

Theodore Gouliouris

List of Publications by Year in descending order

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Version: 2024-02-01

43
papers

3,147
citations

304743

22
h-index

276875

41
g-index

47
all docs

47
docs citations

47
times ranked

6077
citing authors

#	ARTICLE	IF	CITATIONS
1	The Removal of Airborne Severe Acute Respiratory Syndrome Coronavirus 2 (SARS-CoV-2) and Other Microbial Bioaerosols by Air Filtration on Coronavirus Disease 2019 (COVID-19) Surge Units. <i>Clinical Infectious Diseases</i> , 2022, 75, e97-e101.	5.8	52
2	A2B-COVID: A Tool for Rapidly Evaluating Potential SARS-CoV-2 Transmission Events. <i>Molecular Biology and Evolution</i> , 2022, 39, .	8.9	12
3	The interplay between community and hospital <i>Enterococcus faecium</i> clones within health-care settings: a genomic analysis. <i>Lancet Microbe</i> , The, 2022, 3, e133-e141.	7.3	17
4	PowerBacGWAS: a computational pipeline to perform power calculations for bacterial genome-wide association studies. <i>Communications Biology</i> , 2022, 5, 266.	4.4	4
5	Machine Learning for Antimicrobial Resistance Prediction: Current Practice, Limitations, and Clinical Perspective. <i>Clinical Microbiology Reviews</i> , 2022, 35, .	13.6	33
6	Investigation of healthcare-associated SARS-CoV-2 infection: Learning outcomes from an investigative process in the initial phase of the pandemic. <i>Journal of Infection Prevention</i> , 2022, 23, 197-205.	0.9	1
7	Quantifying acquisition and transmission of <i>Enterococcus faecium</i> using genomic surveillance. <i>Nature Microbiology</i> , 2021, 6, 103-111.	13.3	53
8	SARS-CoV-2 evolution during treatment of chronic infection. <i>Nature</i> , 2021, 592, 277-282.	27.8	802
9	Mortality Following <i>Clostridioides difficile</i> Infection in Europe: A Retrospective Multicenter Case-Control Study. <i>Antibiotics</i> , 2021, 10, 299.	3.7	23
10	The global dissemination of hospital clones of <i>Enterococcus faecium</i> . <i>Genome Medicine</i> , 2021, 13, 52.	8.2	33
11	Low diagnostic yield and time to diagnostic confirmation results in prolonged use of antimicrobials in critically ill children. <i>Wellcome Open Research</i> , 2021, 6, 119.	1.8	5
12	Methicillin-resistant <i>Staphylococcus aureus</i> (MRSA) submandibular abscess in a neonate. <i>BMJ Case Reports</i> , 2021, 14, e242258.	0.5	0
13	Applying prospective genomic surveillance to support investigation of hospital-onset COVID-19. <i>Lancet Infectious Diseases</i> , The, 2021, 21, 916-917.	9.1	14
14	Superspreaders drive the largest outbreaks of hospital onset COVID-19 infections. <i>ELife</i> , 2021, 10, .	6.0	34
15	Defining nosocomial transmission of <i>Escherichia coli</i> and antimicrobial resistance genes: a genomic surveillance study. <i>Lancet Microbe</i> , The, 2021, 2, e472-e480.	7.3	39
16	Rapid Assay for Sick Children with Acute Lung infection Study (RASCALS): diagnostic cohort study protocol. <i>BMJ Open</i> , 2021, 11, e056197.	1.9	5
17	A One Health Study of the Genetic Relatedness of <i>Klebsiella pneumoniae</i> and Their Mobile Elements in the East of England. <i>Clinical Infectious Diseases</i> , 2020, 70, 219-226.	5.8	46
18	Rapid implementation of SARS-CoV-2 sequencing to investigate cases of health-care associated COVID-19: a prospective genomic surveillance study. <i>Lancet Infectious Diseases</i> , The, 2020, 20, 1263-1271.	9.1	352

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19	Defining persistent <i>Staphylococcus aureus</i> bacteraemia: secondary analysis of a prospective cohort study. <i>Lancet Infectious Diseases</i> , The, 2020, 20, 1409-1417.	9.1	84
20	One Health Genomic Surveillance of <i>Escherichia coli</i> Demonstrates Distinct Lineages and Mobile Genetic Elements in Isolates from Humans versus Livestock. <i>MBio</i> , 2019, 10, .	4.1	130
21	Detection of vancomycin-resistant <i>Enterococcus faecium</i> hospital-adapted lineages in municipal wastewater treatment plants indicates widespread distribution and release into the environment. <i>Genome Research</i> , 2019, 29, 626-634.	5.5	40
22	Genomic surveillance of <i>Escherichia coli</i> in municipal wastewater treatment plants as an indicator of clinically relevant pathogens and their resistance genes. <i>Microbial Genomics</i> , 2019, 5, .	2.0	29
23	Genome-Based Analysis of <i>Enterococcus faecium</i> Bacteremia Associated with Recurrent and Mixed-Strain Infection. <i>Journal of Clinical Microbiology</i> , 2018, 56, .	3.9	14
24	Duration of exposure to multiple antibiotics is associated with increased risk of VRE bacteraemia: a nested case-control study. <i>Journal of Antimicrobial Chemotherapy</i> , 2018, 73, 1692-1699.	3.0	40
25	Adjunctive rifampicin for <i>Staphylococcus aureus</i> bacteraemia (ARREST): a multicentre, randomised, double-blind, placebo-controlled trial. <i>Lancet</i> , The, 2018, 391, 668-678.	13.7	140
26	Genomic Surveillance of <i>Enterococcus faecium</i> Reveals Limited Sharing of Strains and Resistance Genes between Livestock and Humans in the United Kingdom. <i>MBio</i> , 2018, 9, .	4.1	63
27	Genomic survey of <i>Clostridium difficile</i> reservoirs in the East of England implicates environmental contamination of wastewater treatment plants by clinical lineages. <i>Microbial Genomics</i> , 2018, 4, .	2.0	19
28	Complex Routes of Nosocomial Vancomycin-Resistant <i>Enterococcus faecium</i> Transmission Revealed by Genome Sequencing. <i>Clinical Infectious Diseases</i> , 2017, 64, 886-893.	5.8	93
29	A Retained Stitch in Time Saves 9 - But Does it Increase the Risk of Deep Prosthetic Infection?. <i>HIP International</i> , 2017, 27, 564-566.	1.7	0
30	Within-host evolution of <i>Enterococcus faecium</i> during longitudinal carriage and transition to bloodstream infection in immunocompromised patients. <i>Genome Medicine</i> , 2017, 9, 119.	8.2	26
31	Sharing of carbapenemase-encoding plasmids between Enterobacteriaceae in UK sewage uncovered by MiniON sequencing. <i>Microbial Genomics</i> , 2017, 3, e000114.	2.0	33
32	Impact of a candidaemia care bundle on patient care at a large teaching hospital in England. <i>Journal of Infection</i> , 2016, 72, 501-503.	3.3	17
33	Genome-based characterization of hospital-adapted <i>Enterococcus faecalis</i> lineages. <i>Nature Microbiology</i> , 2016, 1, .	13.3	65
34	Comparison of two chromogenic media for the detection of vancomycin-resistant enterococcal carriage by nursing home residents. <i>Diagnostic Microbiology and Infectious Disease</i> , 2016, 85, 409-412.	1.8	7
35	Comparison of 2 chromogenic media for the detection of extended-spectrum β -lactamase producing Enterobacteriaceae stool carriage in nursing home residents. <i>Diagnostic Microbiology and Infectious Disease</i> , 2016, 84, 181-183.	1.8	16
36	The Protective Role of Albumin in <i>Clostridium difficile</i> Infection: A Step Toward Solving the Puzzle. <i>Infection Control and Hospital Epidemiology</i> , 2015, 36, 1478-1480.	1.8	18

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37	Fecal microbiota transplantation (FMT) for <i>Clostridium difficile</i> infection: Focus on immunocompromised patients. <i>Journal of Infection and Chemotherapy</i> , 2015, 21, 230-237.	1.7	65
38	Impact of infectious diseases consultation on the management of <i>Staphylococcus aureus</i> bacteraemia in children. <i>BMJ Open</i> , 2014, 4, e004659-e004659.	1.9	18
39	Rapid Bacterial Whole-Genome Sequencing to Enhance Diagnostic and Public Health Microbiology. <i>JAMA Internal Medicine</i> , 2013, 173, 1397.	5.1	181
40	Use of Vitek 2 Antimicrobial Susceptibility Profile To Identify <i>mecC</i> in Methicillin-Resistant <i>Staphylococcus aureus</i> . <i>Journal of Clinical Microbiology</i> , 2013, 51, 2732-2734.	3.9	53
41	<i>S. truongyloides stercoralis</i> hyperinfection in a patient treated for multiple myeloma. <i>British Journal of Haematology</i> , 2012, 158, 2-2.	2.5	7
42	Prevention and treatment of <i>Clostridium difficile</i> infection. <i>Clinical Medicine</i> , 2011, 11, 75-79.	1.9	11
43	Spondylodiscitis: update on diagnosis and management. <i>Journal of Antimicrobial Chemotherapy</i> , 2010, 65, iii11-iii24.	3.0	418