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List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3510561/publications.pdf

Version: 2024-02-01

21 papers

2,502 citations

16 h-index 19 g-index

26 all docs

26 docs citations

26 times ranked 2209 citing authors

#	Article	IF	Citations
1	Changes in the incidence of invasive disease due to Streptococcus pneumoniae, Haemophilus influenzae, and Neisseria meningitidis during the COVID-19 pandemic in 26 countries and territories in the Invasive Respiratory Infection Surveillance Initiative: a prospective analysis of surveillance data. The Lancet Digital Health, 2021, 3, e360-e370.	5.9	260
2	Prophages and satellite prophages are widespread in Streptococcus and may play a role in pneumococcal pathogenesis. Nature Communications, 2019, 10, 4852.	5.8	64
3	Vaccination of Icelandic Children with the 10-Valent Pneumococcal Vaccine Leads to a Significant Herd Effect among Adults in Iceland. Journal of Clinical Microbiology, 2019, 57, .	1.8	16
4	Genomic Analyses of >3,100 Nasopharyngeal Pneumococci Revealed Significant Differences Between Pneumococci Recovered in Four Different Geographical Regions. Frontiers in Microbiology, 2019, 10, 317.	1.5	9
5	Effect of Vaccination on Pneumococci Isolated from the Nasopharynx of Healthy Children and the Middle Ear of Children with Otitis Media in Iceland. Journal of Clinical Microbiology, 2018, 56, .	1.8	26
6	Genome Sequencing Reveals a Large and Diverse Repertoire of Antimicrobial Peptides. Frontiers in Microbiology, 2018, 9, 2012.	1.5	34
7	Diverse Streptococcus pneumoniae Strains Drive a Mucosal-Associated Invariant T-Cell Response Through Major Histocompatibility Complex class l–Related Molecule–Dependent and Cytokine-Driven Pathways. Journal of Infectious Diseases, 2018, 217, 988-999.	1.9	59
8	Pneumococcal prophages are diverse, but not without structure or history. Scientific Reports, 2017, 7, 42976.	1.6	62
9	Putatively novel serotypes and the potential for reduced vaccine effectiveness: capsular locus diversity revealed among 5405 pneumococcal genomes. Microbial Genomics, 2016, 2, 000090.	1.0	41
10	Genomics Reveals the Worldwide Distribution of Multidrug-Resistant Serotype 6E Pneumococci. Journal of Clinical Microbiology, 2015, 53, 2271-2285.	1.8	44
11	Genomic analyses of pneumococci reveal a wide diversity of bacteriocins – including pneumocyclicin, a novel circular bacteriocin. BMC Genomics, 2015, 16, 554.	1.2	67
12	Evolutionary and Population Biology of Streptococcus Pneumoniae., 2014, , 117-135.		6
13	Defining the Estimated Core Genome of Bacterial Populations Using a Bayesian Decision Model. PLoS Computational Biology, 2014, 10, e1003788.	1.5	72
14	Population Genetic Structure of Streptococcus pneumoniae in Kilifi, Kenya, Prior to the Introduction of Pneumococcal Conjugate Vaccine. PLoS ONE, 2013, 8, e81539.	1.1	20
15	Pneumococcal genome sequencing tracks a vaccine escape variant formed through a multi-fragment recombination event. Nature Genetics, 2012, 44, 352-355.	9.4	144
16	Temporal and Geographic Stability of the Serogroupâ€Specific Invasive Disease Potential ofStreptococcus pneumoniaein Children. Journal of Infectious Diseases, 2004, 190, 1203-1211.	1.9	312
17	Clonal Relationships between Invasive and CarriageStreptococcus pneumoniaeand Serotype―and Cloneâ€Specific Differences in Invasive Disease Potential. Journal of Infectious Diseases, 2003, 187, 1424-1432.	1.9	563
18	Characterization of German penicillin non-susceptible serotype 23F pneumococci using multilocus sequence typing. Journal of Medical Microbiology, 2003, 52, 981-987.	0.7	10

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19	Geographic Distribution and Clonal Diversity of Streptococcus pneumoniae Serotype 1 Isolates. Journal of Clinical Microbiology, 2003, 41, 4966-4970.	1.8	131
20	Antimicrobial Resistance among Clinical Isolates of Streptococcus pneumoniae in the United States during 1999–2000, Including a Comparison of Resistance Rates since 1994–1995. Antimicrobial Agents and Chemotherapy, 2001, 45, 1721-1729.	1.4	523
21	Pneumococcal Carriage. , 0, , 136-147.		10