

Dennis Nurjadi

List of Publications by Year in descending order

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Version: 2024-02-01

63
papers

1,109
citations

471061

17
h-index

454577

30
g-index

67
all docs

67
docs citations

67
times ranked

1363
citing authors

#	ARTICLE	IF	CITATIONS
1	Rapid Development of Cefiderocol Resistance in Carbapenem-resistant <i>Enterobacter cloacae</i> During Therapy Is Associated With Heterogeneous Mutations in the Catecholate Siderophore Receptor <i>cirA</i> . <i>Clinical Infectious Diseases</i> , 2022, 74, 905-908.	2.9	67
2	Direct-PCR from rectal swabs and environmental reservoirs: A fast and efficient alternative to detect blaOXA-48 carbapenemase genes in an <i>Enterobacter cloacae</i> outbreak setting. <i>Environmental Research</i> , 2022, 203, 111808.	3.7	5
3	Impact of discontinuing contact precautions and enforcement of basic hygiene measures on nosocomial vancomycin-resistant <i>Enterococcus faecium</i> transmission. <i>Journal of Hospital Infection</i> , 2022, 121, 120-127.	1.4	9
4	<i>Acinetobacter baumannii</i> . , 2022, , 113-129.		1
5	<i>Staphylococcus massiliensis</i> isolated from human blood cultures, Germany, 2017–2020. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2022, 41, 663-669.	1.3	0
6	New Delhi Metallo-Beta-Lactamase Facilitates the Emergence of Cefiderocol Resistance in <i>Enterobacter cloacae</i> . <i>Antimicrobial Agents and Chemotherapy</i> , 2022, 66, AAC0201121.	1.4	42
7	Imported Panton-valentine leucocidin (PVL)-positive <i>Staphylococcus aureus</i> skin infections: patients' perspective on quality of life and quality of medical care. <i>Journal of Travel Medicine</i> , 2022, 29, .	1.4	6
8	Comparative Genomic Reveals Clonal Heterogeneity in Persistent <i>Staphylococcus aureus</i> Infection. <i>Frontiers in Cellular and Infection Microbiology</i> , 2022, 12, 817841.	1.8	1
9	Treatment of Infection-Related Non-Unions with Bioactive Glass—A Promising Approach or Just Another Method of Dead Space Management?. <i>Materials</i> , 2022, 15, 1697.	1.3	4
10	Inflammatory Response Against <i>Staphylococcus aureus</i> via Intracellular Sensing of Nucleic Acids in Keratinocytes. <i>Frontiers in Immunology</i> , 2022, 13, 828626.	2.2	13
11	Cefiderocol Protects against Cytokine- and Endotoxin-Induced Disruption of Vascular Endothelial Cell Integrity in an In Vitro Experimental Model. <i>Antibiotics</i> , 2022, 11, 581.	1.5	1
12	Host genetic loci LZTFL1 and CCL2 associated with SARS-CoV-2 infection and severity of COVID-19. <i>International Journal of Infectious Diseases</i> , 2022, 122, 427-436.	1.5	11
13	Pitfalls in genotypic antimicrobial susceptibility testing caused by low expression of <i>bla</i> KPC in <i>Escherichia coli</i> . <i>Journal of Antimicrobial Chemotherapy</i> , 2021, 76, 2795-2801.	1.3	8
14	Genomic structure of ST8-t008 USA300 and USA300-LV MRSA in the Rhine-Neckar Region, Germany, 2012–2018. <i>International Journal of Antimicrobial Agents</i> , 2021, 57, 106312.	1.1	9
15	Maternal Vaginal Colonization and Extended-Spectrum Beta-Lactamase-Producing Bacteria in Vietnamese Pregnant Women. <i>Antibiotics</i> , 2021, 10, 572.	1.5	4
16	The Identification of the SARS-CoV-2 Whole Genome: Nine Cases Among Patients in Banten Province, Indonesia. <i>Journal of Pure and Applied Microbiology</i> , 2021, 15, 936-948.	0.3	0
17	Molecular Detection of Carbapenemases in Enterobacterales: A Comparison of Real-Time Multiplex PCR and Whole-Genome Sequencing. <i>Antibiotics</i> , 2021, 10, 726.	1.5	5
18	Existing evidence supports clinical trials on interventions preventing ventricular assist device infection in patients colonized with <i>Staphylococcus aureus</i> . Letter regarding the article "Association of preoperative infections, nasal <i>Staphylococcus aureus</i> colonization and gut microbiota with left ventricular assist device outcomes". <i>European Journal of Heart Failure</i> , 2021, 23, 1566-1566.	2.9	0

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19	Host factors facilitating SARS-CoV-2 virus infection and replication in the lungs. Cellular and Molecular Life Sciences, 2021, 78, 5953-5976.	2.4	19
20	Systemic Administration of PTH Supports Vascularization in Segmental Bone Defects Filled with Ceramic-Based Bone Graft Substitute. Cells, 2021, 10, 2058.	1.8	6
21	Association Between Soluble Notch Ligand Delta-like Ligand 1 and Bleeding Complications in Patients With Dengue Fever Infection. Journal of Infectious Diseases, 2021, , .	1.9	1
22	Molecular analysis of an increase in trimethoprim/sulfamethoxazole-resistant MRSA reveals multiple introductions into a tertiary care hospital, Germany 2012-19. Journal of Antimicrobial Chemotherapy, 2021, 77, 38-48.	1.3	7
23	Surveillance for Colonization, Transmission, and Infection With Methicillin-Susceptible <i>Staphylococcus aureus</i> in a Neonatal Intensive Care Unit. JAMA Network Open, 2021, 4, e2124938.	2.8	22
24	Molecular detection of blaCTX-M gene to predict phenotypic cephalosporin resistance and clinical outcome of Escherichia coli bloodstream infections in Vietnam. Annals of Clinical Microbiology and Antimicrobials, 2021, 20, 60.	1.7	4
25	Acquisition and Transmission of Carbapenemase-Producing (<i>bla</i> KPC-2) <i>Enterobacter cloacae</i> in a Highly Frequented Outpatient Clinic. Clinical Infectious Diseases, 2021, 72, e158-e161.	2.9	8
26	Phenotypic Detection of Hemin-Inducible Trimethoprim-Sulfamethoxazole Heteroresistance in <i>Staphylococcus aureus</i> . Microbiology Spectrum, 2021, 9, e0151021.	1.2	2
27	Genomic Investigation and Successful Containment of an Intermittent Common Source Outbreak of OXA-48-Producing <i>Enterobacter cloacae</i> Related to Hospital Shower Drains. Microbiology Spectrum, 2021, 9, e0138021.	1.2	8
28	Notch Ligand Delta-Like 1 Is Associated With Loss of Vascular Endothelial Barrier Function. Frontiers in Physiology, 2021, 12, 766713.	1.3	8
29	Invasiveness of <i>Escherichia coli</i> Is Associated with an IncFII Plasmid. Pathogens, 2021, 10, 1645.	1.2	3
30	Comparative genomic analysis reveals a high prevalence of inter-species <i>in vivo</i> transfer of carbapenem-resistance plasmids in patients with haematological malignancies. Clinical Microbiology and Infection, 2020, 26, 780.e1-780.e8.	2.8	21
31	Alteration of antibiotic regimen as an additional control measure in suspected multi-drug-resistant <i>Enterobacter cloacae</i> outbreak in a neonatal intensive care unit. Journal of Hospital Infection, 2020, 104, 144-149.	1.4	13
32	Challenges in interpretation of WGS and epidemiological data to investigate nosocomial transmission of vancomycin-resistant <i>Enterococcus faecium</i> in an endemic region: incorporation of patient movement network and admission screening. Journal of Antimicrobial Chemotherapy, 2020, 75, 1716-1721.	1.3	10
33	Whole-genome sequencing disproves two suspected transmission events of blaNDM between <i>Pseudomonas aeruginosa</i> and <i>Enterobacterales</i> in hospitalized patients. Journal of Hospital Infection, 2020, 106, 372-375.	1.4	1
34	Entry of Pantone-Valentine leukocidin-positive methicillin-resistant <i>Staphylococcus aureus</i> into the hospital: prevalence and population structure in Heidelberg, Germany 2015-2018. Scientific Reports, 2020, 10, 13243.	1.6	22
35	Comparative evaluation of the effect of different growth media on <i>in vitro</i> sensitivity to azithromycin in multi-drug resistant <i>Pseudomonas aeruginosa</i> isolated from cystic fibrosis patients. Antimicrobial Resistance and Infection Control, 2020, 9, 197.	1.5	9
36	Identification and Elimination of the Clinically Relevant Multi-Resistant Environmental Bacteria <i>Ralstonia insidiosa</i> in Primary Cell Culture. Microorganisms, 2020, 8, 1599.	1.6	6

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37	A new sequential animal model for infection-related non-unions with segmental bone defect. <i>BMC Musculoskeletal Disorders</i> , 2020, 21, 329.	0.8	9
38	Nasal colonization with <i>Staphylococcus aureus</i> is a risk factor for ventricular assist device infection in the first year after implantation: A prospective, single-centre, cohort study. <i>Journal of Infection</i> , 2020, 80, 511-518.	1.7	11
39	Integrative Analysis of Whole Genome Sequencing and Phenotypic Resistance Toward Prediction of Trimethoprim-Sulfamethoxazole Resistance in <i>Staphylococcus aureus</i> . <i>Frontiers in Microbiology</i> , 2020, 11, 607842.	1.5	18
40	Low prevalence of combined linezolid- and vancomycin-resistant <i>Enterococcus faecium</i> from hospital admission screening in an endemic region in Germany. <i>Journal of Global Antimicrobial Resistance</i> , 2020, 22, 646-650.	0.9	12
41	Emergence of carbapenem-resistant ST131 <i>Escherichia coli</i> carrying blaOXA-244 in Germany, 2019 to 2020. <i>Eurosurveillance</i> , 2020, 25, .	3.9	11
42	Molecular characterization of carbapenem-resistant <i>Acinetobacter baumannii</i> using WGS revealed missed transmission events in Germany from 2012â€“15. <i>Journal of Antimicrobial Chemotherapy</i> , 2019, 74, 3473-3480.	1.3	15
43	Increase in the prevalence of Pantonâ€“Valentine leukocidin and clonal shift in community-onset methicillin-resistant <i>Staphylococcus aureus</i> causing skin and soft-tissue infections in the Rhine-Neckar Region, Germany, 2012â€“2016. <i>International Journal of Antimicrobial Agents</i> , 2019, 53, 261-267.	1.1	32
44	Import of community-associated, methicillin-resistant <i>Staphylococcus aureus</i> to Europe through skin and soft-tissue infection in intercontinental travellers, 2011â€“2016. <i>Clinical Microbiology and Infection</i> , 2019, 25, 739-746.	2.8	35
45	Toll-like receptor 9 (TLR-9) promotor polymorphisms and gene expression are associated with persistent <i>Staphylococcus aureus</i> nasal carriage. <i>Clinical Microbiology and Infection</i> , 2018, 24, 1210.e7-1210.e12.	2.8	16
46	Significant increase in cultivation of <i>Gardnerella vaginalis</i> , <i>Alloscardovia omnicolens</i> , <i>Actinotignum schaalii</i> , and <i>Actinomyces</i> spp. in urine samples with total laboratory automation. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2018, 37, 1305-1311.	1.3	30
47	Draft Genome Sequence of <i>Staphylococcus aureus</i> Strain HD1410, Isolated from a Persistent Nasal Carrier. <i>Genome Announcements</i> , 2018, 6, .	0.8	4
48	Transmission of ST8-USA300 Latin American Variant Methicillin-Resistant <i>Staphylococcus aureus</i> on a Neonatal Intensive Care Unit: Recurrent Skin and Soft- Tissue Infections as a Marker for Epidemic Community-Associated-MRSA Colonization. <i>Infection Control and Hospital Epidemiology</i> , 2017, 38, 883-885.	1.0	11
49	A Trial of Antibiotics for Smaller Skin Abscesses. <i>New England Journal of Medicine</i> , 2017, 377, e36.	13.9	5
50	Ratio of T-Helper Type 1 (Th1) to Th17 Cytokines in Whole Blood Is Associated With Human Î²-Defensin 3 Expression in Skin and Persistent <i>Staphylococcus aureus</i> Nasal Carriage. <i>Journal of Infectious Diseases</i> , 2016, 214, 1744-1751.	1.9	14
51	Evaluation of antibiotic resistance to orally administrable antibiotics in staphylococcal bone and joint infections in one of the largest university hospitals in Germany: is there a role for fusidic acid?. <i>International Journal of Antimicrobial Agents</i> , 2016, 47, 155-157.	1.1	17
52	Skin and soft tissue infections in intercontinental travellers and the import of multi-resistant <i>Staphylococcus aureus</i> to Europe. <i>Clinical Microbiology and Infection</i> , 2015, 21, 567.e1-567.e10.	2.8	71
53	Predominance of dfrG as determinant of trimethoprim resistance in imported <i>Staphylococcus aureus</i> . <i>Clinical Microbiology and Infection</i> , 2015, 21, 1095.e5-1095.e9.	2.8	35
54	Emergence of trimethoprim resistance gene dfrG in <i>Staphylococcus aureus</i> causing human infection and colonization in sub-Saharan Africa and its import to Europe. <i>Journal of Antimicrobial Chemotherapy</i> , 2014, 69, 2361-2368.	1.3	87

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55	Effectiveness of rifaximin in prevention of diarrhoea in individuals travelling to south and southeast Asia: a randomised, double-blind, placebo-controlled, phase 3 trial. <i>Lancet Infectious Diseases</i> , The, 2013, 13, 946-954.	4.6	28
56	Impaired β -Defensin Expression in Human Skin Links DEFB1 Promoter Polymorphisms With Persistent <i>Staphylococcus aureus</i> Nasal Carriage. <i>Journal of Infectious Diseases</i> , 2013, 207, 666-674.	1.9	59
57	Import and Spread of Panton-Valentine Leukocidin-Positive <i>Staphylococcus aureus</i> Through Nasal Carriage and Skin Infections in Travelers Returning From the Tropics and Subtropics. <i>Clinical Infectious Diseases</i> , 2012, 54, 483-492.	2.9	78
58	Hormonal Contraceptive Use and Persistent <i>Staphylococcus aureus</i> Nasal Carriage. <i>Clinical Infectious Diseases</i> , 2012, 55, 1625-1632.	2.9	31
59	Reply to Mimica. <i>Clinical Infectious Diseases</i> , 2012, 54, 1518-1519.	2.9	0
60	Reliable and rapid characterization of functional FCN2 gene variants reveals diverse geographical patterns. <i>BMC Medical Genetics</i> , 2012, 13, 37.	2.1	17
61	Clonal expansion accounts for an excess of antimicrobial resistance in <i>Staphylococcus aureus</i> colonising HIV-positive individuals in Lagos, Nigeria. <i>International Journal of Antimicrobial Agents</i> , 2012, 40, 268-272.	1.1	32
62	<i>Staphylococcus aureus</i> throat carriage is associated with ABO-/secretor status. <i>Journal of Infection</i> , 2012, 65, 310-317.	1.7	31
63	Persistent Nasal Carriage of <i>Staphylococcus aureus</i> Is Associated with Deficient Induction of Human β -Defensin 3 after Sterile Wounding of Healthy Skin <i>In Vivo</i> . <i>Infection and Immunity</i> , 2011, 79, 2658-2662.	1.0	44