

Franciele M Siqueira

List of Publications by Year in descending order

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56
papers

827
citations

687363
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56
all docs

56
docs citations

56
times ranked

1099
citing authors

#	ARTICLE	IF	CITATIONS
1	Biodegradation potential of oily sludge by pure and mixed bacterial cultures. <i>Bioresource Technology</i> , 2011, 102, 11003-11010.	9.6	238
2	New insights on the biology of swine respiratory tract mycoplasmas from a comparative genome analysis. <i>BMC Genomics</i> , 2013, 14, 175.	2.8	63
3	Antimicrobial activity of propolis extract against <i>Staphylococcus coagulase positive</i> and <i>Malassezia pachydermatis</i> of canine otitis. <i>Veterinary Microbiology</i> , 2010, 142, 432-434.	1.9	44
4	Genomic insights into the versatility of the plant growth-promoting bacterium <i>Azospirillum amazonense</i> . <i>BMC Genomics</i> , 2011, 12, 409.	2.8	43
5	Bioprospection and selection of bacteria isolated from environments contaminated with petrochemical residues for application in bioremediation. <i>World Journal of Microbiology and Biotechnology</i> , 2012, 28, 1203-1222.	3.6	36
6	Unravelling the Transcriptome Profile of the Swine Respiratory Tract Mycoplasmas. <i>PLoS ONE</i> , 2014, 9, e110327.	2.5	34
7	Insights on the virulence of swine respiratory tract mycoplasmas through genome-scale metabolic modeling. <i>BMC Genomics</i> , 2016, 17, 353.	2.8	34
8	Microbiome overview in swine lungs. <i>PLoS ONE</i> , 2017, 12, e0181503.	2.5	33
9	<i>Mycoplasma</i> non-coding RNA: identification of small RNAs and targets. <i>BMC Genomics</i> , 2016, 17, 743.	2.8	27
10	Influence of vaccine strains on the evolution of canine distemper virus. <i>Infection, Genetics and Evolution</i> , 2016, 41, 262-269.	2.3	23
11	Molecular Characterization of <i>Rhodococcus equi</i> from Horse-Breeding Farms by Means of Multiplex PCR for the <i>vap</i> Gene Family. <i>Current Microbiology</i> , 2009, 58, 399-403.	2.2	21
12	<i>Mannheimia haemolytica</i> pleuropneumonia in goats associated with shipping stress. <i>Ciencia Rural</i> , 2019, 49, .	0.5	17
13	Molecular and phylogenetic analyses of <i>Salmonella Gallinarum</i> trace the origin and diversification of recent outbreaks of fowl typhoid in poultry farms. <i>Veterinary Microbiology</i> , 2017, 212, 80-86.	1.9	16
14	Genome analysis reveals insights into high-resistance and virulence of <i>Salmonella Enteritidis</i> involved in foodborne outbreaks. <i>International Journal of Food Microbiology</i> , 2019, 306, 108269.	4.7	15
15	<i>Mycoplasma hyopneumoniae</i> Transcription Unit Organization: Genome Survey and Prediction. <i>DNA Research</i> , 2011, 18, 413-422.	3.4	12
16	Intrinsic terminators in <i>Mycoplasma hyopneumoniae</i> transcription. <i>BMC Genomics</i> , 2015, 16, 273.	2.8	12
17	Characterization of Lactic Acid Bacteria in Raw Buffalo Milk: a Screening for Novel Probiotic Candidates and Their Transcriptional Response to Acid Stress. <i>Probiotics and Antimicrobial Proteins</i> , 2021, 13, 468-483.	3.9	11
18	Virulence factors, antimicrobial resistance, and plasmid content of <i>Escherichia coli</i> isolated in swine commercial farms. <i>Arquivo Brasileiro De Medicina Veterinaria E Zootecnia</i> , 2010, 62, 30-36.	0.4	10

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19	Evaluation of growth and gene expression of <i>Mycoplasma hyopneumoniae</i> and <i>Mycoplasma hyorhinis</i> in defined medium. <i>Molecular Biology Reports</i> , 2018, 45, 2469-2479.	2.3	10
20	Pet Pyometra: Correlating Bacteria Pathogenicity to Endometrial Histological Changes. <i>Pathogens</i> , 2021, 10, 833.	2.8	10
21	Insights on the genetic features of endometrial pathogenic <i>Escherichia coli</i> strains from pyometra in companion animals: Improving the knowledge about pathogenesis. <i>Infection, Genetics and Evolution</i> , 2020, 85, 104453.	2.3	9
22	Genome organization in <i>Mycoplasma hyopneumoniae</i> : identification of promoter-like sequences. <i>Molecular Biology Reports</i> , 2014, 41, 5395-5402.	2.3	8
23	Virulence factors and antimicrobial resistance of <i>Escherichia coli</i> isolated from urinary tract of swine in southern of Brazil. <i>Brazilian Journal of Microbiology</i> , 2008, 39, 741-743.	2.0	7
24	Repetitive Elements in <i>Mycoplasma hyopneumoniae</i> Transcriptional Regulation. <i>PLoS ONE</i> , 2016, 11, e0168626.	2.5	7
25	<i>Chromobacterium violaceum</i> Infection in a Horse. <i>Journal of Comparative Pathology</i> , 2017, 156, 334-338.	0.4	6
26	Histomorphometric study of the anterior latissimus dorsi muscle and evaluation of enzymatic markers of broilers affected with dorsal cranial myopathy. <i>Poultry Science</i> , 2017, 96, 4217-4223.	3.4	6
27	Evaluation of the serum virome in calves persistently infected with Pestivirus A, presenting or not presenting mucosal disease. <i>Virus Genes</i> , 2018, 54, 768-778.	1.6	6
28	Omphalitis, urachocystitis and septicemia by <i>Streptococcus dysgalactiae</i> in a southern right whale calf <i>Eubalaena australis</i> , Brazil. <i>Diseases of Aquatic Organisms</i> , 2018, 131, 227-232.	1.0	6
29	Perception of poultry veterinarians on the use of antimicrobials and antimicrobial resistance in egg production. <i>Poultry Science</i> , 2022, 101, 101987.	3.4	6
30	Global analysis of sRNA target genes in <i>Mycoplasma hyopneumoniae</i> . <i>BMC Genomics</i> , 2018, 19, 767.	2.8	5
31	Caution at choosing a particular colony-forming unit from faecal <i>Escherichia coli</i> : it may not represent the sample profile. <i>Letters in Applied Microbiology</i> , 2020, 70, 130-136.	2.2	5
32	Clonality of <i>Mycoplasma hyopneumoniae</i> in swine farms from Brazil. <i>Veterinary Microbiology</i> , 2019, 238, 108434.	1.9	4
33	Frozen bovine preputial mucus as a suitable sample for the direct molecular diagnosis of <i>Campylobacter fetus</i> subsp. <i>venerealis</i> . <i>Journal of Microbiological Methods</i> , 2020, 179, 106101.	1.6	4
34	Fibrinonecrotic Placentitis and Abortion Associated With <i>Pantoea agglomerans</i> Infection in a Mare. <i>Journal of Equine Veterinary Science</i> , 2020, 92, 103156.	0.9	4
35	Survey of beef bulls in Brazil to assess their role as source of infectious agents related to cow infertility. <i>Journal of Veterinary Diagnostic Investigation</i> , 2022, 34, 54-60.	1.1	4
36	Wild capybaras as reservoir of shiga toxin-producing <i>Escherichia coli</i> in urban Amazonian Region. <i>Letters in Applied Microbiology</i> , 2022, 75, 10-16.	2.2	4

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37	Draft Genome Sequence of the d -Xylose-Fermenting Yeast <i>Spathaspora xylofermentans</i> UFMG-HMD23.3. Genome Announcements, 2017, 5, .	0.8	3
38	Genome sequencing of two <i>Bacillus anthracis</i> strains: a virulent strain and a vaccinal strain. Brazilian Journal of Microbiology, 2018, 49, 18-19.	2.0	3
39	Bovine abortion associated with <i>Staphylococcus aureus</i> infection - characterization of <i>S. aureus</i> strain isolated from fetal tissues. Ciencia Rural, 2020, 50, .	0.5	3
40	Fatal systemic <i>Mortierella wolfii</i> infection in a neonatal calf in southern Brazil. Ciencia Rural, 2020, 50, .	0.5	3
41	Genome Sequence of <i>Mycoplasma hyorhinis</i> Isolated from Cell Cultures. Genome Announcements, 2016, 4, .	0.8	2
42	Draft Genome Sequence of <i>Acholeplasma laidlawii</i> , a Common Contaminant of Cell Cultures. Genome Announcements, 2017, 5, .	0.8	2
43	Non- <i>Lactose</i> -fermenting uropathogenic <i>Escherichia coli</i> from dogs: virulence profile characterization. Letters in Applied Microbiology, 2021, 72, 596-603.	2.2	2
44	The influence of regulatory elements on <i>Mycoplasma hyopneumoniae</i> 7448 transcriptional response during oxidative stress and heat shock. Molecular Biology Reports, 2022, 49, 139-147.	2.3	2
45	Cows' reproductive performances and parity order influences the cervicovaginal fungal community. Microbial Pathogenesis, 2022, 162, 105351.	2.9	2
46	Multidrug-resistant <i>Escherichia coli</i> from free-living pigeons (<i>Columba livia</i>): Insights into antibiotic environmental contamination and detection of resistance genes. Zoonoses and Public Health, 2022, 69, 682-693.	2.2	2
47	Fibrinous pleuropneumonia caused by <i>Pasteurella multocida</i> associated with bovine lymphoma. Ciencia Rural, 2018, 48, .	0.5	1
48	Clinical and microbiological characterization of subclinical bacteriuria and sporadic bacterial cystitis in dogs with spontaneous hypercortisolism. Comparative Immunology, Microbiology and Infectious Diseases, 2021, 75, 101624.	1.6	1
49	Phylogenetic and pathotype analysis of <i>Escherichia coli</i> swine isolates from Southern Brazil. Pesquisa Veterinaria Brasileira, 2012, 32, 374-378.	0.5	1
50	Functional characterization of the putative FAD synthase from <i>Mycoplasma hyopneumoniae</i> . FEMS Microbiology Letters, 2021, 368, .	1.8	0
51	Sequencing and phylogenetic analysis of <i>Clostridium septicum</i> alpha toxin gene from Brazilian field and vaccine strains. African Journal of Microbiology Research, 2012, 6, .	0.4	0
52	Bovine abortion by a vaccine strain of <i>Bacillus anthracis</i> . Ciencia Rural, 2020, 50, .	0.5	0
53	<i>Campylobacter fetus</i> in Abomasal Fluid from Spontaneously Aborted Bovine and Ovine Fetuses. Acta Scientiae Veterinariae, 0, 49, .	0.2	0
54	Molecular detection of respiratory coinfections in pig herds with enzootic pneumonia: a survey in Brazil. Journal of Veterinary Diagnostic Investigation, 2022, , 104063872110695.	1.1	0

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55	Genetic and molecular Omp25 analyses from worldwide <i>Brucella canis</i> strains: Possible mutational influences in protein function. <i>Gene</i> , 2022, 817, 146175.	2.2	0
56	Aerossaccolitis and Pneumonia in an Indian Peafowl Caused by <i>Lactobacillus agilis</i> . <i>Acta Scientiae Veterinariae</i> , 0, 50, .	0.2	0