Adriano Di Pasquale

List of Publications by Year
in descending order

16 papers	**141** citations	1478505 **6** h-index	1281871 **11** g-index
17 all docs	**17** docs citations	**17** times ranked	**249** citing authors

#	Article	IF	Citations
1	A â€œOne-Healthâ€•approach for diagnosis and molecular characterization of SARS-CoV-2 in Italy. One Health, 2020, 10, 100135.	3.4	46
2	Hypo- and Hyper-Virulent Listeria monocytogenes Clones Persisting in Two Different Food Processing Plants of Central Italy. Microorganisms, 2021, 9, 376.	3.6	25
3	SARS-CoV-2 surveillance in Italy through phylogenomic inferences based on Hamming distances derived from pan-SNPs, -MNPs and -InDels. BMC Genomics, 2021, 22, 782.	2.8	12
4	Whole Genome Sequencing Characterization of HEV3-e and HEV3-f Subtypes among the Wild Boar Population in the Abruzzo Region, Italy: First Report. Microorganisms, 2020, 8, 1393.	3.6	11
5	Chapter 6: Access Methods and Query Processing Techniques. Lecture Notes in Computer Science, 2003, , 203-261.	1.3	8
6	Whole-Genome Sequence of a Reemerging Listeria monocytogenes Serovar 1/2a Strain in Central Italy. Microbiology Resource Announcements, 2018, 7, .	0.6	8
7	Proteomic analysis of Brucella melitensis and Brucella ovis for identification of virulence factor using bioinformatics approachs. Molecular and Cellular Probes, 2020, 53, 101581.	2.1	8
8	In vitro and in silico parameters for precise cgMLST typing of Listeria monocytogenes. BMC Genomics, 2022, 23, 235.	2.8	7
9	Distributed Searching of k-Dimensional Data with Almost Constant Costs. Lecture Notes in Computer Science, 2000, , 239-250.	1.3	6
10	Whole-Genome Sequence of Listeria monocytogenes Serovar 1/2a Strain IZSAM_Lm_15_17439_A144, Representative of a Human Outbreak in 2008. Microbiology Resource Announcements, 2019, 8, .	0.6	3
11	ADST: An Order Preserving Scalable Distributed Data Structure with Constant Access Costs. Lecture Notes in Computer Science, 2001, , 211-222.	1.3	2
12	A Very Efficient Order Preserving Scalable Distributed Data Structure. Lecture Notes in Computer Science, 2001, , 186-199.	1.3	2
13	Quality of service in wireless networks. , 0, , .		1
14	Draft Genome Sequences of 19 Salmonella enterica Serovar Typhimurium [4,5:i:â˜˜] Strains Resistant to Nalidixic Acid from a Long-Term Outbreak in Italy. Genome Announcements, 2015, 3, .	0.8	1
15	Development of food safety risk assessment tools based on molecular typing and WGS of CampylobacterÂjejuni genome. EFSA Journal, 2019, 17, e170903.	1.8	1
16	Draft Genome Sequence of Listeria monocytogenes Serovar 1/2a Strain IZSAM_Lm_14-16064, Isolated from an Italian Cooked Ham in 2014. Microbiology Resource Announcements, 2020, 9, .	0.6	0