Sebastian Haller

List of Publications by Year in descending order

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37 papers 8,780 citations

377584 21 h-index 36 g-index

40 all docs

40 docs citations

40 times ranked

7682 citing authors

#	Article	IF	Citations
1	Socioeconomic position and SARS-CoV-2 infections: seroepidemiological findings from a German nationwide dynamic cohort. Journal of Epidemiology and Community Health, 2022, 76, 350-353.	2.0	23
2	Global burden of bacterial antimicrobial resistance in 2019: a systematic analysis. Lancet, The, 2022, 399, 629-655.	6.3	4,915
3	SARS-CoV-2 outbreaks in hospitals and long-term care facilities in Germany: a national observational study. Lancet Regional Health - Europe, The, 2022, 14, 100303.	3.0	17
4	Combined Clinical, Epidemiological, and Genome-Based Analysis Identified a Nationwide Outbreak of <i>Burkholderia cepacia</i> Complex Infections Caused by Contaminated Mouthwash Solutions. Open Forum Infectious Diseases, 2022, 9, ofac114.	0.4	4
5	Bugs That Can Resist Antibiotics but Not Men: Gender-Specific Differences in Notified Infections and Colonisations in Germany, 2010–2019. Microorganisms, 2021, 9, 894.	1.6	9
6	Socioeconomic inequalities in the prevalence and perceived dangerousness of SARS-CoV-2 infections in two early German hotspots: findings from a seroepidemiological study. BMC Research Notes, 2021, 14, 375.	0.6	11
7	Healthcare-associated foodborne outbreaks in high-income countries: a literature review and surveillance study, 16 OECD countries, 2001 to 2019*. Eurosurveillance, 2021, 26, .	3.9	10
8	Seroepidemiological study on the spread of SARS-CoV-2 in populations in especially affected areas in Germany - Study protocol of the CORONA-MONITORING lokal study , 2020, 5, 2-16.		12
9	Cross-border spread of blaNDM-1- and blaOXA-48-positive Klebsiella pneumoniae: a European collaborative analysis of whole genome sequencing and epidemiological data, 2014 to 2019. Eurosurveillance, 2020, 25, .	3.9	26
10	Rapid spread of OXA-244-producing Escherichia coli ST38 in Germany: insights from an integrated molecular surveillance approach; 2017 to January 2020. Eurosurveillance, 2020, 25, .	3.9	16
11	Providing actionable evidence in Public Health - The 2018 international workshop on evidence-based public health at the Robert Koch Institute, Berlin , 2020, 5, 3-6.		1
12	Summary of World Café Discussions Table 2: Evaluation , 2020, 5, 22.		1
13	The rise in vancomycin-resistant Enterococcus faecium in Germany: data from the German Antimicrobial Resistance Surveillance (ARS). Antimicrobial Resistance and Infection Control, 2019, 8, 147.	1.5	83
14	Overestimation of an Outbreak of Enterobacter cloacae in a Neonatal Intensive Care Unit in Germany, 2015. Pediatric Infectious Disease Journal, 2019, 38, 631-637.	1.1	3
15	Attributable deaths and disability-adjusted life-years caused by infections with antibiotic-resistant bacteria in the EU and the European Economic Area in 2015: a population-level modelling analysis. Lancet Infectious Diseases, The, 2019, 19, 56-66.	4.6	1,908
16	Application of a new methodology and R package reveals a high burden of healthcare-associated infections (HAI) in Germany compared to the average in the European Union/European Economic Area, 2011 to 2012. Eurosurveillance, 2019, 24, .	3.9	20
17	Extensively drug-resistant Klebsiella pneumoniae ST307 outbreak, north-eastern Germany, June to October 2019. Eurosurveillance, 2019, 24, .	3.9	46
18	Routine screening for colonization by Gram-negative bacteria in neonates at intensive care units for the prediction of sepsis: systematic review and meta-analysis. Journal of Hospital Infection, 2018, 99, 367-380.	1.4	27

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19	Sepsis prediction during outbreaks at neonatal intensive care units through body surface screening for Gram-negative bacteria: systematic review and meta-analysis. BMC Research Notes, 2018, 11, 917.	0.6	2
20	Healthcare-associated pneumonia in acute care hospitals in European Union/European Economic Area countries: an analysis of data from a point prevalence survey, 2011 to 2012. Eurosurveillance, 2018, 23, .	3.9	48
21	Whole Genome Sequence Analysis of CTX-M-15 Producing Klebsiella Isolates Allowed Dissecting a Polyclonal Outbreak Scenario. Frontiers in Microbiology, 2018, 9, 322.	1.5	40
22	Global outbreak of severe Mycobacterium chimaera disease after cardiac surgery: a molecular epidemiological study. Lancet Infectious Diseases, The, 2017, 17, 1033-1041.	4.6	198
23	Burden of Six Healthcare-Associated Infections on European Population Health: Estimating Incidence-Based Disability-Adjusted Life Years through a Population Prevalence-Based Modelling Study. PLoS Medicine, 2016, 13, e1002150.	3.9	436
24	Contamination during production of heater-cooler units by Mycobacterium chimaera potential cause for invasive cardiovascular infections: results of an outbreak investigation in Germany, April 2015 to February 2016. Eurosurveillance, 2016, 21, .	3.9	109
25	Neurological sequelae of healthcare-associated sepsis in very-low-birthweight infants: Umbrella review and evidence-based outcome tree. Eurosurveillance, 2016, 21, 30143.	3.9	30
26	Effectiveness of Routine and Booster Pertussis Vaccination in Children and Adolescents, Federal State of Brandenburg, Germany, 2002–2012. Pediatric Infectious Disease Journal, 2015, 34, 513-519.	1.1	24
27	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.	0.8	62
28	Incidence of invasive meticillin-resistant Staphylococcus aureus infections in Germany, 2010 to 2014. Eurosurveillance, 2015, 20, .	3.9	18
29	How many outbreaks of nosocomial infections occur in German neonatal intensive care units annually?. Infection, 2014, 42, 73-78.	2.3	25
30	Results from the First 12 Months of the National Surveillance of Healthcare Associated Outbreaks in Germany, 2011/2012. PLoS ONE, 2014, 9, e98100.	1.1	21
31	Large multistate outbreak of norovirus gastroenteritis associated with frozen strawberries, Germany, 2012. Eurosurveillance, 2014, 19, 20719.	3.9	109
32	Polymorphisms in FTO and MAF Genes and Birth Weight, BMI, Ponderal Index, Weight Gain in a Large Cohort of Infants with a Birth Weight below 1500 Grams. PLoS ONE, 2013, 8, e66331.	1.1	8
33	Avoidance of mechanical ventilation by surfactant treatment of spontaneously breathing preterm infants (AMV): an open-label, randomised, controlled trial. Lancet, The, 2011, 378, 1627-1634.	6.3	408
34	Association of <i>Filaggrin</i> lossâ€ofâ€functionâ€mutations with atopic dermatitis and asthma in the Early Treatment of the Atopic Child (ETAC) population. Pediatric Allergy and Immunology, 2009, 20, 358-361.	1.1	27
35	<i>IL13</i> variants are associated with total serum IgE and early sensitization to food allergens in children with atopic dermatitis. Pediatric Allergy and Immunology, 2009, 20, 551-555.	1.1	24
36	<i>MBL2</i> variants in relation to common childhood infections and atopyâ€related phenotypes in a large German birth cohort. Pediatric Allergy and Immunology, 2007, 18, 665-670.	1.1	16

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#	ARTICLE	IF	CITATIONS
37	No association of histamine- N-methyltransferase polymorphism with asthma or bronchial hyperresponsiveness in two German pediatric populations. Pediatric Allergy and Immunology, 2005, 16, 40-42.	1.1	29