

Lutz von MÃ¼ller

List of Publications by Year in descending order

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45
papers

1,430
citations

304602

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48
docs citations

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times ranked

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#	ARTICLE	IF	CITATIONS
1	Quality assurance for genotyping and resistance testing of <i>Clostridium</i> (<i>Clostridioides</i>) <i>difficile</i> isolates - Experiences from the first inter-laboratory ring trial in four German speaking countries. <i>Anaerobe</i> , 2020, 61, 102093.	1.0	10
2	LRPAP1 is a frequent proliferation-inducing antigen of BCRs of mantle cell lymphomas and can be used for specific therapeutic targeting. <i>Leukemia</i> , 2019, 33, 148-158.	3.3	23
3	Integrating Culture-based Antibiotic Resistance Profiles with Whole-genome Sequencing Data for 11,087 Clinical Isolates. <i>Genomics, Proteomics and Bioinformatics</i> , 2019, 17, 169-182.	3.0	8
4	Hospital outbreak due to <i>Clostridium difficile</i> ribotype 018 (RT018) in Southern Germany. <i>International Journal of Medical Microbiology</i> , 2019, 309, 189-193.	1.5	16
5	Comparing genome versus proteome-based identification of clinical bacterial isolates. <i>Briefings in Bioinformatics</i> , 2018, 19, bbw122.	3.2	7
6	Molecular characterization, toxin detection and resistance testing of human clinical <i>Clostridium difficile</i> isolates from Lebanon. <i>International Journal of Medical Microbiology</i> , 2018, 308, 358-363.	1.5	36
7	Prevalence and Strain Characterization of <i>Clostridioides</i> (<i>Clostridium</i>) <i>difficile</i> in Representative Regions of Germany, Ghana, Tanzania and Indonesia – A Comparative Multi-Center Cross-Sectional Study. <i>Frontiers in Microbiology</i> , 2018, 9, 1843.	1.5	26
8	Community-Associated <i>Staphylococcus aureus</i> from Sub-Saharan Africa and Germany: A Cross-Sectional Geographic Correlation Study. <i>Scientific Reports</i> , 2017, 7, 154.	1.6	26
9	Molecular Characterization of Community Acquired <i>Staphylococcus aureus</i> Bacteremia in Young Children in Southern Mozambique, 2001–2009. <i>Frontiers in Microbiology</i> , 2017, 8, 730.	1.5	7
10	Multicentre investigation of carbapenemase-producing <i>Escherichia coli</i> and <i>Klebsiella pneumoniae</i> in German hospitals. <i>International Journal of Medical Microbiology</i> , 2016, 306, 415-420.	1.5	47
11	Detecting <i>Staphylococcus aureus</i> Virulence and Resistance Genes: a Comparison of Whole-Genome Sequencing and DNA Microarray Technology. <i>Journal of Clinical Microbiology</i> , 2016, 54, 1008-1016.	1.8	40
12	Experiences and Lessons from a Multicountry NIDIAG Study on Persistent Digestive Disorders in the Tropics. <i>PLoS Neglected Tropical Diseases</i> , 2016, 10, e0004818.	1.3	11
13	Methicillin-Resistant <i>Staphylococcus aureus</i> in Saarland, Germany: The Long-Term Care Facility Study. <i>PLoS ONE</i> , 2016, 11, e0153030.	1.1	19
14	An assessment on DNA microarray and sequence-based methods for the characterization of methicillin-susceptible <i>Staphylococcus aureus</i> from Nigeria. <i>Frontiers in Microbiology</i> , 2015, 6, 1160.	1.5	19
15	Molecular and culture-based diagnosis of <i>Clostridium difficile</i> isolates from Côte d'Ivoire after prolonged storage at disrupted cold chain conditions. <i>Transactions of the Royal Society of Tropical Medicine and Hygiene</i> , 2015, 109, 660-668.	0.7	8
16	Real-time PCR for detection of <i>Strongyloides stercoralis</i> in human stool samples from Côte d'Ivoire: Diagnostic accuracy, inter-laboratory comparison and patterns of hookworm co-infection. <i>Acta Tropica</i> , 2015, 150, 210-217.	0.9	48
17	Diagnosis of neglected tropical diseases among patients with persistent digestive disorders (diarrhoea and/or abdominal pain ≥14 days): a multi-country, prospective, non-experimental case-control study. <i>BMC Infectious Diseases</i> , 2015, 15, 338.	1.3	16
18	Whole genome sequence typing and microarray profiling of nasal and blood stream methicillin-resistant <i>Staphylococcus aureus</i> isolates: Clues to phylogeny and invasiveness. <i>Infection, Genetics and Evolution</i> , 2015, 36, 475-482.	1.0	7

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19	Epidemiology of Clostridium difficile in Germany based on a single center long-term surveillance and German-wide genotyping of recent isolates provided to the advisory laboratory for diagnostic reasons. International Journal of Medical Microbiology, 2015, 305, 807-813.	1.5	29
20	Tracing the Spread of Clostridium difficile Ribotype 027 in Germany Based on Bacterial Genome Sequences. PLoS ONE, 2015, 10, e0139811.	1.1	40
21	In Reply. Deutsches Ärzteblatt International, 2015, 112, 346.	0.6	0
22	Clostridium Difficile Infection. Deutsches Ärzteblatt International, 2014, 111, 723-31.	0.6	70
23	A Pseudo-Outbreak due to Acinetobacter Species (GIM-1) Contamination of the Pneumatic Transport System of a Large University Hospital. Infection Control and Hospital Epidemiology, 2014, 35, 1364-1372.	1.0	5
24	Underdiagnosis of Clostridium difficile across Europe: the European, multicentre, prospective, biannual, point-prevalence study of Clostridium difficile infection in hospitalised patients with diarrhoea (EUCLID). Lancet Infectious Diseases, The, 2014, 14, 1208-1219.	4.6	308
25	LA-MRSA CC398 differ from classical community acquired-MRSA and hospital acquired-MRSA lineages: Functional analysis of infection and colonization processes. International Journal of Medical Microbiology, 2014, 304, 777-786.	1.5	38
26	Persistent digestive disorders in the tropics: causative infectious pathogens and reference diagnostic tests. BMC Infectious Diseases, 2013, 13, 37.	1.3	69
27	Staphylococcal disease in Africa: another neglected "tropical" disease. Future Microbiology, 2013, 8, 17-26.	1.0	26
28	Analysis of the Airway Microbiota of Healthy Individuals and Patients with Chronic Obstructive Pulmonary Disease by T-RFLP and Clone Sequencing. PLoS ONE, 2013, 8, e68302.	1.1	90
29	Methicillin-Resistant Staphylococcus aureus in Saarland, Germany: A Statewide Admission Prevalence Screening Study. PLoS ONE, 2013, 8, e73876.	1.1	30
30	Matched-Cohort DNA Microarray Diversity Analysis of Methicillin Sensitive and Methicillin Resistant Staphylococcus aureus Isolates from Hospital Admission Patients. PLoS ONE, 2012, 7, e52487.	1.1	30
31	Blockade of gC1qR/p33, a receptor for C1q, inhibits adherence of Staphylococcus aureus to the microvascular endothelium. Microvascular Research, 2011, 82, 66-72.	1.1	14
32	Lues maligna in a patient with unknown HIV infection. BMJ Case Reports, 2011, 2011, bcr0520114221-bcr0520114221.	0.2	14
33	Direct inhibitory effects of Ganciclovir on ICAM-1 expression and proliferation in human coronary vascular cells (SI/MPL-ratio: >1). Medical Science Monitor, 2011, 17, PI1-PI6.	0.5	4
34	Leukocytoclastic vasculitis and myocardial infarction as presenting manifestations of infective endocarditis: a case report. Clinical Research in Cardiology, 2010, 99, 59-61.	1.5	4
35	Characterisation of Clostridium difficile isolates by slpA and tcdC gene sequencing. International Journal of Antimicrobial Agents, 2009, 33, S13-S18.	1.1	27
36	Human cytomegalovirus infection and antiviral immunity in septic patients without canonical immunosuppression. Medical Microbiology and Immunology, 2008, 197, 75-82.	2.6	17

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37	Pharmacokinetics and Pharmacodynamics of Mycophenolic Acid After Enteric-Coated Mycophenolate Versus Mycophenolate Mofetil in Patients With Progressive IgA Nephritis. <i>Journal of Clinical Pharmacology</i> , 2007, 47, 850-859.	1.0	22
38	Cellular Immunity and Active Human Cytomegalovirus Infection in Patients with Septic Shock. <i>Journal of Infectious Diseases</i> , 2007, 196, 1288-1295.	1.9	65
39	Sirolimus inhibits key events of restenosis in vitro/ex vivo: evaluation of the clinical relevance of the data by SI/MPL- and SI/DES-ratio's. <i>BMC Cardiovascular Disorders</i> , 2007, 7, 15.	0.7	4
40	Effects of abciximab on key pattern of human coronary restenosis in vitro: impact of the SI/MPL-ratio. <i>BMC Cardiovascular Disorders</i> , 2006, 6, 14.	0.7	3
41	Severe graft rejection, increased immunosuppression, and active CMV infection in renal transplantation. <i>Journal of Medical Virology</i> , 2006, 78, 394-399.	2.5	19
42	Effects of mycophenolate mofetil on key pattern of coronary restenosis: a cascade of in vitro and ex vivo models. <i>BMC Cardiovascular Disorders</i> , 2005, 5, 9.	0.7	8
43	Bacteremia in an Immunocompromised Patient Caused by a Commensal <i>Neisseria meningitidis</i> Strain Harboring the Capsule Null Locus (cnl). <i>Journal of Clinical Microbiology</i> , 2004, 42, 2898-2901.	1.8	46
44	Polyoma virus-associated interstitial nephritis in a patient with acute myeloid leukaemia and peripheral blood stem cell transplantation. <i>Nephrology Dialysis Transplantation</i> , 2003, 18, 2431-2433.	0.4	37
45	High Variability between Results of Different In-House Tests for Cytomegalovirus (CMV) Monitoring and a Standardized Quantitative Plasma CMV PCR Assay. <i>Journal of Clinical Microbiology</i> , 2002, 40, 2285-2287.	1.8	31