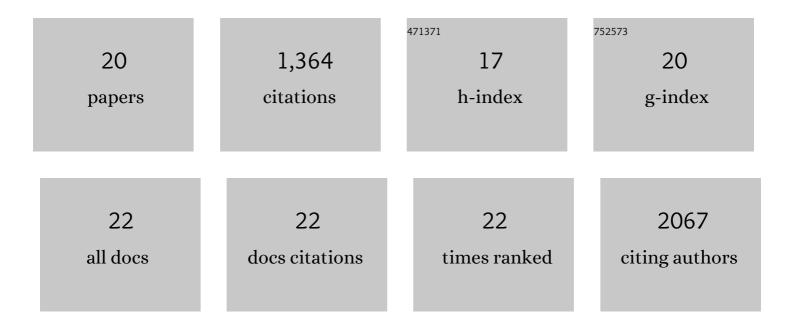
Lejla Imamovic

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7977042/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Expansion and persistence of antibiotic-specific resistance genes following antibiotic treatment. Gut Microbes, 2021, 13, 1-19.	4.3	24
2	Forecasting the dissemination of antibiotic resistance genes across bacterial genomes. Nature Communications, 2021, 12, 2435.	5.8	111
3	Mining, analyzing, and integrating viral signals from metagenomic data. Microbiome, 2019, 7, 42.	4.9	58
4	Complete Genome Sequence of <i>Escherichia coli</i> Strain WG5. Genome Announcements, 2018, 6, .	0.8	10
5	Drug-Driven Phenotypic Convergence Supports Rational Treatment Strategies of Chronic Infections. Cell, 2018, 172, 121-134.e14.	13.5	131
6	Bluephage: A rapid method for the detection of somatic coliphages used as indicators of fecal pollution in water. Water Research, 2018, 128, 10-19.	5.3	29
7	The Environmental Exposures and Inner- and Intercity Traffic Flows of the Metro System May Contribute to the Skin Microbiome and Resistome. Cell Reports, 2018, 24, 1190-1202.e5.	2.9	56
8	Rapid resistome mapping using nanopore sequencing. Nucleic Acids Research, 2017, 45, gkw1328.	6.5	62
9	Heterogeneity in phage induction enables the survival of the lysogenic population. Environmental Microbiology, 2016, 18, 957-969.	1.8	28
10	BaeSR, Involved in Envelope Stress Response, Protects against Lysogenic Conversion by Shiga Toxin 2-Encoding Phages. Infection and Immunity, 2015, 83, 1451-1457.	1.0	4
11	Evolution of a Self-Inducible Cytolethal Distending Toxin Type V-Encoding Bacteriophage from Escherichia coli O157:H7 to Shigella sonnei. Journal of Virology, 2013, 87, 13665-13675.	1.5	18
12	Use of Collateral Sensitivity Networks to Design Drug Cycling Protocols That Avoid Resistance Development. Science Translational Medicine, 2013, 5, 204ra132.	5.8	368
13	Characterizing RecA-Independent Induction of Shiga toxin2-Encoding Phages by EDTA Treatment. PLoS ONE, 2012, 7, e32393.	1.1	87
14	Bacteriophages and genetic mobilization in sewage and faecally polluted environments. Microbial Biotechnology, 2011, 4, 725-734.	2.0	40
15	Quantification and Evaluation of Infectivity of Shiga Toxin-Encoding Bacteriophages in Beef and Salad. Applied and Environmental Microbiology, 2011, 77, 3536-3540.	1.4	28
16	Bacteriophages Carrying Antibiotic Resistance Genes in Fecal Waste from Cattle, Pigs, and Poultry. Antimicrobial Agents and Chemotherapy, 2011, 55, 4908-4911.	1.4	136
17	Quantification of Shiga toxin 2â€encoding bacteriophages, by realâ€time PCR and correlation with phage infectivity. Journal of Applied Microbiology, 2010, 108, 1105-1114.	1.4	24
18	Quantification of Shiga Toxin-Converting Bacteriophages in Wastewater and in Fecal Samples by Real-Time Quantitative PCR. Applied and Environmental Microbiology, 2010, 76, 5693-5701.	1.4	58

#	Article	IF	CITATIONS
19	OI-57, a Genomic Island of <i>Escherichia coli</i> O157, Is Present in Other Seropathotypes of Shiga Toxin-Producing <i>E. coli</i> Associated with Severe Human Disease. Infection and Immunity, 2010, 78, 4697-4704.	1.0	35
20	Phage-Mediated Shiga Toxin 2 Gene Transfer in Food and Water. Applied and Environmental Microbiology, 2009, 75, 1764-1768.	1.4	55