic nightmare. Journal of Antimicrobial Chemotherapy, 2012, 67, 810-818.	3.0	597
3	Treatment of Nontuberculous Mycobacterial Pulmonary Disease: An Official ATS/ERS/ESCMID/IDSA Clinical Practice Guideline. Clinical Infectious Diseases, 2020, 71, e1-e36.	<b>5.</b> 8	367
4	Treatment of Nontuberculous Mycobacterial Pulmonary Disease: An Official ATS/ERS/ESCMID/IDSA Clinical Practice Guideline. Clinical Infectious Diseases, 2020, 71, 905-913.	5 <b>.</b> 8	357
5	Treatment of nontuberculous mycobacterial pulmonary disease: an official ATS/ERS/ESCMID/IDSA clinical practice guideline. European Respiratory Journal, 2020, 56, 2000535.	6.7	336
6	Temporal Trends in Infective Endocarditis in the Context of Prophylaxis Guideline Modifications. Journal of the American College of Cardiology, 2012, 59, 1968-1976.	2.8	327
7	Sixty-three Cases of Mycobacterium marinum Infection. Archives of Internal Medicine, 2002, 162, 1746.	3.8	324
8	Assessment of Clarithromycin Susceptibility in Strains Belonging to the <i>Mycobacterium abscessus</i> Group by <i>erm</i> (41) and <i>rrl</i> Sequencing. Antimicrobial Agents and Chemotherapy, 2011, 55, 775-781.	<b>3.</b> 2	291
9	A systematic review of gyrase mutations associated with fluoroquinolone-resistant Mycobacterium tuberculosis and a proposed gyrase numbering system. Journal of Antimicrobial Chemotherapy, 2012, 67, 819-831.	3.0	221
10	Mycobacterium tuberculosis DNA Gyrase: Interaction with Quinolones and Correlation with Antimycobacterial Drug Activity. Antimicrobial Agents and Chemotherapy, 2004, 48, 1281-1288.	3.2	217
11	Novel Gyrase Mutations in Quinolone-Resistant and -Hypersusceptible Clinical Isolates of Mycobacterium tuberculosis : Functional Analysis of Mutant Enzymes. Antimicrobial Agents and Chemotherapy, 2006, 50, 104-112.	3.2	176
12	Treatment outcome definitions in nontuberculous mycobacterial pulmonary disease: an NTM-NET consensus statement. European Respiratory Journal, 2018, 51, 1800170.	6.7	159
13	Evaluation of a New Test, GenoType HelicoDR, for Molecular Detection of Antibiotic Resistance in <i>Helicobacter pylori</i> Journal of Clinical Microbiology, 2009, 47, 3600-3607.	3.9	151
14	Mycobacterium tuberculosis drug-resistance testing: challenges, recent developments and perspectives. Clinical Microbiology and Infection, 2017, 23, 154-160.	6.0	150
15	Multilocus Sequence Analysis and <i>rpoB</i> Sequencing of Mycobacterium abscessus (Sensu Lato) Strains. Journal of Clinical Microbiology, 2011, 49, 491-499.	3.9	137
16	Genetic Basis for Natural and Acquired Resistance to the Diarylquinoline R207910 in Mycobacteria. Antimicrobial Agents and Chemotherapy, 2006, 50, 2853-2856.	<b>3.</b> 2	125
17	Type II Topoisomerase Mutations in Ciprofloxacin-Resistant Strains of <i>Pseudomonas aeruginosa</i> Antimicrobial Agents and Chemotherapy, 1999, 43, 62-66.	3.2	120
18	Antimicrobial resistance in leprosy: results of the first prospective open survey conducted by a WHO surveillance network for the period 2009–15. Clinical Microbiology and Infection, 2018, 24, 1305-1310.	6.0	113

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19	Correlation between Quinolone Susceptibility Patterns and Sequences in the A and B Subunits of DNA Gyrase in Mycobacteria. Antimicrobial Agents and Chemotherapy, 1998, 42, 2084-2088.	3.2	106
20	Antibiotic Susceptibility Pattern of <i>Mycobacterium marinum</i> . Antimicrobial Agents and Chemotherapy, 2000, 44, 3133-3136.	<b>3.2</b>	105
21	Steps towards the discovery of Mycobacterium tuberculosis by Robert Koch, 1882. Clinical Microbiology and Infection, 2014, 20, 196-201.	6.0	101
22	First Evidence of Amoebae–Mycobacteria Association in Drinking Water Network. Environmental Science & Environmental Science	10.0	99
23	<i>Mycobacterium marinum</i> . Microbiology Spectrum, 2017, 5, .	3.0	92
24	Update on fluoroquinolone resistance in Helicobacter pylori: new mutations leading to resistance and first description of a gyrA polymorphism associated with hypersusceptibility. International Journal of Antimicrobial Agents, 2007, 29, 389-396.	2.5	90
25	In Vivo Evaluation of Antibiotic Activity Against Mycobacterium abscessus. Journal of Infectious Diseases, 2014, 209, 905-912.	4.0	89
26	Epidemic and pandemic viral infections: impact on tuberculosis and the lung. European Respiratory Journal, 2020, 56, 2001727.	6.7	89
27	First functional characterization of a singly expressed bacterial type II topoisomerase: The enzyme from Mycobacterium tuberculosis. Biochemical and Biophysical Research Communications, 2006, 348, 158-165.	2.1	87
28	Multidrug-resistance to dapsone, rifampicin, and ofloxacin in Mycobacterium leprae. Lancet, The, 1997, 349, 103-104.	13.7	86
29	Comparing Mycobacterium massiliense and Mycobacterium abscessus lung infections in cystic fibrosis patients. Journal of Cystic Fibrosis, 2015, 14, 63-69.	0.7	80
30	Molecular Detection of Rifampin and Ofloxacin Resistance for Patients Who Experience Relapse of Multibacillary Leprosy. Clinical Infectious Diseases, 2002, 34, 39-45.	5.8	75
31	Classification Algorithm for Subspecies Identification within the Mycobacterium abscessus Species, Based on Matrix-Assisted Laser Desorption Ionization–Time of Flight Mass Spectrometry. Journal of Clinical Microbiology, 2014, 52, 3362-3369.	3.9	75
32	Impact of Low-Level Resistance to Fluoroquinolones Due to <i>qnrA1</i> and <i>qnrS1</i> Genes or a <i>gyrA</i> Mutation on Ciprofloxacin Bactericidal Activity in a Murine Model of <i>Escherichia coli</i> Urinary Tract Infection. Antimicrobial Agents and Chemotherapy, 2009, 53, 4292-4297.	3.2	71
33	Target specificity of the new fluoroquinolone besifloxacin in Streptococcus pneumoniae, Staphylococcus aureus and Escherichia coli. Journal of Antimicrobial Chemotherapy, 2009, 63, 443-450.	3.0	70
34	Clinical Management of Rapidly Growing Mycobacterial Cutaneous Infections in Patients after Mesotherapy. Clinical Infectious Diseases, 2009, 49, 1358-1364.	5.8	66
35	Cardiac surgery during the acute phase of infective endocarditis: discrepancies between European Society of Cardiology guidelines and practices. European Heart Journal, 2016, 37, 840-848.	2.2	64
36	Outbreak of Nontuberculous Mycobacterial Subcutaneous Infections Related to Multiple Mesotherapy Injections. Journal of Clinical Microbiology, 2009, 47, 1961-1964.	3.9	63

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37	Infections caused by (i) Mycobacterium abscessus (i): epidemiology, diagnostic tools and treatment. Expert Review of Anti-Infective Therapy, 2016, 14, 1139-1154.	4.4	63
38	Design, synthesis and activity against Toxoplasma gondii, Plasmodium spp., and Mycobacterium tuberculosis of new 6-fluoroquinolones. European Journal of Medicinal Chemistry, 2006, 41, 1478-1493.	5 <b>.</b> 5	62
39	Rapid detection of qnr and qepA plasmid-mediated quinolone resistance genes using real-time PCR. Diagnostic Microbiology and Infectious Disease, 2011, 70, 253-259.	1.8	58
40	Low selection of topoisomerase mutants from strains of Escherichia coli harbouring plasmid-borne qnr genes. Journal of Antimicrobial Chemotherapy, 2008, 61, 1007-1015.	3.0	57
41	Standardized interpretation of antibiotic susceptibility testing and resistance genotyping for i>Mycobacterium abscessus / i>with regard to subspecies and i>erm41 / i>sequevar. Journal of Antimicrobial Chemotherapy, 2016, 71, 2208-2212.	3.0	54
42	Selection of Resistance to Clarithromycin in Mycobacterium abscessus Subspecies. Antimicrobial Agents and Chemotherapy, 2017, 61, .	3.2	52
43	The Pentapeptide Repeat Proteins MfpA <sub>Mt</sub> and QnrB4 Exhibit Opposite Effects on DNA Gyrase Catalytic Reactions and on the Ternary Gyrase-DNA-Quinolone Complex. Journal of Bacteriology, 2009, 191, 1587-1594.	2.2	51
44	Consensus management recommendations for less common non-tuberculous mycobacterial pulmonary diseases. Lancet Infectious Diseases, The, 2022, 22, e178-e190.	9.1	51
45	Multilocus sequence typing scheme for the Mycobacterium abscessus complex. Research in Microbiology, 2014, 165, 82-90.	2.1	49
46	Antimicrobial susceptibility testing of Mycobacterium tuberculosis complex isolates – the EUCAST broth microdilution reference method for MIC determination. Clinical Microbiology and Infection, 2020, 26, 1488-1492.	6.0	49
47	Identification of Mycobacterial Species by PCR Sequencing of Quinolone Resistance-Determining Regions of DNA Gyrase Genes. Journal of Clinical Microbiology, 2003, 41, 1311-1315.	3.9	48
48	Comparison of Culture Methods for Isolation of Nontuberculous Mycobacteria from Surface Waters. Applied and Environmental Microbiology, 2010, 76, 3514-3520.	3.1	48
49	Selective reporting of antibiotic susceptibility test results in European countries: an ESCMID cross-sectional survey. International Journal of Antimicrobial Agents, 2017, 49, 162-166.	2.5	48
50	Dihydropteroate Synthase Mutations in the folP1 Gene Predict Dapsone Resistance in Relapsed Cases of Leprosy. Clinical Infectious Diseases, 2006, 42, 238-241.	5.8	47
51	Functional Analysis of DNA Gyrase Mutant Enzymes Carrying Mutations at Position 88 in the A Subunit Found in Clinical Strains of Mycobacterium tuberculosis Resistant to Fluoroquinolones. Antimicrobial Agents and Chemotherapy, 2006, 50, 4170-4173.	3.2	45
52	Evaluation of the new GenoType NTM-DR kit for the molecular detection of antimicrobial resistance in non-tuberculous mycobacteria. Journal of Antimicrobial Chemotherapy, 2017, 72, 1669-1677.	3.0	44
53	Type II topoisomerase mutations in clinical isolates of Enterobacter cloacae and other enterobacterial species harbouring the qnrA gene. International Journal of Antimicrobial Agents, 2007, 29, 402-409.	2.5	43
54	Amplification and nucleotide sequence of the quinolone resistance-determining region in thegyrAgene of mycobacteria. FEMS Microbiology Letters, 1994, 116, 49-54.	1.8	40

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55	Mobile Insertion Cassette Elements Found in Small Non-Transmissible Plasmids in Proteeae May Explain qnrD Mobilization. PLoS ONE, 2014, 9, e87801.	2.5	40
56	A Plasmid-Borne <i>Shewanella algae</i> Gene, <i>qnrA3</i> , and Its Possible Transfer In Vivo between <i>Kluyvera ascorbata</i> and <i>Klebsiella pneumoniae</i> Journal of Bacteriology, 2008, 190, 5217-5223.	2.2	38
57	Absence of <i>Mycobacterium tuberculosis</i> in Arterial Lesions from Patients with Takayasu's Arteritis. Journal of Rheumatology, 2009, 36, 1682-1685.	2.0	38
58	atpE gene as a new useful specific molecular target to quantify Mycobacterium in environmental samples. BMC Microbiology, 2013, 13, 277.	3.3	38
59	<i>Mycobacterium</i> Behavior in Wastewater Treatment Plant, A Bacterial Model Distinct From <i>Escherichia coli</i> and Enterococci. Environmental Science & Environmental Sc	10.0	37
60	Detection of Antibiotic Resistance in Leprosy Using GenoType LepraeDR, a Novel Ready-To-Use Molecular Test. PLoS Neglected Tropical Diseases, 2012, 6, e1739.	3.0	37
61	Impact of highly active antiretroviral therapy on onset of Mycobacterium avium complex infection and cytomegalovirus disease in patients with AIDS. Aids, 2000, 14, 2593-2596.	2.2	36
62	An enhanced regimen as post-exposure chemoprophylaxis for leprosy: PEP++. BMC Infectious Diseases, 2018, 18, 506.	2.9	36
63	Comparison of Saramis 4.12 and IVD 3.0 Vitek MS Matrix-Assisted Laser Desorption Ionization–Time of Flight Mass Spectrometry for Identification of Mycobacteria from Solid and Liquid Culture Media. Journal of Clinical Microbiology, 2017, 55, 2045-2054.	3.9	35
64	Ciprofloxacin Treatment Failure in a Murine Model of Pyelonephritis Due to an AAC(6′)-lb-cr-Producing Escherichia coli Strain Susceptible to Ciprofloxacin ⟨i⟩In Vitro⟨ i⟩. Antimicrobial Agents and Chemotherapy, 2013, 57, 5830-5835.	3.2	34
65	Molecular epidemiology and mechanisms of resistance of azithromycin-resistantNeisseria gonorrhoeaeisolated in France during 2013–14. Journal of Antimicrobial Chemotherapy, 2016, 71, 2471-2478.	3.0	34
66	Description of a 2,683-Base-Pair Plasmid Containing <i>qnrD</i> in Two Providencia rettgeri Isolates. Antimicrobial Agents and Chemotherapy, 2012, 56, 565-568.	3.2	33
67	Human infections due to nontuberculous mycobacteria: the infectious diseases and clinical microbiology specialists' point of view. Future Microbiology, 2015, 10, 1467-1483.	2.0	33
68	Development of a Real-Time qPCR Method for Detection and Enumeration of <i>Mycobacterium</i> spp. in Surface Water. Applied and Environmental Microbiology, 2010, 76, 7348-7351.	3.1	32
69	Diversity of Individual Dynamic Patterns of Emergence of Resistance to Quinolones in Escherichia coli From the Fecal Flora of Healthy Volunteers Exposed to Ciprofloxacin. Journal of Infectious Diseases, 2012, 206, 1399-1406.	4.0	31
70	Negligible risk of inducing resistance in Mycobacterium tuberculosis with single-dose rifampicin as post-exposure prophylaxis for leprosy. Infectious Diseases of Poverty, 2016, 5, 46.	3.7	31
71	Purification and inhibition by quinolones of DNA gyrases from Mycobacterium avium, Mycobacterium smegmatis and Mycobacterium fortuitum bv. peregrinum. Microbiology (United Kingdom), 1999, 145, 2527-2532.	1.8	30
72	Relation between presence of extended-spectrum $\hat{l}^2$ -lactamase-producing Enterobacteriaceae in systematic rectal swabs and respiratory tract specimens in ICU patients. Annals of Intensive Care, 2017, 7, 13.	4.6	29

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73	In vivo selection during ofloxacin therapy of Escherichia coli with combined topoisomerase mutations that confer high resistance to ofloxacin but susceptibility to nalidixic acid. Journal of Antimicrobial Chemotherapy, 2006, 58, 1054-1057.	3.0	28
74	Evaluation of the New MB Redox System for Detection of Growth of Mycobacteria. Journal of Clinical Microbiology, 1999, 37, 2013-2015.	3.9	27
75	Are All the DNA Gyrase Mutations Found in Mycobacterium leprae Clinical Strains Involved in Resistance to Fluoroquinolones?. Antimicrobial Agents and Chemotherapy, 2008, 52, 745-747.	3.2	26
76	Performance and economic evaluation of the molecular detection of pathogens for patients with severe infections: the EVAMICA open-label, cluster-randomised, interventional crossover trial. Intensive Care Medicine, 2017, 43, 1613-1625.	8.2	26
77	Plasmidic qnrA3 Enhances Escherichia coli Fitness in Absence of Antibiotic Exposure. PLoS ONE, 2011, 6, e24552.	2.5	26
78	Expression and Purification of an Active Form of the Mycobacterium leprae DNA Gyrase and Its Inhibition by Quinolones. Antimicrobial Agents and Chemotherapy, 2007, 51, 1643-1648.	3.2	25
79	Diagnostic criteria for urinary tract infection in hospitalized elderly patients over 75 years of age: A multicenter cross-sectional study. Médecine Et Maladies Infectieuses, 2013, 43, 189-194.	5.0	25
80	Mutagenesis in the $\hat{l}\pm3\hat{l}\pm4$ GyrA Helix and in the Toprim Domain of GyrB Refines the Contribution of <i>Mycobacterium tuberculosis</i> DNA Gyrase to Intrinsic Resistance to Quinolones. Antimicrobial Agents and Chemotherapy, 2008, 52, 2909-2914.	3.2	24
81	Ceftriaxone-Resistant Neisseria gonorrhoeae Isolates (2010 to 2014) in France Characterized by Using Whole-Genome Sequencing. Antimicrobial Agents and Chemotherapy, 2016, 60, 6962-6964.	3.2	24
82	High-resolution melting analysis for rapid characterization of qnr alleles in clinical isolates and detection of two novel alleles, qnrB25 and qnrB42. Journal of Antimicrobial Chemotherapy, 2012, 67, 2635-2639.	3.0	22
83	Trend of plasmid-mediated quinolone resistance genes at the Children's Hospital in Tunisia. Journal of Medical Microbiology, 2014, 63, 195-202.	1.8	21
84	Determination of Escherichia coli phylogroups in elderly patients with urinary tract infection or asymptomatic bacteriuria. Clinical Microbiology and Infection, 2019, 25, 839-844.	6.0	21
85	Outbreak in a haematology unit involving an unusual strain of glycopeptide-resistant Enterococcus faecium carrying both vanA and vanB genes. Journal of Antimicrobial Chemotherapy, 2014, 69, 500-505.	3.0	19
86	Clinical Utility of an Amplification Test Based on Ligase Chain Reaction in Pulmonary Tuberculosis. American Journal of Respiratory and Critical Care Medicine, 1998, 158, 1096-1101.	5.6	18
87	Genetic Diversity and Population Structure of Mycobacterium marinum: New Insights into Host and Environmental Specificities. Journal of Clinical Microbiology, 2012, 50, 3627-3634.	3.9	18
88	A case of postoperative breast infection by Mycobacterium fortuitum associated with the hospital water supply. American Journal of Infection Control, 2015, 43, 406-408.	2.3	18
89	Benefits of Polymerase Chain Reaction Combined With Culture for the Diagnosis of Bone and Joint Infections: A Prospective Test Performance Study. Open Forum Infectious Diseases, 2019, 6, ofz511.	0.9	18
90	Untreated highly viraemic pregnant women from Asia or subâ€Saharan Africa often transmit hepatitis B virus despite serovaccination to newborns. Liver International, 2015, 35, 409-416.	3.9	17

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91	Prosthetic knee arthritis due to Granulicatella adiacens after dental treatment. Journal of Medical Microbiology, 2013, 62, 1624-1627.	1.8	16
92	New Insights into the Geographic Distribution of Mycobacterium leprae SNP Genotypes Determined for Isolates from Leprosy Cases Diagnosed in Metropolitan France and French Territories. PLoS Neglected Tropical Diseases, 2015, 9, e0004141.	3.0	15
93	Mycobacterium llatzerense, a waterborne Mycobacterium, that resists phagocytosis by Acanthamoeba castellanii. Scientific Reports, 2017, 7, 46270.	3.3	15
94	Bacterial biofilm in adenoids of children with chronic otitis media. Part I: a case control study of prevalence of biofilms in adenoids, risk factors and middle ear biofilms. Acta Oto-Laryngologica, 2019, 139, 345-350.	0.9	15
95	Population Genomics of Mycobacterium leprae Reveals a New Genotype in Madagascar and the Comoros. Frontiers in Microbiology, 2020, 11, 711.	3.5	15
96	What is the role of the EUCAST reference method for MIC testing of the Mycobacterium tuberculosis complex?. Clinical Microbiology and Infection, 2020, 26, 1453-1455.	6.0	14
97	Clonal Relationship and Differentiation among Mycobacterium abscessus Isolates as Determined Using the Semiautomated Repetitive Extragenic Palindromic Sequence PCR-Based DiversiLab System. Journal of Clinical Microbiology, 2014, 52, 1969-1977.	3.9	11
98	Use of Andromas and Bruker MALDI-TOF MS in the identification of Neisseria. European Journal of Clinical Microbiology and Infectious Diseases, 2018, 37, 2273-2277.	2.9	11
99	A systematic review of Mycobacterium leprae DNA gyrase mutations and their impact on fluoroquinolone resistance. Clinical Microbiology and Infection, 2021, 27, 1601-1612.	6.0	11
100	Assessing Primary and Secondary Resistance to Clarithromycin and Amikacin in Infections Due to Mycobacterium avium Complex. Antimicrobial Agents and Chemotherapy, 2015, 59, 7153-7155.	3.2	10
101	Current mentorship practices in the training of the next generation of clinical microbiology and infectious disease specialists: an international cross-sectional survey. European Journal of Clinical Microbiology and Infectious Diseases, 2019, 38, 659-665.	2.9	10
102	Association of Healthcare and Aesthetic Procedures with Infections Caused by Nontuberculous Mycobacteria, France, 2012â€'2020. Emerging Infectious Diseases, 2022, 28, 518-526.	4.3	10
103	Multicentre testing of the EUCAST broth microdilution reference method for MIC determination on Mycobacterium tuberculosis. Clinical Microbiology and Infection, 2021, 27, 288.e1-288.e4.	6.0	9
104	Investigating drug resistance of Mycobacterium leprae in the Comoros: an observational deep-sequencing study. Lancet Microbe, The, 2022, 3, e693-e700.	7.3	9
105	First detection of plasmid-mediated quinolone resistance in the community setting and in hospitalized patients in Switzerland. Journal of Antimicrobial Chemotherapy, 2008, 62, 1151-1152.	3.0	8
106	Infective Endocarditis Related to Unusual Microorganisms: A Prospective Population-Based Study. Open Forum Infectious Diseases, 2020, 7, ofaa127.	0.9	8
107	Disseminated Mycobacterium chimaera Following Open-Heart Surgery, the Heater–Cooler Unit Worldwide Outbreak: Case Report and Minireview. Frontiers in Medicine, 2020, 7, 243.	2.6	8
108	Prevalence of Oropharyngeal Candidiasis in Geriatric Inpatients. Journal of the American Geriatrics Society, 2001, 49, 1741-1742.	2.6	7

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109	â€~Antimicrobial resistance in leprosy: results of the first prospective open survey conducted by a WHO surveillance network for the period 2009–2015' – Author's reply. Clinical Microbiology and Infection, 2019, 25, 646-647.	6.0	7
110	Antimicrobials that affect the synthesis and conformation of nucleic acids. OIE Revue Scientifique Et Technique, 2012, 31, 77-87.	1.2	7
111	Impact of a 24/7 multiplex-PCR on the management of patients with confirmed viral meningitis. Journal of Infection, 2021, 83, 650-655.	3.3	7
112	In vivo selection of a complex mutant TEM (CMT) from an inhibitor-resistant TEM (IRT) during ceftazidime therapy. Journal of Antimicrobial Chemotherapy, 2013, 68, 2792-2796.	3.0	6
113	A new workflow for the microbiological diagnosis of febrile neutropenia in patients with a central venous catheter. Journal of Antimicrobial Chemotherapy, 2013, 68, 943-946.	3.0	6
114	Xpert GBS Assay for Rapid Detection of Group B Streptococcus in Gastric Fluid Samples from Newborns. Journal of Clinical Microbiology, 2014, 52, 657-659.	3.9	6
115	A Case of Fluoroquinolone-Resistant Leprosy Discovered after 9 Years of Misdiagnosis. Case Reports in Infectious Diseases, 2016, 2016, 1-4.	0.5	6
116	Mycobacterium marinum., 0,, 735-752.		6
117	The inÂvitro mechanisms of isoniazid and ethionamide resistance poorly reflect those inÂvivo in Mycobacterium tuberculosis. Tuberculosis, 2016, 101, 144-145.	1.9	5
118	First genetic characterisation of multidrug-resistant Mycobacterium tuberculosis isolates from Algeria. Journal of Global Antimicrobial Resistance, 2019, 19, 301-307.	2.2	5
119	Management of Tuberculosis: Are the Practices Homogeneous in High-Income Countries?. Frontiers in Public Health, 2020, 8, 443.	2.7	5
120	<i>Mycobacterium chimaera</i> Genomics With Regard to Epidemiological and Clinical Investigations Conducted for an Open Chest Postsurgical <i>Mycobacterium chimaera</i> Infection Outbreak. Open Forum Infectious Diseases, 2021, 8, ofab192.	0.9	5
121	Management of patients with pulmonary mycobacteriosis in France: a multicenter retrospective cohort study. BMC Pulmonary Medicine, 2021, 21, 333.	2.0	5
122	Multi-pathogen real-time PCR system adds benefit for my patients: yes. Intensive Care Medicine, 2015, 41, 528-530.	8.2	4
123	<i>qnrA6</i> genetic environment and quinolone resistance conferred on <i>Proteus mirabilis</i> Journal of Antimicrobial Chemotherapy, 2016, 71, 903-908.	3.0	4
124	Bacterial biofilm in adenoids of children with chronic otitis media. Part II: a case–control study of nasopharyngeal microbiota, virulence, and resistance of biofilms in adenoids. Acta Oto-Laryngologica, 2020, 140, 220-224.	0.9	4
125	Contribution of the ATP Binding Site of ParE to Susceptibility to Novobiocin and Quinolones in Streptococcus pneumoniae. Journal of Bacteriology, 2005, 187, 1536-1540.	2.2	3
126	Association of the 16S rRNA methylase gene rmtB with a novel insertion sequence element belonging to the IS L3 family. International Journal of Antimicrobial Agents, 2017, 49, 117-118.	2.5	3

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127	Native joint septic arthritis due to Clostridium tarantellae. Anaerobe, 2019, 56, 46-48.	2.1	3
128	Inadequate Therapeutic Response to a Recommended Antituberculosis Fixed-Dose Combination Regimen in an Overweight Patient with <i>Mycobacterium bovis</i> Infection. Annals of Pharmacotherapy, 2013, 47, e4-e4.	1.9	1
129	Visualizing viable Mycobacterium tuberculosis in sputum to monitor isolation measures. Journal of Infection, 2017, 74, 207-210.	3.3	1
130	Cause Analysis of an Infection in Facelift Surgery Due to Mycobacterium chelonae. Frontiers in Medicine, 2019, 6, 243.	2.6	1
131	Decreased susceptibility to cephalosporins among gonococci?. Lancet Infectious Diseases, The, 2014, 14, 184-185.	9.1	O
132	Mycobactéries et eau. Revue Francophone Des Laboratoires, 2014, 2014, 69-74.	0.0	0
133	Impact of a 24/7 Rapid Molecular Assay for Influenza Detection on the Prescription of Oseltamivir. Open Forum Infectious Diseases, 2020, 7, ofaa374.	0.9	0