

Lorena Sacchini

List of Publications by Year in descending order

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19
papers

444
citations

759233

12
h-index

794594

19
g-index

22
all docs

22
docs citations

22
times ranked

685
citing authors

#	ARTICLE	IF	CITATIONS
1	Core Genome Multilocus Sequence Typing and Single Nucleotide Polymorphism Analysis in the Epidemiology of <i>Brucella melitensis</i> Infections. <i>Journal of Clinical Microbiology</i> , 2018, 56, .	3.9	58
2	Antimicrobial resistance genotypes and phenotypes of <i>Campylobacter jejuni</i> isolated in Italy from humans, birds from wild and urban habitats, and poultry. <i>PLoS ONE</i> , 2019, 14, e0223804.	2.5	55
3	Distribution of <i>Brucella</i> field strains isolated from livestock, wildlife populations, and humans in Italy from 2007 to 2015. <i>PLoS ONE</i> , 2019, 14, e0213689.	2.5	47
4	Characterization of Antimicrobial Resistance Patterns and Detection of Virulence Genes in <i>Campylobacter</i> Isolates in Italy. <i>Sensors</i> , 2014, 14, 3308-3322.	3.8	46
5	Cases of human brucellosis in Sweden linked to Middle East and Africa. <i>BMC Research Notes</i> , 2016, 9, 277.	1.4	32
6	Phylogeography and epidemiology of <i>Brucella suis</i> biovar 2 in wildlife and domestic swine. <i>Veterinary Microbiology</i> , 2019, 233, 68-77.	1.9	29
7	Origins and global context of <i>Brucella abortus</i> in Italy. <i>BMC Microbiology</i> , 2017, 17, 28.	3.3	28
8	Rapid and safe one-step extraction method for the identification of <i>Brucella</i> strains at genus and species level by MALDI-TOF mass spectrometry. <i>PLoS ONE</i> , 2018, 13, e0197864.	2.5	25
9	Whole Genome Sequencing for Tracing Geographical Origin of Imported Cases of Human Brucellosis in Sweden. <i>Microorganisms</i> , 2019, 7, 398.	3.6	23
10	Outbreak of unusual <i>Salmonella enterica</i> serovar Typhimurium monophasic variant 1,4 [5],12:i:-, Italy, June 2013 to September 2014. <i>Eurosurveillance</i> , 2016, 21, .	7.0	22
11	Detection of methicillin-resistant <i>Staphylococcus aureus</i> in dairy cow farms. <i>Food Control</i> , 2014, 46, 532-538.	5.5	17
12	First isolation, identification and genetic characterization of <i>Brucella abortus</i> biovar 3 from dairy cattle in Bangladesh. <i>Veterinary Medicine and Science</i> , 2019, 5, 556-562.	1.6	11
13	Detection and Genotyping of <i>Campylobacter jejuni</i> and <i>Campylobacter coli</i> by Use of DNA Oligonucleotide Arrays. <i>Molecular Biotechnology</i> , 2013, 53, 182-188.	2.4	10
14	First outbreak of food poisoning caused by <i>Salmonella enterica</i> subspecies <i>enterica</i> serovar Berta in Italy. <i>Letters in Applied Microbiology</i> , 2012, 55, 122-127.	2.2	7
15	Molecular typing of <i>Salmonella enterica</i> subspecies <i>enterica</i> serovar Typhimurium isolated in Abruzzo region (Italy) from 2008 to 2010. <i>Veterinaria Italiana</i> , 2014, 50, 31-9.	0.5	7
16	The prevalence, characterisation, and antimicrobial resistance of <i>Yersinia enterocolitica</i> in pigs from Central Italy. <i>Veterinaria Italiana</i> , 2018, 54, 115-123.	0.5	6
17	Variability of the microbiota in traditional Caciocavallo, Scamorza and Caciotta cheeses manufactured with raw milk and natural cultures. <i>International Journal of Dairy Technology</i> , 2021, 74, 564-574.	2.8	3
18	Characterization of Lactic Acid Bacteria in Pecorino di Farindola Cheese and Manufacturing with a <i>Lactisacibacillus paracasei</i> Autochthonous Culture. <i>Applied Sciences (Switzerland)</i> , 2021, 11, 7897.	2.5	3

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19	Draft Genome Sequences of 19 Salmonella enterica Serovar Typhimurium [4,5:i:âˆ²] Strains Resistant to Nalidixic Acid from a Long-Term Outbreak in Italy. Genome Announcements, 2015, 3, .	0.8	1