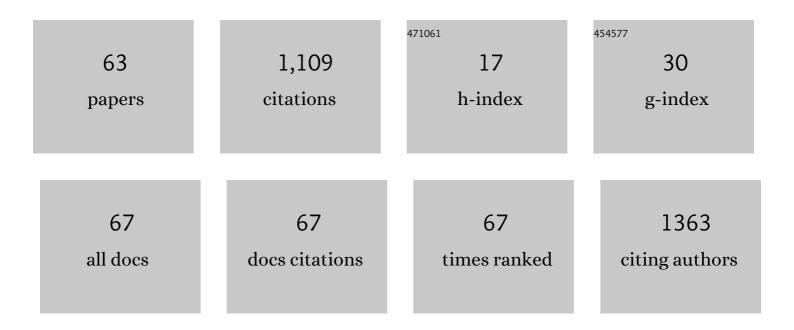
## Dennis Nurjadi

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Emergence of trimethoprim resistance gene dfrG in Staphylococcus aureus causing human infection and colonization in sub-Saharan Africa and its import to Europe. Journal of Antimicrobial Chemotherapy, 2014, 69, 2361-2368.	1.3	87
2	Import and Spread of Panton-Valentine Leukocidin–Positive Staphylococcus aureus Through Nasal Carriage and Skin Infections in Travelers Returning From the Tropics and Subtropics. Clinical Infectious Diseases, 2012, 54, 483-492.	2.9	78
3	Skin and soft tissue infections in intercontinental travellers and the import of multi-resistant Staphylococcus aureus to Europe. Clinical Microbiology and Infection, 2015, 21, 567.e1-567.e10.	2.8	71
4	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> . Clinical Infectious Diseases, 2022, 74, 905-908.	2.9	67
5	Impaired β-Defensin Expression in Human Skin Links DEFB1 Promoter Polymorphisms With Persistent Staphylococcus aureus Nasal Carriage. Journal of Infectious Diseases, 2013, 207, 666-674.	1.9	59
6	Persistent Nasal Carriage of Staphylococcus aureus Is Associated with Deficient Induction of Human β-Defensin 3 after Sterile Wounding of Healthy Skin <i>In Vivo</i> . Infection and Immunity, 2011, 79, 2658-2662.	1.0	44
7	New Delhi Metallo-Beta-Lactamase Facilitates the Emergence of Cefiderocol Resistance in Enterobacter cloacae. Antimicrobial Agents and Chemotherapy, 2022, 66, AAC0201121.	1.4	42
8	Predominance of dfrG as determinant of trimethoprim resistance in imported Staphylococcus aureus. Clinical Microbiology and Infection, 2015, 21, 1095.e5-1095.e9.	2.8	35
9	Import of community-associated, methicillin-resistant Staphylococcus aureus to Europe through skin and soft-tissue infection in intercontinental travellers, 2011–2016. Clinical Microbiology and Infection, 2019, 25, 739-746.	2.8	35
10	Clonal expansion accounts for an excess of antimicrobial resistance in Staphylococcus aureus colonising HIV-positive individuals in Lagos, Nigeria. International Journal of Antimicrobial Agents, 2012, 40, 268-272.	1.1	32
11	Increase in the prevalence of Panton–Valentine leukocidin and clonal shift in community-onset methicillin-resistant Staphylococcus aureus causing skin and soft-tissue infections in the Rhine-Neckar Region, Germany, 2012–2016. International Journal of Antimicrobial Agents, 2019, 53, 261-267.	1.1	32
12	Hormonal Contraceptive Use and Persistent Staphylococcus aureus Nasal Carriage. Clinical Infectious Diseases, 2012, 55, 1625-1632.	2.9	31
13	Staphylococcus aureus throat carriage is associated with ABO-/secretor status. Journal of Infection, 2012, 65, 310-317.	1.7	31
14	Significant increase in cultivation of Gardnerella vaginalis, Alloscardovia omnicolens, Actinotignum schaalii, and Actinomyces spp. in urine samples with total laboratory automation. European Journal of Clinical Microbiology and Infectious Diseases, 2018, 37, 1305-1311.	1.3	30
15	Effectiveness of rifaximin in prevention of diarrhoea in individuals travelling to south and southeast Asia: a randomised, double-blind, placebo-controlled, phase 3 trial. Lancet Infectious Diseases, The, 2013, 13, 946-954.	4.6	28
16	Entry of Panton–Valentine leukocidin-positive methicillin-resistant Staphylococcus aureus into the hospital: prevalence and population structure in Heidelberg, Germany 2015–2018. Scientific Reports, 2020, 10, 13243.	1.6	22
17	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
18	Comparative genomic analysis reveals a high prevalence of inter-species inÂvivo transfer of carbapenem-resistance plasmids in patients with haematological malignancies. Clinical Microbiology and Infection, 2020, 26, 780.e1-780.e8.	2.8	21

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19	Host factors facilitating SARSâ€CoVâ€2 virus infection and replication in the lungs. Cellular and Molecular Life Sciences, 2021, 78, 5953-5976.	2.4	19
20	Integrative Analysis of Whole Genome Sequencing and Phenotypic Resistance Toward Prediction of Trimethoprim-Sulfamethoxazole Resistance in Staphylococcus aureus. Frontiers in Microbiology, 2020, 11, 607842.	1.5	18
21	Reliable and rapid characterization of functional FCN2 gene variants reveals diverse geographical patterns. BMC Medical Genetics, 2012, 13, 37.	2.1	17
22	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?. International Journal of Antimicrobial Agents, 2016, 47, 155-157.	1.1	17
23	Toll-like receptor 9 (TLR-9) promotor polymorphisms and gene expression are associated with persistent Staphylococcus aureus nasalAcarriage. Clinical Microbiology and Infection, 2018, 24, 1210.e7-1210.e12.	2.8	16
24	Molecular characterization of carbapenem-resistant Acinetobacter baumannii using WGS revealed missed transmission events in Germany from 2012–15. Journal of Antimicrobial Chemotherapy, 2019, 74, 3473-3480.	1.3	15
25	Ratio of T-Helper Type 1 (Th1) to Th17 Cytokines in Whole Blood Is Associated With Human Î2-Defensin 3 Expression in Skin and PersistentStaphylococcus aureusNasal Carriage. Journal of Infectious Diseases, 2016, 214, 1744-1751.	1.9	14
26	Alteration of antibiotic regimen as an additional control measure in suspected multi-drug-resistant Enterobacter cloacae outbreak in a neonatal intensive care unit. Journal of Hospital Infection, 2020, 104, 144-149.	1.4	13
27	Inflammatory Response Against Staphylococcus aureus via Intracellular Sensing of Nucleic Acids in Keratinocytes. Frontiers in Immunology, 2022, 13, 828626.	2.2	13
28	Low prevalence of combined linezolid- and vancomycin-resistant Enterococcus faecium from hospital admission screening in an endemic region in Germany. Journal of Global Antimicrobial Resistance, 2020, 22, 646-650.	0.9	12
29	Transmission of ST8-USA300 Latin American Variant Methicillin-Resistant Staphylococcus aureus on a Neonatal Intensive Care Unit: Recurrent Skin and Soft- Tissue Infections as a Marker for Epidemic Community-Associated-MRSA Colonization. Infection Control and Hospital Epidemiology, 2017, 38, 883-885.	1.0	11
30	Nasal colonization with Staphylococcus aureus is a risk factor for ventricular assist device infection in the first year after implantation: A prospective, single-centre, cohort study. Journal of Infection, 2020, 80, 511-518.	1.7	11
31	Emergence of carbapenem-resistant ST131 Escherichia coli carrying blaOXA-244 in Germany, 2019 to 2020. Eurosurveillance, 2020, 25, .	3.9	11
32	Host genetic loci LZTFL1 and CCL2 associated with SARS-CoV-2 infection and severity of COVID-19. International Journal of Infectious Diseases, 2022, 122, 427-436.	1.5	11
33	Challenges in interpretation of WGS and epidemiological data to investigate nosocomial transmission of vancomycin-resistant Enterococcus faecium in an endemic region: incorporation of patient movement network and admission screening. Journal of Antimicrobial Chemotherapy, 2020, 75, 1716-1721.	1.3	10
34	Comparative evaluation of the effect of different growth media on in vitro sensitivity to azithromycin in multi-drug resistant Pseudomonas aeruginosa isolated from cystic fibrosis patients. Antimicrobial Resistance and Infection Control, 2020, 9, 197.	1.5	9
35	A new sequential animal model for infection-related non-unions with segmental bone defect. BMC Musculoskeletal Disorders, 2020, 21, 329.	0.8	9
36	Genomic structure of ST8-t008 USA300 and USA300-LV MRSA in the Rhine-Neckar Region, Germany, 2012–2018. International Journal of Antimicrobial Agents, 2021, 57, 106312.	1.1	9

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37	Impact of discontinuing contact precautions and enforcement of basic hygiene measures on nosocomial vancomycin-resistant Enterococcus faecium transmission. Journal of Hospital Infection, 2022, 121, 120-127.	1.4	9
38	Pitfalls in genotypic antimicrobial susceptibility testing caused by low expression of <i>bla</i> KPC in <i>Escherichia coli</i> . Journal of Antimicrobial Chemotherapy, 2021, 76, 2795-2801.	1.3	8
39	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
40	Genomic Investigation and Successful Containment of an Intermittent Common Source Outbreak of OXA-48-Producing Enterobacter cloacae Related to Hospital Shower Drains. Microbiology Spectrum, 2021, 9, e0138021.	1.2	8
41	Notch Ligand Delta-Like 1 Is Associated With Loss of Vascular Endothelial Barrier Function. Frontiers in Physiology, 2021, 12, 766713.	1.3	8
42	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
43	Identification and Elimination of the Clinically Relevant Multi-Resistant Environmental Bacteria Ralstonia insidiosa in Primary Cell Culture. Microorganisms, 2020, 8, 1599.	1.6	6
44	Systemic Administration of PTH Supports Vascularization in Segmental Bone Defects Filled with Ceramic-Based Bone Graft Substitute. Cells, 2021, 10, 2058.	1.8	6
45	Imported Panton-valentine leucocidin (PVL)-positive <i>Staphylococcus aureus</i> skin infections: patients' perspective on quality of life and quality of medical care. Journal of Travel Medicine, 2022, 29, .	1.4	6
46	A Trial of Antibiotics for Smaller Skin Abscesses. New England Journal of Medicine, 2017, 377, e36.	13.9	5
47	Molecular Detection of Carbapenemases in Enterobacterales: A Comparison of Real-Time Multiplex PCR and Whole-Genome Sequencing. Antibiotics, 2021, 10, 726.	1.5	5
48	Direct-PCR from rectal swabs and environmental reservoirs: A fast and efficient alternative to detect blaOXA-48 carbapenemase genes in an Enterobacter cloacae outbreak setting. Environmental Research, 2022, 203, 111808.	3.7	5
49	Draft Genome Sequence of Staphylococcus aureus Strain HD1410, Isolated from a Persistent Nasal Carrier. Genome Announcements, 2018, 6, .	0.8	4
50	Maternal Vaginal Colonization and Extended-Spectrum Beta-Lactamase-Producing Bacteria in Vietnamese Pregnant Women. Antibiotics, 2021, 10, 572.	1.5	4
51	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
52	Treatment of Infection-Related Non-Unions with Bioactive Glass—A Promising Approach or Just Another Method of Dead Space Management?. Materials, 2022, 15, 1697.	1.3	4
53	Invasiveness of Escherichia coli Is Associated with an IncFII Plasmid. Pathogens, 2021, 10, 1645.	1.2	3
54	Phenotypic Detection of Hemin-Inducible Trimethoprim-Sulfamethoxazole Heteroresistance in Staphylococcus aureus. Microbiology Spectrum, 2021, 9, e0151021.	1.2	2

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55	Whole-genome sequencing disproves two suspected transmission events of blaNDM between Pseudomonas aeruginosa and Enterobacterales in hospitalized patients. Journal of Hospital Infection, 2020, 106, 372-375.	1.4	1
56	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
57	Acinetobacter baumannii. , 2022, , 113-129.		1
58	Comparative Genomic Reveals Clonal Heterogeneity in Persistent Staphylococcus aureus Infection. Frontiers in Cellular and Infection Microbiology, 2022, 12, 817841.	1.8	1
59	Cefiderocol Protects against Cytokine- and Endotoxin-Induced Disruption of Vascular Endothelial Cell Integrity in an In Vitro Experimental Model. Antibiotics, 2022, 11, 581.	1.5	1
60	Reply to Mimica. Clinical Infectious Diseases, 2012, 54, 1518-1519.	2.9	0
61	The Identification of the SARS-CoV-2 Whole Genome: Nine Cases Among Patients in Banten Province, Indonesia. Journal of Pure and Applied Microbiology, 2021, 15, 936-948.	0.3	0
62	Existing evidence supports clinical trials on interventions preventing ventricular assist device infection in patients colonized with <scp><i>Staphylococcus aureus</i></scp> . Letter regarding the article â€Association of preoperative infections, nasal <scp><i>Staphylococcus aureus</i></scp> colonization and gut microbiota with left ventricular assist device outcomes'. European Journal of	2.9	0
63	Heart Failure, 2021, 23, 1566-1566. Staphylococcus massiliensis isolated from human blood cultures, Germany, 2017–2020. European Journal of Clinical Microbiology and Infectious Diseases, 2022, 41, 663-669.	1.3	Ο