Emmanuelle Cambau

List of Publications by Year
in descending order

133 papers	9,860 citations	53794 45 h-index	38395 95 g-index
139 all docs	139 docs citations	139 times ranked	9382 citing authors

#	Article	IF	Citations
1	Consensus management recommendations for less common non-tuberculous mycobacterial pulmonary diseases. Lancet Infectious Diseases, The, 2022, 22, e178-e190.	9.1	51
2	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
3	Investigating drug resistance of Mycobacterium leprae in the Comoros: an observational deep-sequencing study. Lancet Microbe, The, 2022, 3, e693-e700.	7.3	9
4	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
5	Mycobacterium chimaera Genomics With Regard to Epidemiological and Clinical Investigations Conducted for an Open Chest Postsurgical Mycobacterium chimaera Infection Outbreak. Open Forum Infectious Diseases, 2021, 8, ofab192.	0.9	5
6	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
7	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
8	Management of patients with pulmonary mycobacteriosis in France: a multicenter retrospective cohort study. BMC Pulmonary Medicine, 2021, 21, 333.	2.0	5
9	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
10	Antimicrobial susceptibility testing of Mycobacterium tuberculosis complex isolates â€“ the EUCAST broth microdilution reference method for MIC determination. Clinical Microbiology and Infection, 2020, 26, 1488-1492.	6.0	49
11	Management of Tuberculosis: Are the Practices Homogeneous in High-Income Countries?. Frontiers in Public Health, 2020, 8, 443.	2.7	5
12	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
13	Treatment of Nontuberculous Mycobacterial Pulmonary Disease: An Official ATS/ERS/ESCMID/IDSA Clinical Practice Guideline. Clinical Infectious Diseases, 2020, 71, 905-913.	5.8	357
14	Infective Endocarditis Related to Unusual Microorganisms: A Prospective Population-Based Study. Open Forum Infectious Diseases, 2020, 7, ofaa127.	0.9	8
15	Population Genomics of Mycobacterium leprae Reveals a New Genotype in Madagascar and the Comoros. Frontiers in Microbiology, 2020, 11, 711.	3.5	15
16	Disseminated Mycobacterium chimaera Following Open-Heart Surgery, the Heaterâ€"Cooler Unit Worldwide Outbreak: Case Report and Minireview. Frontiers in Medicine, 2020, 7, 243.	2.6	8
17	Treatment of nontuberculous mycobacterial pulmonary disease: an official ATS/ERS/ESCMID/IDSA clinical practice guideline. European Respiratory Journal, 2020, 56, 2000535.	6.7	336
18	Epidemic and pandemic viral infections: impact on tuberculosis and the lung. European Respiratory Journal, 2020, 56, 2001727.	6.7	89

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19	Treatment of Nontuberculous Mycobacterial Pulmonary Disease: An Official ATS/ERS/ESCMID/IDSA Clinical Practice Guideline. Clinical Infectious Diseases, 2020, 71, e1-e36.	5.8	367
20	Bacterial biofilm in adenoids of children with chronic otitis media. Part II: a caseâ€“control study of nasopharyngeal microbiota, virulence, and resistance of biofilms in adenoids. Acta Oto-Laryngologica, 2020, 140, 220-224.	0.9	4
21	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
22	â€˜Antimicrobial resistance in leprosy: results of the first prospective open survey conducted by a WHO surveillance network for the period 2009â€“2015â€™ â€“ Author's reply. Clinical Microbiology and Infection, 2019, 25, 646-647.	6.0	7
23	First genetic characterisation of multidrug-resistant Mycobacterium tuberculosis isolates from Algeria. Journal of Global Antimicrobial Resistance, 2019, 19, 301-307.	2.2	5
24	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
25	Native joint septic arthritis due to Clostridium tarantellae. Anaerobe, 2019, 56, 46-48.	2.1	3
26	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
27	Cause Analysis of an Infection in Facelift Surgery Due to Mycobacterium chelonae. Frontiers in Medicine, 2019, 6, 243.	2.6	1
28	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
29	Antimicrobial resistance in leprosy: results of the first prospective open survey conducted by a WHO surveillance network for the period 2009â€“15. Clinical Microbiology and Infection, 2018, 24, 1305-1310.	6.0	113
30	Treatment outcome definitions in nontuberculous mycobacterial pulmonary disease: an NTM-NET consensus statement. European Respiratory Journal, 2018, 51, 1800170.	6.7	159
31	An enhanced regimen as post-exposure chemoprophylaxis for leprosy: PEP++. BMC Infectious Diseases, 2018, 18, 506.	2.9	36
32	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
33	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
34	Relation between presence of extended-spectrum Î²-lactamase-producing Enterobacteriaceae in systematic rectal swabs and respiratory tract specimens in ICU patients. Annals of Intensive Care, 2017, 7, 13.	4.6	29
35	Mycobacterium llatzerense, a waterborne Mycobacterium, that resists phagocytosis by Acanthamoeba castellanii. Scientific Reports, 2017, 7, 46270.	3.3	15
36	Comparison of Saramis 4.12 and IVD 3.0 Vitek MS Matrix-Assisted Laser Desorption Ionizationâ€“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

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37	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
38	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
39	Mycobacterium marinum. Microbiology Spectrum, 2017, 5, .	3.0	92
40	Visualizing viable Mycobacterium tuberculosis in sputum to monitor isolation measures. Journal of Infection, 2017, 74, 207-210.	3.3	1
41	Selection of Resistance to Clarithromycin in Mycobacterium abscessus Subspecies. Antimicrobial Agents and Chemotherapy, 2017, 61, .	3.2	52
42	Mycobacterium tuberculosis drug-resistance testing: challenges, recent developments and perspectives. Clinical Microbiology and Infection, 2017, 23, 154-160.	6.0	150
43	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
44	A Case of Fluoroquinolone-Resistant Leprosy Discovered after 9 Years of Misdiagnosis. Case Reports in Infectious Diseases, 2016, 2016, 1-4.	0.5	6
45	Standardized interpretation of antibiotic susceptibility testing and resistance genotyping forMycobacterium abscessuswith regard to subspecies anderm41sequevar. Journal of Antimicrobial Chemotherapy, 2016, 71, 2208-2212.	3.0	54
46	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
47	The inÂvitro mechanisms of isoniazid and ethionamide resistance poorly reflect those inÂvivo in Mycobacterium tuberculosis. Tuberculosis, 2016, 101, 144-145.	1.9	5
48	Infections caused byMycobacterium abscessus: epidemiology, diagnostic tools and treatment. Expert Review of Anti-Infective Therapy, 2016, 14, 1139-1154.	4.4	63
49	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
50	Molecular epidemiology and mechanisms of resistance of azithromycin-resistantNeisseria gonorrhoeaeisolated in France during 2013â€“14. Journal of Antimicrobial Chemotherapy, 2016, 71, 2471-2478.	3.0	34
51	qnrA6genetic environment and quinolone resistance conferred onProteus mirabilis. Journal of Antimicrobial Chemotherapy, 2016, 71, 903-908.	3.0	4
52	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
53	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
54	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

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55	Multi-pathogen real-time PCR system adds benefit for my patients: yes. Intensive Care Medicine, 2015, 41, 528-530.	8.2	4
56	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
57	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
58	Human infections due to nontuberculous mycobacteria: the infectious diseases and clinical microbiology specialistsâ€™ point of view. Future Microbiology, 2015, 10, 1467-1483.	2.0	33
59	Comparing Mycobacterium massiliense and Mycobacterium abscessus lung infections in cystic fibrosis patients. Journal of Cystic Fibrosis, 2015, 14, 63-69.	0.7	80
60	Steps towards the discovery of Mycobacterium tuberculosis by Robert Koch, 1882. Clinical Microbiology and Infection, 2014, 20, 196-201.	6.0	101
61	First Evidence of Amoebaeâ€“Mycobacteria Association in Drinking Water Network. Environmental Science & Technology, 2014, 48, 11872-11882.	10.0	99
62	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
63	Decreased susceptibility to cephalosporins among gonococci?. Lancet Infectious Diseases, The, 2014, 14, 184-185.	9.1	0
64	Multilocus sequence typing scheme for the Mycobacterium abscessus complex. Research in Microbiology, 2014, 165, 82-90.	2.1	49
65	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
66	Trend of plasmid-mediated quinolone resistance genes at the Childrenâ€™s Hospital in Tunisia. Journal of Medical Microbiology, 2014, 63, 195-202.	1.8	21
67	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
68	In Vivo Evaluation of Antibiotic Activity Against Mycobacterium abscessus. Journal of Infectious Diseases, 2014, 209, 905-912.	4.0	89
69	Classification Algorithm for Subspecies Identification within the Mycobacterium abscessus Species, Based on Matrix-Assisted Laser Desorption Ionizationâ€“Time of Flight Mass Spectrometry. Journal of Clinical Microbiology, 2014, 52, 3362-3369.	3.9	75
70	MycobactÃ©ries et eau. Revue Francophone Des Laboratoires, 2014, 2014, 69-74.	0.0	0
71	Mobile Insertion Cassette Elements Found in Small Non-Transmissible Plasmids in Proteeae May Explain qnrD Mobilization. PLoS ONE, 2014, 9, e87801.	2.5	40
72	Diagnostic criteria for urinary tract infection in hospitalized elderly patients over 75 years of age: A multicenter cross-sectional study. MÃ©decine Et Maladies Infectieuses, 2013, 43, 189-194.	5.0	25

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73	atpE gene as a new useful specific molecular target to quantify Mycobacterium in environmental samples. BMC Microbiology, 2013, 13, 277.	3.3	38
74	Prosthetic knee arthritis due to Granulicatella adiacens after dental treatment. Journal of Medical Microbiology, 2013, 62, 1624-1627.	1.8	16
75	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
76	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
77	Inadequate Therapeutic Response to a Recommended Antituberculosis Fixed-Dose Combination Regimen in an Overweight Patient with Mycobacterium bovis Infection. Annals of Pharmacotherapy, 2013, 47, e4-e4.	1.9	1
78	Ciprofloxacin Treatment Failure in a Murine Model of Pyelonephritis Due to an AAC(6â€²)-Ib-cr-Producing Escherichia coli Strain Susceptible to Ciprofloxacin In Vitro. Antimicrobial Agents and Chemotherapy, 2013, 57, 5830-5835.	3.2	34
79	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
80	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
81	Description of a 2,683-Base-Pair Plasmid Containing qnrD in Two Providencia rettgeri Isolates. Antimicrobial Agents and Chemotherapy, 2012, 56, 565-568.	3.2	33
82	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
83	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
84	Mycobacterium abscessus: a new antibiotic nightmare. Journal of Antimicrobial Chemotherapy, 2012, 67, 810-818.	3.0	597
85	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
86	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
87	Antimicrobials that affect the synthesis and conformation of nucleic acids. OIE Revue Scientifique Et Technique, 2012, 31, 77-87.	1.2	7
88	Mycobacterium Behavior in Wastewater Treatment Plant, A Bacterial Model Distinct From Escherichia coli and Enterococci. Environmental Science & Technology, 2011, 45, 5380-5386.	10.0	37
89	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
90	Assessment of Clarithromycin Susceptibility in Strains Belonging to the Mycobacterium abscessus Group by erm (41) and rrl Sequencing. Antimicrobial Agents and Chemotherapy, 2011, 55, 775-781.	3.2	291

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91	Multilocus Sequence Analysis and rpoB Sequencing of Mycobacterium abscessus (Sensu Lato) Strains. Journal of Clinical Microbiology, 2011, 49, 491-499.	3.9	137
92	Plasmidic qnrA3 Enhances Escherichia coli Fitness in Absence of Antibiotic Exposure. PLoS ONE, 2011, 6, e24552.	2.5	26
93	Comparison of Culture Methods for Isolation of Nontuberculous Mycobacteria from Surface Waters. Applied and Environmental Microbiology, 2010, 76, 3514-3520.	3.1	48
94	Development of a Real-Time qPCR Method for Detection and Enumeration of Mycobacterium spp. in Surface Water. Applied and Environmental Microbiology, 2010, 76, 7348-7351.	3.1	32
95	Absence ofMycobacterium tuberculosisin Arterial Lesions from Patients with Takayasuâ€™s Arteritis. Journal of Rheumatology, 2009, 36, 1682-1685.	2.0	38
96	Evaluation of a New Test, GenoType HelicoDR, for Molecular Detection of Antibiotic Resistance inHelicobacter pylori. Journal of Clinical Microbiology, 2009, 47, 3600-3607.	3.9	151
97	Outbreak of Nontuberculous Mycobacterial Subcutaneous Infections Related to Multiple Mesotherapy Injections. Journal of Clinical Microbiology, 2009, 47, 1961-1964.	3.9	63
98	The Pentapeptide Repeat Proteins MfpA _{Mt} 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
99	Impact of Low-Level Resistance to Fluoroquinolones Due to qnrA1 and qnrS1 Genes or a gyrA Mutation on Ciprofloxacin Bactericidal Activity in a Murine Model of Escherichia coli Urinary Tract Infection. Antimicrobial Agents and Chemotherapy, 2009, 53, 4292-4297.	3.2	71
100	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
101	Clinical Management of Rapidly Growing Mycobacterial Cutaneous Infections in Patients after Mesotherapy. Clinical Infectious Diseases, 2009, 49, 1358-1364.	5.8	66
102	Mutagenesis in the Î±3Î±4 GyrA Helix and in the Toprim Domain of GyrB Refines the Contribution of Mycobacterium tuberculosis DNA Gyrase to Intrinsic Resistance to Quinolones. Antimicrobial Agents and Chemotherapy, 2008, 52, 2909-2914.	3.2	24
103	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
104	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
105	A Plasmid-Borne Shewanella algae Gene, qnrA3 , and Its Possible Transfer In Vivo between Kluyvera ascorbata and Klebsiella pneumoniae. Journal of Bacteriology, 2008, 190, 5217-5223.	2.2	38
106	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
107	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
108	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

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109	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
110	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
111	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.	5.5	62
112	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
113	Genetic Basis for Natural and Acquired Resistance to the Diarylquinoline R207910 in Mycobacteria. Antimicrobial Agents and Chemotherapy, 2006, 50, 2853-2856.	3.2	125
114	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
115	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
116	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
117	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
118	A Diarylquinoline Drug Active on the ATP Synthase of Mycobacterium tuberculosis. Science, 2005, 307, 223-227.	12.6	1,907
119	Mycobacterium tuberculosis DNA Gyrase: Interaction with Quinolones and Correlation with Antimycobacterial Drug Activity. Antimicrobial Agents and Chemotherapy, 2004, 48, 1281-1288.	3.2	217
120	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
121	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
122	Sixty-three Cases of Mycobacterium marinum Infection. Archives of Internal Medicine, 2002, 162, 1746.	3.8	324
123	Prevalence of Oropharyngeal Candidiasis in Geriatric Inpatients. Journal of the American Geriatrics Society, 2001, 49, 1741-1742.	2.6	7
124	Antibiotic Susceptibility Pattern of Mycobacterium marinum. Antimicrobial Agents and Chemotherapy, 2000, 44, 3133-3136.	3.2	105
125	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
126	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

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127	Type II Topoisomerase Mutations in Ciprofloxacin-Resistant Strains of Pseudomonas aeruginosa. Antimicrobial Agents and Chemotherapy, 1999, 43, 62-66.	3.2	120
128	Evaluation of the New MB Redox System for Detection of Growth of Mycobacteria. Journal of Clinical Microbiology, 1999, 37, 2013-2015.	3.9	27
129	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
130	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
131	Multidrug-resistance to dapsone, rifampicin, and ofloxacin in Mycobacterium leprae. Lancet, The, 1997, 349, 103-104.	13.7	86
132	Amplification and nucleotide sequence of the quinolone resistance-determining region in thegyrAgene of mycobacteria. FEMS Microbiology Letters, 1994, 116, 49-54.	1.8	40
133	Mycobacterium marinum. , 0, , 735-752.		6