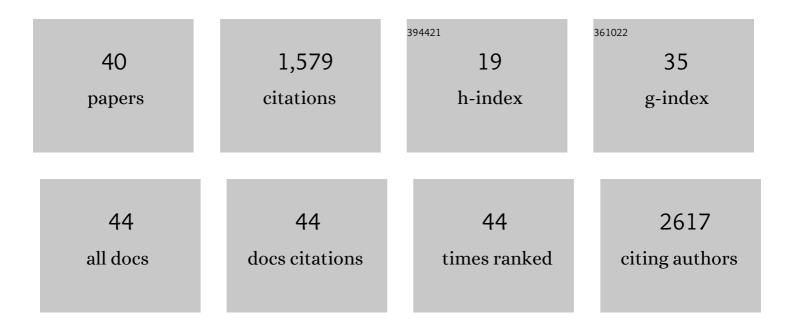
Medini K Annavajhala

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Prolonged outbreak of clonal, mupirocin-resistant methicillin-resistant Staphylococcus aureus in a neonatal intensive care unit: association with personnel and a possible environmental reservoir, analyzed using whole genome sequencing. American Journal of Infection Control, 2022, 50, 680-685.	2.3	3
2	Prolonged severe acute respiratory syndrome coronavirus 2 persistence, attenuated immunologic response, and viral evolution in a solid organ transplant patient. American Journal of Transplantation, 2022, 22, 649-653.	4.7	9
3	Population structure of <i>bla</i> KPC-harbouring IncN plasmids at a New York City medical centre and evidence for multi-species horizontal transmission. Journal of Antimicrobial Chemotherapy, 2022, 77, 1873-1882.	3.0	7
4	Klebsiella pneumoniae induces host metabolic stress that promotes tolerance to pulmonary infection. Cell Metabolism, 2022, 34, 761-774.e9.	16.2	36
5	Development and performance of a point-of-care rapid antigen test for detection of SARS-COV-2 variants. Journal of Clinical Virology Plus, 2022, 2, 100080.	1.0	3
6	Partial ORF1ab Gene Target Failure with Omicron BA.2.12.1. Journal of Clinical Microbiology, 2022, 60, e0060022.	3.9	11
7	Time to act–assessing variations in qPCR analyses in biological nitrogen removal with examples from partial nitritation/anammox systems. Water Research, 2021, 190, 116604.	11.3	8
8	Carbapenemase-producing Enterobacterales causing secondary infections during the COVID-19 crisis at a New York City hospital. Journal of Antimicrobial Chemotherapy, 2021, 76, 380-384.	3.0	58
9	Closing The Brief Case: A Maggot Mystery—Ignatzschineria larvae Sepsis Secondary to an Infested Wound. Journal of Clinical Microbiology, 2021, 59, .	3.9	1
10	The Brief Case: A Maggot Mystery—Ignatzschineria larvae Sepsis Secondary to an Infested Wound. Journal of Clinical Microbiology, 2021, 59, .	3.9	1
11	Staphylococcus aureus induces an itaconate-dominated immunometabolic response that drives biofilm formation. Nature Communications, 2021, 12, 1399.	12.8	72
12	<i>Spa</i> Typing of <i>Staphylococcus aureus</i> in a Neonatal Intensive Care Unit During Routine Surveillance. Journal of the Pediatric Infectious Diseases Society, 2021, 10, 766-773.	1.3	4
13	An acquired acyltransferase promotes Klebsiella pneumoniae ST258 respiratory infection. Cell Reports, 2021, 35, 109196.	6.4	15
14	Associations between urinary 3-indoxyl sulfate, a gut microbiome-derived biomarker, and patient outcomes after intensive care unit admission. Journal of Critical Care, 2021, 63, 15-21.	2.2	4
15	Emergence and expansion of SARS-CoV-2 B.1.526 after identification in New York. Nature, 2021, 597, 703-708.	27.8	103
16	Oral Microbiome Alterations and SARS-CoV-2 Saliva Viral Load in Patients with COVID-19. Microbiology Spectrum, 2021, 9, e0005521.	3.0	31
17	Emergence of Polymyxin Resistance in Clinical Klebsiella pneumoniae Through Diverse Genetic Adaptations: A Genomic, Retrospective Cohort Study. Clinical Infectious Diseases, 2020, 70, 2084-2091.	5.8	45
18	CrrB Positively Regulates High-Level Polymyxin Resistance and Virulence in Klebsiella pneumoniae. Cell Reports, 2020, 33, 108313.	6.4	39

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19	TAML- and Buffer-Catalyzed Oxidation of Picric Acid by H2O2: Products, Kinetics, DFT, and the Mechanism of Dual Catalysis. Inorganic Chemistry, 2020, 59, 13223-13232.	4.0	4
20	Subgingival microbiome and clinical periodontal status in an elderly cohort: The WHICAP ancillary study of oral health. Journal of Periodontology, 2020, 91, S56-S67.	3.4	31
21	Oral and Gut Microbial Diversity and Immune Regulation in Patients with HIV on Antiretroviral Therapy. MSphere, 2020, 5, .	2.9	41
22	Relationship of the Esophageal Microbiome and Tissue Gene Expression and Links to the Oral Microbiome: A Randomized Clinical Trial. Clinical and Translational Gastroenterology, 2020, 11, e00235.	2.5	13
23	Colonizing multidrug-resistant bacteria and the longitudinal evolution of the intestinal microbiome after liver transplantation. Nature Communications, 2019, 10, 4715.	12.8	70
24	Transfusional iron overload and intravenous iron infusions modify the mouse gut microbiota similarly to dietary iron. Npj Biofilms and Microbiomes, 2019, 5, 26.	6.4	35
25	Escherichia coli Harboring mcr-1 in a Cluster of Liver Transplant Recipients: Detection through Active Surveillance and Whole-Genome Sequencing. Antimicrobial Agents and Chemotherapy, 2019, 63, .	3.2	8
26	Metabolic Adaptation in Methicillin-Resistant <i>Staphylococcus aureus</i> Pneumonia. American Journal of Respiratory Cell and Molecular Biology, 2019, 61, 185-197.	2.9	34
27	Multidrug-Resistant Enterobacter cloacae Complex Emerging as a Global, Diversifying Threat. Frontiers in Microbiology, 2019, 10, 44.	3.5	211
28	Comammox Functionality Identified in Diverse Engineered Biological Wastewater Treatment Systems. Environmental Science and Technology Letters, 2018, 5, 110-116.	8.7	118
29	Successive Emergence of Ceftazidime-Avibactam Resistance through Distinct Genomic Adaptations in <i>bla</i> _{KPC-2} -Harboring Klebsiella pneumoniae Sequence Type 307 Isolates. Antimicrobial Agents and Chemotherapy, 2018, 62, .	3.2	162
30	1249. Emergence of Diverse Carbapenem-Resistant Enterobacteriaceae (CRE) in the Dominican Republic. Open Forum Infectious Diseases, 2018, 5, S380-S380.	0.9	0
31	Metatranscriptomic Investigation of Adaptation in NO and N2O Production From a Lab-Scale Nitrification Process Upon Repeated Exposure to Anoxic–Aerobic Cycling. Frontiers in Microbiology, 2018, 9, 3012.	3.5	10
32	Structural and Functional Interrogation of Selected Biological Nitrogen Removal Systems in the United States, Denmark, and Singapore Using Shotgun Metagenomics. Frontiers in Microbiology, 2018, 9, 2544.	3.5	7
33	Genomic and Geographic Context for the Evolution of High-Risk Carbapenem-Resistant <i>Enterobacter cloacae</i> Complex Clones ST171 and ST78. MBio, 2018, 9, .	4.1	67
34	Mo1941 - A Randomized Controlled Trial to Assess the Effects of an Antimicrobial Mouthwash on the Oral and Esophageal Microbiome. Gastroenterology, 2018, 154, S-859.	1.3	0
35	Pathogen colonization of the gastrointestinal microbiome at intensive care unit admission and risk for subsequent death or infection. Intensive Care Medicine, 2018, 44, 1203-1211.	8.2	121
36	791 - Gastrointestinal Bacterial Pathogen Colonization and Risk for Subsequent Infection in the Intensive Care Unit. Gastroenterology, 2018, 154, S-165.	1.3	0

#	Article	IF	CITATIONS
37	Clonal Background, Resistance Gene Profile, and Porin Gene Mutations Modulate <i>In Vitro</i> Susceptibility to Imipenem-Relebactam in Diverse Enterobacteriaceae. Antimicrobial Agents and Chemotherapy, 2018, 62, .	3.2	34
38	Presence and functional potential of comammox in full-scale wastewater treatment systems across the globe. Proceedings of the Water Environment Federation, 2017, 2017, 4060-4068.	0.0	0
39	Metagenomics of Anaerobic Food Waste Fermentation. Proceedings of the Water Environment Federation, 2017, 2017, 4041-4047.	0.0	Ο
40	Experimental and Theoretical Evidence for Multiple Fe ^{IV} Reactive Intermediates in TAMLâ€Activator Catalysis: Rationalizing a Counterintuitive Reactivity Order. Chemistry - A European Journal, 2012, 18, 10244-10249.	3.3	22