

# Franz Allerberger

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2416310/publications.pdf>

Version: 2024-02-01

18  
papers

1,047  
citations

759233

12  
h-index

839539

18  
g-index

23  
all docs

23  
docs citations

23  
times ranked

2386  
citing authors

#	ARTICLE	IF	CITATIONS
1	Diversity of West Nile and Usutu virus strains in mosquitoes at an international airport in Austria. <i>Transboundary and Emerging Diseases</i> , 2022, 69, 2096-2109.	3.0	10
2	First records of <i>Aedes pulcritarsis</i> (Rondani, 1872) (Diptera: Culicidae) in Austria. <i>Parasitology Research</i> , 2022, 121, 765-768.	1.6	2
3	Viral variant-resolved wastewater surveillance of SARS-CoV-2 at national scale. <i>Nature Biotechnology</i> , 2022, 40, 1814-1822.	17.5	82
4	SARS-CoV-2 mutations in MHC-I-restricted epitopes evade CD8 <sup>+</sup> T cell responses. <i>Science Immunology</i> , 2021, 6, .	11.9	143
5	Multiplexed detection of SARS-CoV-2 and other respiratory infections in high throughput by SARSeq. <i>Nature Communications</i> , 2021, 12, 3132.	12.8	32
6	Presence of $\beta$ -Lactamase-producing Enterobacterales and Salmonella Isolates in Marine Mammals. <i>International Journal of Molecular Sciences</i> , 2021, 22, 5905.	4.1	10
7	The First Report of mcr-1-Carrying <i>Escherichia coli</i> Originating from Animals in Serbia. <i>Antibiotics</i> , 2021, 10, 1063.	3.7	7
8	Population structure-guided profiling of antibiotic resistance patterns in clinical <i>Listeria monocytogenes</i> isolates from Germany identifies <i>pbpB3</i> alleles associated with low levels of cephalosporin resistance. <i>Emerging Microbes and Infections</i> , 2020, 9, 1804-1813.	6.5	18
9	Genomic epidemiology of superspreading events in Austria reveals mutational dynamics and transmission properties of SARS-CoV-2. <i>Science Translational Medicine</i> , 2020, 12, .	12.4	203
10	Large Nationwide Outbreak of Invasive Listeriosis Associated with Blood Sausage, Germany, 2018–2019. <i>Emerging Infectious Diseases</i> , 2020, 26, 1456-1464.	4.3	40
11	Broad-Spectrum Cephalosporin-Resistant <i>Klebsiella</i> spp. Isolated from Diseased Horses in Austria. <i>Animals</i> , 2020, 10, 332.	2.3	15
12	Whole genome sequencing reveals resemblance between ESBL-producing and carbapenem resistant <i>Klebsiella pneumoniae</i> isolates from Austrian rivers and clinical isolates from hospitals. <i>Science of the Total Environment</i> , 2019, 662, 227-235.	8.0	60
13	A Syndrome-Based Surveillance System for Infectious Diseases Among Asylum Seekers in Austrian Reception Centers, 2015-2018: Analysis of Reported Data. <i>JMIR Public Health and Surveillance</i> , 2019, 5, e11465.	2.6	4
14	Integrated analysis of human-animal-vector surveillance: West Nile virus infections in Austria, 2015–2016. <i>Emerging Microbes and Infections</i> , 2018, 7, 1-15.	6.5	22
15	Characterization of a community-acquired-MRSA USA300 isolate from a river sample in Austria and whole genome sequence based comparison to a diverse collection of USA300 isolates. <i>Scientific Reports</i> , 2018, 8, 9467.	3.3	22
16	Usutu virus infections among blood donors, Austria, July and August 2017 – Raising awareness for diagnostic challenges. <i>Eurosurveillance</i> , 2017, 22, .	7.0	57
17	West Nile Virus Positive Blood Donation and Subsequent Entomological Investigation, Austria, 2014. <i>PLoS ONE</i> , 2015, 10, e0126381.	2.5	24
18	Defining and Evaluating a Core Genome Multilocus Sequence Typing Scheme for Whole-Genome Sequence-Based Typing of <i>Listeria monocytogenes</i> . <i>Journal of Clinical Microbiology</i> , 2015, 53, 2869-2876.	3.9	266