Martin Kaase

List of Publications by Year in descending order

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414414 471509 1,224 32 17 32 citations h-index g-index papers 32 32 32 1892 citing authors all docs docs citations times ranked

#	Article	IF	CITATIONS
1	NDM-2 carbapenemase in Acinetobacter baumannii from Egypt. Journal of Antimicrobial Chemotherapy, 2011, 66, 1260-1262.	3.0	189
2	An outbreak of carbapenem-resistant OXA-48 – producing Klebsiella pneumonia associated to duodenoscopy. Antimicrobial Resistance and Infection Control, 2015, 4, 8.	4.1	121
3	Genome-based analysis of Carbapenemase-producing Klebsiella pneumoniae isolates from German hospital patients, 2008-2014. Antimicrobial Resistance and Infection Control, 2018, 7, 62.	4.1	100
4	Fosfomycin Susceptibility in Carbapenem-Resistant Enterobacteriaceae from Germany. Journal of Clinical Microbiology, 2014, 52, 1893-1897.	3.9	90
5	Identification of molecularly defined Staphylococcus aureus strains using matrix-assisted laser desorption/ionization time of flight mass spectrometry and the Biotyper 2.0 database. Journal of Medical Microbiology, 2010, 59, 787-790.	1.8	87
6	Detection of Carbapenemases in Enterobacteriaceae by a Commercial Multiplex PCR. Journal of Clinical Microbiology, 2012, 50, 3115-3118.	3.9	79
7	What caused the outbreak of ESBL-producing Klebsiella pneumoniae in a neonatal intensive care unit, Germany 2009 to 2012? Reconstructing transmission with epidemiological analysis and whole-genome sequencing. BMJ Open, 2015, 5, e007397-e007397.	1.9	62
8	IS26-Mediated Transfer of blaNDM–1 as the Main Route of Resistance Transmission During a Polyclonal, Multispecies Outbreak in a German Hospital. Frontiers in Microbiology, 2019, 10, 2817.	3.5	57
9	Multicentre investigation of carbapenemase-producing Escherichia coli and Klebsiella pneumoniae in German hospitals. International Journal of Medical Microbiology, 2016, 306, 415-420.	3.6	47
10	The Washing Machine as a Reservoir for Transmission of Extended-Spectrum-Beta-Lactamase (CTX-M-15)-Producing Klebsiella oxytoca ST201 to Newborns. Applied and Environmental Microbiology, 2019, 85, .	3.1	41
11	Whole Genome Sequence Analysis of CTX-M-15 Producing Klebsiella Isolates Allowed Dissecting a Polyclonal Outbreak Scenario. Frontiers in Microbiology, 2018, 9, 322.	3.5	40
12	Comparison of Phenotypic Tests and an Immunochromatographic Assay and Development of a New Algorithm for Detection of OXA-48-like Carbapenemases. Journal of Clinical Microbiology, 2017, 55, 877-883.	3.9	33
13	Description of the metallo- \hat{l}^2 -lactamase GIM-1 in Acinetobacter pittii. Journal of Antimicrobial Chemotherapy, 2014, 69, 81-84.	3.0	31
14	Stalking a lethal superbug by whole-genome sequencing and phylogenetics: Influence on unraveling a major hospital outbreak of carbapenem-resistant Klebsiella pneumoniae. American Journal of Infection Control, 2018, 46, 54-59.	2.3	27
15	Cross-border comparison of the Dutch and German guidelines on multidrug-resistant Gram-negative microorganisms. Antimicrobial Resistance and Infection Control, 2015, 4, 7.	4.1	25
16	Evaluation of species-specific score cut-off values for various Staphylococcus species using a MALDI Biotyper-based identification. Journal of Medical Microbiology, 2012, 61, 1409-1416.	1.8	21
17	Genetic and biochemical characterization of HMB-1, a novel subclass B1 metallo- \hat{l}^2 -lactamase found in a < i > Pseudomonas aeruginosa < / i > clinical isolate. Journal of Antimicrobial Chemotherapy, 2017, 72, dkw554.	3.0	18
18	Molecular epidemiology of VIM-1 producing Escherichia coli from Germany referred to the National Reference Laboratory. International Journal of Medical Microbiology, 2015, 305, 784-789.	3.6	17

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19	Performance of MicroScan WalkAway and Vitek 2 for Detection of Oxacillin Resistance in a Set of Methicillin-Resistant <i>Staphylococcus aureus < i> Isolates with Diverse Genetic Backgrounds. Journal of Clinical Microbiology, 2009, 47, 2623-2625.</i>	3.9	15
20	Dissemination ofblaOXA-58inProteus mirabilisisolates from Germany. Journal of Antimicrobial Chemotherapy, 2017, 72, dkw566.	3.0	15
21	Community-acquired adult Escherichia coli meningitis leading to diagnosis of unrecognized retropharyngeal abscess and cervical spondylodiscitis: a case report. BMC Infectious Diseases, 2015, 15, 567.	2.9	14
22	Novel multiplex PCRs for detection of the most prevalent carbapenemase genes in Gram-negative bacteria within Germany. Journal of Medical Microbiology, 2021, 70, .	1.8	14
23	Description of IMP-31, a novel metallo- \hat{l}^2 -lactamase found in an ST235 <i>Pseudomonas aeruginosa </i> strain in Western Germany. Journal of Antimicrobial Chemotherapy, 2015, 70, 1973-1980.	3.0	13
24	Complete Nucleotide Sequence of a Citrobacter freundii Plasmid Carrying KPC-2 in a Unique Genetic Environment. Genome Announcements, 2014, 2, .	0.8	12
25	False non-susceptible results of tigecycline susceptibility testing against Enterobacteriaceae by an automated system: a multicentre study. Journal of Medical Microbiology, 2016, 65, 877-881.	1.8	11
26	Carbapenem-resistant Gram-negative bacteria - analysis of the data obtained through a mandatory reporting system in the Rhine-Main region, Germany, 2012-2015. GMS Hygiene and Infection Control, 2016, 11, Doc10.	0.3	10
27	Genome Analysis of the Carbapenem- and Colistin-Resistant Escherichia coli Isolate NRZ14408 Reveals Horizontal Gene Transfer Pathways towards Panresistance and Enhanced Virulence. Antimicrobial Agents and Chemotherapy, 2017, 61, .	3.2	9
28	Characterization of mutations in <i>Escherichia coli</i> PBP2 leading to increased carbapenem MICs. Journal of Antimicrobial Chemotherapy, 2019, 74, 571-576.	3.0	9
29	Protracted Regional Dissemination of GIM-1-Producing Serratia marcescens in Western Germany. Antimicrobial Agents and Chemotherapy, 2017, 61, .	3.2	6
30	MRGN: New classification for multidrug-resistant Gram-negative bacteria. Laboratoriums Medizin, 2016, 37, .	0.6	6
31	Overestimation of an Outbreak of Enterobacter cloacae in a Neonatal Intensive Care Unit in Germany, 2015. Pediatric Infectious Disease Journal, 2019, 38, 631-637.	2.0	3
32	MRGN: neue Klassifikation für multiresistente gramnegative Bakterien. Laboratoriums Medizin, 2013, 37,	0.6	2