## Surbhi Malhotra-Kumar

List of Publications by Year in descending order

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122 papers 11,990 citations

35 h-index 30081 103 g-index

133 all docs

133
docs citations

times ranked

133

15012 citing authors

#	Article	IF	CITATIONS
1	Discovery, research, and development of new antibiotics: the WHO priority list of antibiotic-resistant bacteria and tuberculosis. Lancet Infectious Diseases, The, 2018, 18, 318-327.	9.1	3,672
2	ResFinder 4.0 for predictions of phenotypes from genotypes. Journal of Antimicrobial Chemotherapy, 2020, 75, 3491-3500.	3.0	1,523
3	Methicillin-resistant Staphylococcus aureus. Nature Reviews Disease Primers, 2018, 4, 18033.	30.5	833
4	Identification of a novel plasmid-mediated colistin-resistance gene, mcr-2, in Escherichia coli, Belgium, June 2016. Eurosurveillance, 2016, 21, .	7.0	648
5	Effect of azithromycin and clarithromycin therapy on pharyngeal carriage of macrolide-resistant streptococci in healthy volunteers: a randomised, double-blind, placebo-controlled study. Lancet, The, 2007, 369, 482-490.	13.7	465
6	Multiplex PCR for detection of plasmid-mediated colistin resistance determinants, mcr-1, mcr-2, mcr-3, mcr-4 and mcr-5 for surveillance purposes. Eurosurveillance, 2018, 23, .	7.0	431
7	European Surveillance of Antimicrobial Consumption (ESAC): outpatient antibiotic use in Europe. Journal of Antimicrobial Chemotherapy, 2006, 58, 401-407.	3.0	336
8	Interventions to reduce colonisation and transmission of antimicrobial-resistant bacteria in intensive care units: an interrupted time series study and cluster randomised trial. Lancet Infectious Diseases, The, 2014, 14, 31-39.	9.1	297
9	Surveillance for control of antimicrobial resistance. Lancet Infectious Diseases, The, 2018, 18, e99-e106.	9.1	235
10	In vivo and In vitro Interactions between Pseudomonas aeruginosa and Staphylococcus spp Frontiers in Cellular and Infection Microbiology, 2017, 7, 106.	3.9	193
11	Colistin resistance gene mcr-1 harboured on a multidrug resistant plasmid. Lancet Infectious Diseases, The, 2016, 16, 283-284.	9.1	153
12	Multiplex PCR for Simultaneous Detection of Macrolide and Tetracycline Resistance Determinants in Streptococci. Antimicrobial Agents and Chemotherapy, 2005, 49, 4798-4800.	3.2	144
13	MLST reveals potentially high-risk international clones of Enterobacter cloacae*. Journal of Antimicrobial Chemotherapy, 2015, 70, 48-56.	3.0	131
14	Decontamination Strategies and Bloodstream Infections With Antibiotic-Resistant Microorganisms in Ventilated Patients. JAMA - Journal of the American Medical Association, 2018, 320, 2087.	7.4	127
15	Colistin-resistant Escherichia coli harbouring mcr-1 isolated from food animals in Hanoi, Vietnam. Lancet Infectious Diseases, The, 2016, 16, 286-287.	9.1	109
16	Current Trends in Rapid Diagnostics for Methicillin-Resistant <i>Staphylococcus aureus </i> and Glycopeptide-Resistant <i>Enterococcus </i> Species. Journal of Clinical Microbiology, 2008, 46, 1577-1587.	3.9	107
17	Proposal for assignment of allele numbers for mobile colistin resistance (mcr) genes. Journal of Antimicrobial Chemotherapy, 2018, 73, 2625-2630.	3.0	101
18	Current Trends in Culture-Based and Molecular Detection of Extended-Spectrum-Â-Lactamase-Harboring and Carbapenem-Resistant Enterobacteriaceae. Journal of Clinical Microbiology, 2012, 50, 1140-1146.	3.9	98

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19	Consolidating and Exploring Antibiotic Resistance Gene Data Resources. Journal of Clinical Microbiology, 2016, 54, 851-859.	3.9	94
20	European Surveillance of Antimicrobial Consumption (ESAC): outpatient quinolone use in Europe. Journal of Antimicrobial Chemotherapy, 2006, 58, 423-427.	3.0	90
21	The effects of antibiotic cycling and mixing on antibiotic resistance in intensive care units: a cluster-randomised crossover trial. Lancet Infectious Diseases, The, 2018, 18, 401-409.	9.1	65
22	Complete sequence of an IncFII plasmid harbouring the colistin resistance gene <i>mcr-1</i> isolated from Belgian pig farms. Journal of Antimicrobial Chemotherapy, 2016, 71, 2342-2344.	3.0	63
23	Evaluation of colistin stability in agar and comparison of four methods for MIC testing of colistin. European Journal of Clinical Microbiology and Infectious Diseases, 2018, 37, 345-353.	2.9	61
24	Effect of Amoxicillin Dose and Treatment Duration on the Need for Antibiotic Re-treatment in Children With Community-Acquired Pneumonia. JAMA - Journal of the American Medical Association, 2021, 326, 1713.	7.4	57
25	Evaluation of Chromogenic Media for Detection of Methicillin-Resistant <i>Staphylococcus aureus </i> . Journal of Clinical Microbiology, 2010, 48, 1040-1046.	3.9	56
26	Addressing the challenges in antisepsis: focus on povidone iodine. International Journal of Antimicrobial Agents, 2020, 56, 106064.	2.5	56
27	European Surveillance of Antimicrobial Consumption (ESAC): outpatient macrolide, lincosamide and streptogramin (MLS) use in Europe. Journal of Antimicrobial Chemotherapy, 2006, 58, 418-422.	3.0	51
28	The endotracheal tube microbiome associated with Pseudomonas aeruginosa or Staphylococcus epidermidis. Scientific Reports, 2016, 6, 36507.	3.3	51
29	Comparison of strategies to reduce meticillin-resistant <i>Staphylococcus aureus</i> rates in surgical patients: a controlled multicentre intervention trial. BMJ Open, 2013, 3, e003126.	1.9	49
30	Effect of outpatient antibiotics for urinary tract infections on antimicrobial resistance among commensal Enterobacteriaceae: a multinational prospective cohort study. Clinical Microbiology and Infection, 2018, 24, 972-979.	6.0	49
31	Characterization of carbapenemase-producing Enterobacteriaceae from colonized patients in a university hospital in Madrid, Spain, during the R-GNOSIS project depicts increased clonal diversity over time with maintenance of high-risk clones. Journal of Antimicrobial Chemotherapy, 2018, 73, 3039-3043.	3.0	47
32	Rapid evolution and host immunity drive the rise and fall of carbapenem resistance during an acute Pseudomonas aeruginosa infection. Nature Communications, 2021, 12, 2460.	12.8	47
33	Comparison of Biofilm Formation between Major Clonal Lineages of Methicillin Resistant Staphylococcus aureus. PLoS ONE, 2014, 9, e104561.	2.5	43
34	Microbiota-based markers predictive of development of Clostridioides difficile infection. Nature Communications, 2021, 12, 2241.	12.8	40
35	Active liquid degassing in microfluidic systems. Lab on A Chip, 2013, 13, 4366.	6.0	38
36	Colistin-Resistant Acinetobacter baumannii Clinical Strains with Deficient Biofilm Formation. Antimicrobial Agents and Chemotherapy, 2016, 60, 1892-1895.	3.2	38

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37	Clonal spread of fluoroquinolone non-susceptible Streptococcus pyogenes. Journal of Antimicrobial Chemotherapy, 2005, 55, 320-325.	3.0	37
38	Macrolide- and Telithromycin-resistant <i>Streptococcus pyogenes</i> , Belgium, 1999–2003 <u></u> , lemerging Infectious Diseases, 2005, 11, 939-942.	4.3	34
39	Estimating the association between antibiotic exposure and colonization with extended-spectrum $\hat{l}^2$ -lactamase-producing Gram-negative bacteria using machine learning methods: a multicentre, prospective cohort study. Clinical Microbiology and Infection, 2020, 26, 87-94.	6.0	34
40	A Lateral Flow Immunoassay for the Rapid Identification of CTX-M-Producing Enterobacterales from Culture Plates and Positive Blood Cultures. Diagnostics, 2020, 10, 764.	2.6	33
41	The global dissemination of hospital clones of Enterococcus faecium. Genome Medicine, 2021, 13, 52.	8.2	33
42	An <i>In Vitro</i> Deletion in <i>ribE</i> Encoding Lumazine Synthase Contributes to Nitrofurantoin Resistance in Escherichia coli. Antimicrobial Agents and Chemotherapy, 2014, 58, 7225-7233.	3.2	32
43	Integrated DNA and RNA extraction and purification on an automated microfluidic cassette from bacterial and viral pathogens causing community-acquired lower respiratory tract infections. Lab on A Chip, 2014, 14, 1519-1526.	6.0	32
44	Antimicrobial Drug Use and Macrolide-Resistant <i>Streptococcus pyogenes</i> , Belgium. Emerging Infectious Diseases, 2012, 18, 1515-1518.	4.3	31
45	EFFECT OF GTS-21, AN ALPHA7 NICOTINIC ACETYLCHOLINE RECEPTOR AGONIST, ON CLP-INDUCED INFLAMMATORY, GASTROINTESTINAL MOTILITY, AND COLONIC PERMEABILITY CHANGES IN MICE. Shock, 2016, 45, 450-459.	2.1	31
46	BacPipe: A Rapid, User-Friendly Whole-Genome Sequencing Pipeline for Clinical Diagnostic Bacteriology. IScience, 2020, 23, 100769.	4.1	31
47	Oropharyngeal carriage of macrolide-resistant viridans group streptococci: a prevalence study among healthy adults in Belgium. Journal of Antimicrobial Chemotherapy, 2004, 53, 271-276.	3.0	29
48	Impact of amoxicillin therapy on resistance selection in patients with community-acquired lower respiratory tract infections: a randomized, placebo-controlled study. Journal of Antimicrobial Chemotherapy, 2016, 71, 3258-3267.	3.0	29
49	Animal models of hospital-acquired pneumonia: current practices and future perspectives. Annals of Translational Medicine, 2017, 5, 132-132.	1.7	29
50	Nasopharyngeal s. pneumoniae carriage and density in Belgian infants after 9†years of pneumococcal conjugate vaccine programme. Vaccine, 2018, 36, 15-22.	3.8	29
51	Bacitracin-Resistant Clone of Streptococcus pyogenes Isolated from Pharyngitis Patients in Belgium. Journal of Clinical Microbiology, 2003, 41, 5282-5284.	3.9	28
52	Metagenomic analysis of the impact of nitrofurantoin treatment on the human faecal microbiota. Journal of Antimicrobial Chemotherapy, 2015, 70, 1989-1992.	3.0	28
53	The COMPARE Data Hubs. Database: the Journal of Biological Databases and Curation, 2019, 2019, .	3.0	28
54	Emergence of high-level fluoroquinolone resistance in emm6 Streptococcus pyogenes and in vitro resistance selection with ciprofloxacin, levofloxacin and moxifloxacin. Journal of Antimicrobial Chemotherapy, 2009, 63, 886-894.	3.0	27

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55	Evaluation of Molecular Assays for Rapid Detection of Methicillin-Resistant <i>Staphylococcus aureus</i> . Journal of Clinical Microbiology, 2010, 48, 4598-4601.	3.9	27
56	A multinational study of colonization with extended spectrum $\hat{l}^2$ -lactamase-producing Enterobacteriaceae in healthcare personnel and family members of carrier patients hospitalized in rehabilitation centres. Clinical Microbiology and Infection, 2014, 20, O516-O523.	6.0	27
57	Fatty acid kinase A is an important determinant of biofilm formation in Staphylococcus aureus USA300. BMC Genomics, 2015, 16, 861.	2.8	26
58	Prospective One Health genetic surveillance in Vietnam identifies distinct blaCTX-M-harbouring Escherichia coli in food-chain and human-derived samples. Clinical Microbiology and Infection, 2021, 27, 1515.e1-1515.e8.	6.0	26
59	AMRmap: An Interactive Web Platform for Analysis of Antimicrobial Resistance Surveillance Data in Russia. Frontiers in Microbiology, 2021, 12, 620002.	3.5	26
60	Comparison of GeneXpert MRSA/SA ETA assay with semi-quantitative and quantitative cultures and nuc gene-based qPCR for detection of Staphylococcus aureus in endotracheal aspirate samples. Antimicrobial Resistance and Infection Control, 2019, 8, 4.	4.1	25
61	Antimicrobial Resistance Following Azithromycin Mass Drug Administration: Potential Surveillance Strategies to Assess Public Health Impact. Clinical Infectious Diseases, 2020, 70, 1501-1508.	5.8	25
62	Faropenem Consumption is Increasing in India. Clinical Infectious Diseases, 2016, 62, 1050.2-1052.	5.8	24
63	Follow-up of serotype distribution and antimicrobial susceptibility of Streptococcus pneumoniae in child carriage after a PCV13-to-PCV10 vaccine switch in Belgium. Vaccine, 2019, 37, 1080-1086.	3.8	23
64	A dynamic mucin mRNA signature associates with COVID-19 disease presentation and severity. JCI Insight, 2021, $6$ , .	5.0	23
65	Susceptibility profiles and resistance genomics of <i>Pseudomonas aeruginosa</i> isolates from European ICUs participating in the ASPIRE-ICU trial. Journal of Antimicrobial Chemotherapy, 2022, 77, 1862-1872.	3.0	23
66	Enzymes Catalyzing the TCA- and Urea Cycle Influence the Matrix Composition of Biofilms Formed by Methicillin-Resistant Staphylococcus aureus USA300. Microorganisms, 2018, 6, 113.	3.6	21
67	Association of <i>Staphylococcus aureus</i> Colonization and Pneumonia in the Intensive Care Unit. JAMA Network Open, 2020, 3, e2012741.	5.9	21
68	Incidence and predictive biomarkers of Clostridioides difficile infection in hospitalized patients receiving broad-spectrum antibiotics. Nature Communications, 2021, 12, 2240.	12.8	21
69	How nasopharyngeal pneumococcal carriage evolved during and after a PCV13-to-PCV10 vaccination programme switch in Belgium, 2016 to 2018. Eurosurveillance, 2020, 25, .	7.0	21
70	Quantifying antibiotic impact on within-patient dynamics of extended-spectrum beta-lactamase resistance. ELife, 2020, 9, .	6.0	21
71	Beneficial Effects of Anti-Interleukin-6 Antibodies on Impaired Gastrointestinal Motility, Inflammation and Increased Colonic Permeability in a Murine Model of Sepsis Are Most Pronounced When Administered in a Preventive Setup. PLoS ONE, 2016, 11, e0152914.	2.5	20
72	Interspecies Recombination Occurs Frequently in Quinolone Resistance-Determining Regions of Clinical Isolates of <i>Streptococcus pyogenes</i> Antimicrobial Agents and Chemotherapy, 2008, 52, 4191-4193.	3.2	19

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<b>7</b> 3	Unusual resistance patterns in macrolide-resistant Streptococcus pyogenes harbouring erm(A). Journal of Antimicrobial Chemotherapy, 2009, 63, 42-46.	3.0	19
74	Increase in fluoroquinolone non-susceptibility among clinical Streptococcus pyogenes in Belgium during 2007-10. Journal of Antimicrobial Chemotherapy, 2012, 67, 2602-2605.	3.0	19
75	Biofilm-Induced Type 2 Innate Immunity in a Cystic Fibrosis Model of Pseudomonas aeruginosa. Frontiers in Cellular and Infection Microbiology, 2017, 7, 274.	3.9	19
76	Molecular pathways to high-level azithromycin resistance in <i>Neisseria gonorrhoeae</i> Antimicrobial Chemotherapy, 2021, 76, 1752-1758.	3.0	19
77	Characterizing the in vitro biofilm phenotype of Staphylococcus epidermidis isolates from central venous catheters. Journal of Microbiological Methods, 2016, 127, 95-101.	1.6	18
78	The interplay between community and hospital Enterococcus faecium clones within health-care settings: a genomic analysis. Lancet Microbe, The, 2022, 3, e133-e141.	7.3	17
79	No SARSâ€CoVâ€2 carriage observed in children attending daycare centers during the intialÂweeks of the epidemic in Belgium. Journal of Medical Virology, 2021, 93, 1828-1831.	5.0	16
80	Whole-genome typing and characterization ofblaVIM19-harbouring ST383Klebsiella pneumoniaeby PFGE, whole-genome mapping and WGS. Journal of Antimicrobial Chemotherapy, 2016, 71, 1501-1509.	3.0	14
81	Remarkable Genome Stability among emm1 Group A Streptococcus in Belgium over 19 Years. Genome Biology and Evolution, 2019, 11, 1432-1439.	2.5	14
82	Emergence of colistin resistance during treatment of recurrent pneumonia caused by carbapenemase producing Klebsiella pneumoniae. Diagnostic Microbiology and Infectious Disease, 2019, 94, 407-409.	1.8	14
83	Remarkable geographical variations between India and Europe in carriage of the staphylococcal surface protein-encoding sasX/sesI and in the population structure of methicillin-resistant Staphylococcus aureus belonging to clonal complex 8. Clinical Microbiology and Infection, 2019, 25, 628.e1-628.e7.	6.0	14
84	Effects of intestinal alkaline phosphatase on intestinal barrier function in a cecal ligation and puncture (CLP)â€induced mouse model for sepsis. Neurogastroenterology and Motility, 2020, 32, e13754.	3.0	14
85	In-depth analysis of pneumococcal serotypes in Belgian children (2015–2018): Diversity, invasive disease potential, and antimicrobial susceptibility in carriage and disease. Vaccine, 2021, 39, 372-379.	3.8	14
86	Complete Genome Sequences of Two Prolific Biofilm-Forming Staphylococcus aureus Isolates Belonging to USA300 and EMRSA-15 Clonal Lineages. Genome Announcements, 2014, 2, .	0.8	13
87	Rationale and design of ASPIRE-ICU: a prospective cohort study on the incidence and predictors of Staphylococcus aureus and Pseudomonas aeruginosa pneumonia in the ICU. BMC Infectious Diseases, 2017, 17, 643.	2.9	13
88	Exploring Virulence Factors and Alternative Therapies against Staphylococcus aureus Pneumonia. Toxins, 2020, 12, 721.	3.4	13
89	Employing whole genome mapping for optimal de novo assembly of bacterial genomes. BMC Research Notes, 2014, 7, 484.	1.4	12
90	Optimization of an in vitro gut microbiome biotransformation platform with chlorogenic acid as model compound: From fecal sample to biotransformation product identification. Journal of Pharmaceutical and Biomedical Analysis, 2019, 175, 112768.	2.8	12

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91	Mechanical Ventilation Impairs IL-17 Cytokine Family Expression in Ventilator-Associated Pneumonia. International Journal of Molecular Sciences, 2019, 20, 5072.	4.1	12
92	Whole genome mapping as a fast-track tool to assess genomic stability of sequenced Staphylococcus aureus strains. BMC Research Notes, 2014, 7, 704.	1.4	11
93	Host Immunity Influences the Composition of Murine Gut Microbiota. Frontiers in Immunology, 2022, 13, 828016.	4.8	11
94	Complete Genome Sequences of Nitrofurantoin-Sensitive and -Resistant Escherichia coli ST540 and ST2747 Strains. Genome Announcements, 2014, 2, .	0.8	10
95	A longitudinal study of Staphylococcus aureus colonization in pigs in Ireland. Veterinary Microbiology, 2014, 174, 504-513.	1.9	10
96	Blood Cytokine Analysis Suggests That SARS-CoV-2 Infection Results in a Sustained Tumour Promoting Environment in Cancer Patients. Cancers, 2021, 13, 5718.	3.7	10
97	The effects of topical antibiotics on eradication and acquisition of third-generation cephalosporin and carbapenem-resistant Gram-negative bacteria in ICU patients; a post hoc analysis from a multicentre cluster-randomized trial. Clinical Microbiology and Infection, 2020, 26, 485-491.	6.0	9
98	Emergence of ST654 Pseudomonas aeruginosa co-harbouring blaNDM-1 and blaGES-5 in novel class I integron In1884 from Bulgaria. Journal of Global Antimicrobial Resistance, 2020, 22, 672-673.	2.2	6
99	Mechanical Ventilation Induces Interleukin 4 Secretion in Lungs and Reduces the Phagocytic Capacity of Lung Macrophages. Journal of Infectious Diseases, 2018, 217, 1645-1655.	4.0	5
100	Identification of mcr-8 in Clinical Isolates From Qatar and Evaluation of Their Antimicrobial Profiles. Frontiers in Microbiology, 2020, 11, 1954.	3.5	5
101	Streptococcus pneumoniae Serotypes Carried by Young Children and Their Association With Acute Otitis Media During the Period 2016–2019. Frontiers in Pediatrics, 2021, 9, 664083.	1.9	5
102	Sub-Inhibitory Concentrations of Chlorhexidine Induce Resistance to Chlorhexidine and Decrease Antibiotic Susceptibility in Neisseria gonorrhoeae. Frontiers in Microbiology, 2021, 12, 776909.	3.5	5
103	Molecular characteristics of community-associated methicillin-resistant Staphylococcus aureus colonizing surgical patients in Greece. Diagnostic Microbiology and Infectious Disease, 2012, 74, 420-422.	1.8	4
104	Chlorogenic Acid as a Model Compound for Optimization of an In Vitro Gut Microbiome-Metabolism Model. Proceedings (mdpi), 2019, 11, 31.	0.2	4
105	Potential in vivo transfer of a blaCTX-M14-harbouring plasmid established by combining long- and short-read sequencing. Journal of Microbiological Methods, 2019, 159, 1-4.	1.6	4
106	Detection and characterization of two NDM-1-producing <i>Klebsiella pneumoniae</i> Bulgaria. Journal of Antimicrobial Chemotherapy, 2016, 71, 1428-1430.	3.0	3
107	Detection of colistin resistance in the highly virulent Escherichia coli ST131 H30Rx clone in Greece. Journal of Global Antimicrobial Resistance, 2020, 20, 31-32.	2.2	3
108	Editorial Commentary: Mass Azithromycin Distribution and Emerging Resistance: Taking a Minimum Harms Approach. Clinical Infectious Diseases, 2013, 56, 1527-1529.	5.8	2

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109	Clonal transmission of multidrug-resistant Acinetobacter baumannii harbouring blaOXA-24-like and blaOXA-23-like genes in a tertiary hospital in Albania. Journal of Global Antimicrobial Resistance, 2020, 23, 79-81.	2.2	2
110	Evaluation of GeneXpert PA assay compared to genomic and (semi-)quantitative culture methods for direct detection of Pseudomonas aeruginosa in endotracheal aspirates. Antimicrobial Resistance and Infection Control, 2021, 10, 110.	4.1	2
111	Assessing the Impact of Flavophospholipol and Virginiamycin Supplementation on the Broiler Microbiota: a Prospective Controlled Intervention Study. MSystems, 2021, 6, e0038121.	3.8	2
112	Detection of SARS-CoV-2 in young children attending day-care centres in Belgium, May 2020 to February 2022. Eurosurveillance, 2022, 27, .	7.0	2
113	Novel composite SCCmec type III element in ST239 MRSA isolated from an Indian hospital. Journal of Antimicrobial Chemotherapy, 2018, 74, 264-266.	3.0	1
114	Analysis from the NeoOBS Global Neonatal Sepsis Prospective Observational Cohort Study Across 19 Hospitals in 11 Countries; Clinical Presentation, Treatment, Mortality Outcomes and Development of the NeoSEP Sepsis Severity Score. SSRN Electronic Journal, 0, , .	0.4	1
115	Obesity influences the microbiotic biotransformation of chlorogenic acid. Journal of Pharmaceutical and Biomedical Analysis, 2022, 211, 114550.	2.8	1
116	Screening of Anorectal and Oropharyngeal Samples Fails to Detect Bacteriophages Infecting Neisseria gonorrhoeae. Antibiotics, 2022, $11,268$ .	3.7	1
117	Identification of Potential Urinary Metabolite Biomarkers of <i>Pseudomonas aeruginosa</i> Ventilator-Associated Pneumonia. Biomarker Insights, 2022, 17, 117727192210991.	2.5	1
118	Sall72 - Effects of the Non-Selective Protease Inhibitor Nafamostat Mesylate on Intestinal Permeability and Bacterial Translocation in a Murine Model of Sepsis. Gastroenterology, 2018, 154, S-267-S-268.	1.3	0
119	COMBACTE LAB-Net: building a European laboratory network for clinical trials on anti-infectives. Future Microbiology, 2021, 16, 635-647.	2.0	0
120	Clinical Presentation and 28-Day Mortality in Hospitalized Neonates and Young Infants with Clinical Sepsis: The Global NeoOBS Observational Cohort. SSRN Electronic Journal, 0, , .	0.4	0
121	BacPipe: A Rapid, User-Friendly Whole Genome Sequencing Pipeline for Clinical Diagnostic Bacteriology. SSRN Electronic Journal, 0, , .	0.4	0
122	1486. Phylogenetic and alpha toxin variant analyses of Staphylococcus aureus strains isolated from patients during the SAATELLITE study. Open Forum Infectious Diseases, 2020, 7, S744-S745.	0.9	0