Sohail Naushad

List of Publications by Year
in descending order

18	1,873	471509 17	794594 19
papers	citations	h-index	g-index

19	19	19	2174
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Non-aureus Staphylococci and Bovine Udder Health: Current Understanding and Knowledge Gaps. Frontiers in Veterinary Science, 2021, 8, 658031.	2.2	52
2	Genomic Analysis of Bovine Staphylococcus aureus Isolates from Milk To Elucidate Diversity and Determine the Distributions of Antimicrobial and Virulence Genes and Their Association with Mastitis. MSystems, 2020, 5, .	3.8	35
3	A New Whole Genome Culture-Independent Diagnostic Test (WG-CIDT) for Rapid Detection of Salmonella in Lettuce. Frontiers in Microbiology, 2020, 11, 602.	3.5	8
4	Comprehensive Virulence Gene Profiling of Bovine Non- aureus Staphylococci Based on Whole-Genome Sequencing Data. MSystems, 2019, 4, .	3.8	32
5	Staphylococcus debuckii sp. nov., a coagulase-negative species from bovine milk. International Journal of Systematic and Evolutionary Microbiology, 2019, 69, 2239-2249.	1.7	10
6	Virulence gene profiles: alpha-hemolysin and clonal diversity in Staphylococcus aureus isolates from bovine clinical mastitis in China. BMC Veterinary Research, 2018, 14, 63.	1.9	38
7	Associations between digital dermatitis lesion grades in dairy cattle and the quantities of four Treponema species. Veterinary Research, 2018, 49, 111.	3.0	28
8	Prevalence and Genetic Basis of Antimicrobial Resistance in Non-aureus Staphylococci Isolated from Canadian Dairy Herds. Frontiers in Microbiology, 2018, 9, 256.	3.5	52
9	Prevalence of non-aureus staphylococci species causing intramammary infections in Canadian dairy herds. Journal of Dairy Science, 2017, 100, 5592-5612.	3.4	70
10	Comparison of treatment records and inventory of empty drug containers to quantify antimicrobial usage in dairy herds. Journal of Dairy Science, 2017, 100, 9736-9745.	3.4	44
11	Bacteriocins of Non-aureus Staphylococci Isolated from Bovine Milk. Applied and Environmental Microbiology, 2017, 83, .	3.1	46
12	Comprehensive Phylogenetic Analysis of Bovine Non-aureus Staphylococci Species Based on Whole-Genome Sequencing. Frontiers in Microbiology, 2016, 7, 1990.	3.5	49
13	A phylogenomic reappraisal of family-level divisions within the class Halobacteria: proposal to divide the order Halobacteriales into the families Halobacteriaceae, Haloarculaceae fam. nov., and Halococcaceae fam. nov., and the order Haloferacales into the families, Haloferacaceae and Halorubraceae fam nov.. Antonie Van Leeuwenhoek, 2016, 109, 565-587.	1.7	127
14	Genome-based phylogeny and taxonomy of the â€˜Enterobacterialesâ€™: proposal for Enterobacterales ord. nov. divided into the families Enterobacteriaceae, Erwiniaceae fam. nov., Pectobacteriaceae fam. nov., Yersiniaceae fam. nov., Hafniaceae fam. nov., Morganellaceae fam. nov., and Budviciaceae fam. nov.. International Journal of Systematic and Evolutionary Microbiology, 2016, 66, 5575-5599.	1.7	792
15	Phylogenomic and Molecular Demarcation of the Core Members of the PolyphyleticPasteurellaceaeGeneraActinobacillus,Haemophilus, andPasteurella. International Journal of Genomics, 2015, 2015, 1-15.	1.6	19
16	A phylogenomic and molecular marker based taxonomic framework for the order Xanthomonadales: proposal to transfer the families Algiphilaceae and Solimonadaceae to the order Nevskiales ord. nov. and to create a new family within the order Xanthomonadales, the family Rhodanobacteraceae fam. nov., containing the genus Rhodanobacter and its closest relatives. Antonie Van Leeuwenhoek, 2015, 107, 467-485.	1.7	135
17	A phylogenomic and molecular markers based analysis of the phylum Chlamydiae: proposal to divide the class Chlamydiia into two orders, Chlamydiales and Parachlamydiales ord. nov., and emended description of the class Chlamydiia. Antonie Van Leeuwenhoek, 2015, 108, 765-781.	1.7	38
18	Phylogenomic analyses and molecular signatures for the class Halobacteria and its two major clades: a proposal for division of the class Halobacteria into an emended order Halobacteriales and two new orders, Haloferacales ord. nov. and Natrialbales ord. nov., containing the novel families Haloferacaceae fam. nov. and Natrialbaceae fam. nov.. International Journal of Systematic and Evolutionary Microbiology, 2015, 65, 1050-1069.	1.7	260