Fabiana Quoos Mayer

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6857278/publications.pdf

Version: 2024-02-01

98 papers

1,493 citations

19 h-index 395702 33 g-index

102 all docs

102 docs citations

times ranked

102

2239 citing authors

#	Article	IF	CITATIONS
1	Detection of coronavirus in vampire bats (<i>Desmodus rotundus</i>) in southern Brazil. Transboundary and Emerging Diseases, 2022, 69, 2384-2389.	3.0	18
2	Genomic characterization and production of antimicrobial lipopeptides by Bacillus velezensis P45 growing on feather by-products. Journal of Applied Microbiology, 2022, 132, 2067-2079.	3.1	4
3	Cows' reproductive performances and parity order influences the cervicovaginal fungal community. Microbial Pathogenesis, 2022, 162, 105351.	2.9	2
4	Short-chain fatty acid acetate triggers antiviral response mediated by RIC-I in cells from infants with respiratory syncytial virus bronchiolitis. EBioMedicine, 2022, 77, 103891.	6.1	37
5	The virome of the white-winged vampire bat Diaemus youngi is rich in circular DNA viruses. Virus Genes, 2022, 58, 214-226.	1.6	1
6	Multidrugâ€resistant <i>Escherichia coli</i> from freeâ€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
7	Molecular survey of porcine respiratory disease complex pathogens in Brazilian wild boars. Preventive Veterinary Medicine, 2022, 206, 105698.	1.9	5
8	Complete genome characterization of porcine circovirus 3 recovered from wild boars in Southern Brazil. Transboundary and Emerging Diseases, 2021, 68, 240-247.	3.0	5
9	A plate of viruses: Viral metagenomics of supermarket chicken, pork and beef from Brazil. Virology, 2021, 552, 1-9.	2.4	16
10	Detection of multiple viruses in oropharyngeal samples from Brazilian free-tailed bats (Tadarida) Tj ETQq0 0 0 rgE	3T /Overloo 2.1	ck 10 Tf 50 38
11	Tuberculosis outbreak in intensive swine farming from southern Brazil. Ciencia Rural, 2021, 51, .	0.5	O
12	Genomic analysis on Brazilian strains of Anaplasma marginale. Brazilian Journal of Veterinary Parasitology, 2021, 30, e000421.	0.7	3
13	Molecular identification of Mycobacterium spp. isolated from Brazilian wild boars. Molecular Biology Reports, 2021, 48, 1025-1031.	2.3	3
14	A new highly divergent copiparvovirus in sheep. Archives of Virology, 2021, 166, 1517-1520.	2.1	2
15	Molecular characterization of the bacterial communities present in sheep's milk and cheese produced in South Brazilian Region via 16S rRNA gene metabarcoding sequencing. LWT - Food Science and Technology, 2021, 147, 111579.	5.2	11
16	The genetic diversity of "papillomavirome―in bovine teat papilloma lesions. Animal Microbiome, 2021, 3, 51.	3.8	5
17	Investigation of Mycobacterium bovis and Metastrongylus sp. co-infection and its relationship to tuberculosis lesions' occurrence in wild boars. Comparative Immunology, Microbiology and Infectious Diseases, 2021, 77, 101674.	1.6	1
18	Spleen and lung virome analysis of South American fur seals (Arctocephalus australis) collected on the southern Brazilian coast. Infection, Genetics and Evolution, 2021, 92, 104862.	2.3	2

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19	Antimicrobial Resistance of Coagulase-positive Staphylococcus Isolated From Healthy Crioulo Horses and Associated Risk Factors. Journal of Equine Veterinary Science, 2021, 107, 103779.	0.9	3
20	Molecular characterization of bacterial communities of two neotropical tick species (Amblyomma) Tj ETQq0 0 0 r 2021, 12, 101746.	gBT /Overl 2.7	lock 10 Tf 50 5
21	SARS-CoV-2 introduction and lineage dynamics across three epidemic peaks in Southern Brazil: massive spread of P.1. Infection, Genetics and Evolution, 2021, 96, 105144.	2.3	14
22	A variety of highly divergent eukaryotic ssDNA viruses in sera of pigs. Journal of General Virology, 2021, 102, .	2.9	3
23	Clinicopathological characteristics and papillomavirus types in cutaneous warts in bovine. Brazilian Journal of Microbiology, 2020, 51, 395-401.	2.0	6
24	Mycobacterium tuberculosis var. tuberculosis infection in two captive black capuchins (Sapajus) Tj ETQq0 0 0 rgt	BT /Overloo	ck ₆ 10 Tf 50 5
25	A molecular strategy to optimize bovine tuberculosis post-mortem diagnosis and the exposure to Mycobacterium tuberculosis variant bovis. Molecular Biology Reports, 2020, 47, 7291-7296.	2.3	4
26	Penile Tuberculosis in a Bull. Journal of Comparative Pathology, 2020, 180, 5-8.	0.4	0
27	Investigation on porcine circovirus type 3 in serum of farrowing sows with stillbirths. Microbial Pathogenesis, 2020, 149, 104316.	2.9	10
28	Virome of crab-eating (Cerdocyon thous) and pampas foxes (Lycalopex gymnocercus) from southern Brazil and Uruguay. Infection, Genetics and Evolution, 2020, 85, 104421.	2.3	11
29	Viral metagenomics in Brazilian Pekin ducks identifies two gyrovirus, including a new species, and the potentially pathogenic duck circovirus. Virology, 2020, 548, 101-108.	2.4	10
30	Canine papillomavirus type 16 associated to squamous cell carcinoma in a dog: virological and pathological findings. Brazilian Journal of Microbiology, 2020, 51, 2087-2094.	2.0	9
31	Diversity of cyclic antimicrobial lipopeptides from Bacillus P34 revealed by functional annotation and comparative genome analysis. Microbiological Research, 2020, 238, 126515.	5.3	17
32	Viral DNA genomes in sera of farrowing sows with or without stillbirths. PLoS ONE, 2020, 15, e0230714.	2.5	11
33	Novel Gyrovirus genomes recovered from free-living pigeons in Southern Brazil. Virology, 2020, 548, 132-135.	2.4	9
34	Insights on the genetic features of endometrial pathogenic Escherichia coli strains from pyometra in companion animals: Improving the knowledge about pathogenesis. Infection, Genetics and Evolution, 2020, 85, 104453.	2.3	9
35	Liver virome of healthy pigs reveals diverse small ssDNA viral genomes. Infection, Genetics and Evolution, 2020, 81, 104203.	2.3	16
36	Phylogenetic and evolutionary analysis of HoBiâ€like pestivirus: Insights into origin and dispersal. Transboundary and Emerging Diseases, 2020, 67, 1909.	3.0	10

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37	Bovine abortion by a vaccine strain of Bacillus anthracis. Ciencia Rural, 2020, 50, .	0.5	O
38	Valvular endocarditis associated with Helcococcus ovis in a cow in Southern Brazil. Ciencia Rural, 2020, 50, .	0.5	0
39	Genome analysis reveals insights into high-resistance and virulence of Salmonella Enteritidis involved in foodborne outbreaks. International Journal of Food Microbiology, 2019, 306, 108269.	4.7	15
40	Microbiota-derived acetate protects against respiratory syncytial virus infection through a GPR43-type 1 interferon response. Nature Communications, 2019, 10, 3273.	12.8	234
41	Characterization of the viral genomes present in commercial batches of horse serum obtained by high-throughput sequencing. Biologicals, 2019, 61, 1-7.	1.4	9
42	Highly divergent cattle hepacivirus N in Southern Brazil. Archives of Virology, 2019, 164, 3133-3136.	2.1	5
43	Genomic mosaicism: A neglected factor that promotes variability in asthma diagnosis. Medical Hypotheses, 2019, 127, 112-115.	1.5	0
44	Finding factors associated with nasal shedding of <i>Mycobacterium tuberculosis</i> variant <i>bovis</i> in wild boar. Veterinary Record, 2019, 185, 627-628.	0.3	0
45	The intestinal virome of malabsorption syndrome-affected and unaffected broilers through shotgun metagenomics. Virus Research, 2019, 261, 9-20.	2.2	64
46	A technical assessment of the porcine ejaculated spermatozoa for a sperm-specific RNA-seq analysis. Systems Biology in Reproductive Medicine, 2018, 64, 291-303.	2.1	45
47	Tuberculosis in Southern Brazilian wild boars (<i>Sus scrofa</i>): First epidemiological findings. Transboundary and Emerging Diseases, 2018, 65, 518-526.	3.0	27
48	Full-Genome Sequence of Porcine Circovirus type 3 recovered from serum of sows with stillbirths in Brazil. Transboundary and Emerging Diseases, 2018, 65, 5-9.	3.0	114
49	Genome sequencing of two Bacillus anthracis strains: a virulent strain and a vaccinal strain. Brazilian Journal of Microbiology, 2018, 49, 18-19.	2.0	3
50	Characterization of dog serum virome from Northeastern Brazil. Virology, 2018, 525, 192-199.	2.4	21
51	Evaluation of the serum virome in calves persistently infected with Pestivirus A, presenting or not presenting mucosal disease. Virus Genes, 2018, 54, 768-778.	1.6	6
52	First detection of Pseudomonas aeruginosa ST2963 from hospital effluent: A draft genome analysis. Journal of Global Antimicrobial Resistance, 2018, 14, 275-276.	2.2	1
53	Columbid circoviruses detected in free ranging pigeons from Southern Brazil: insights on PiCV evolution. Archives of Virology, 2018, 163, 3083-3090.	2.1	11
54	New polyomavirus species identified in nutria, Myocastor coypus polyomavirus 1. Archives of Virology, 2018, 163, 3203-3206.	2.1	5

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55	Mamastrovirus 5 detected in a crab-eating fox (Cerdocyon thous): Expanding wildlife host range of astroviruses. Comparative Immunology, Microbiology and Infectious Diseases, 2018, 58, 36-43.	1.6	6
56	Progressive heart disease in mucopolysaccharidosis type I mice may be mediated by increased cathepsin B activity. Cardiovascular Pathology, 2017, 27, 45-50.	1.6	19
57	Genome sequence of bubaline alphaherpesvirus 1 (BuHV1) isolated in Australia in 1972. Archives of Virology, 2017, 162, 1169-1176.	2.1	8
58	Pathogenic Leptospira spp. in bats: Molecular investigation in Southern Brazil. Comparative Immunology, Microbiology and Infectious Diseases, 2017, 52, 14-18.	1.6	18
59	Genomic characterization of a bovine viral diarrhea virus subtype 1i in Brazil. Archives of Virology, 2017, 162, 1119-1123.	2.1	12
60	Co-occurrence of mcr-1 and blaKPC-2 in a clinical isolate of Escherichia coli in Brazil. Journal of Antimicrobial Chemotherapy, 2017, 72, 2404-2406.	3.0	26
61	Draft Genome Sequence of the d -Xylose-Fermenting Yeast Spathaspora xylofermentans UFMG-HMD23.3. Genome Announcements, 2017, 5, .	0.8	3
62	Genomic and antigenic relationships between two †HoBi†IIII-like strains and other members of the Pestivirus genus. Archives of Virology, 2017, 162, 3025-3034.	2.1	10
63	Genome characterization of a bovine papillomavirus type 5 from cattle in the Amazon region, Brazil. Virus Genes, 2017, 53, 130-133.	1.6	6
64	Draft Genome Sequence of Acholeplasma laidlawii, a Common Contaminant of Cell Cultures. Genome Announcements, $2017, 5, .$	0.8	2
65	Faecal virome of healthy chickens reveals a large diversity of the eukaryote viral community, including novel circular ssDNA viruses. Journal of General Virology, 2017, 98, 690-703.	2.9	50
66	Long-term restoration of alpha-L-iduronidase activity in fibroblasts from patients with mucopolysaccharidosis type I after non-viral gene transfer. Clinical and Biomedical Research, 2017, 37, 330-333.	0.1	0
67	Nasal swab real-time PCR is not suitable for in vivo diagnosis of bovine tuberculosis. Pesquisa Veterinaria Brasileira, 2017, 37, 549-554.	0.5	1
68	Complete genome sequence of Deltapapillomavirus 4 (bovine papillomavirus 2) from a bovine papillomavirus lesion in Amazon Region, Brazil. Memorias Do Instituto Oswaldo Cruz, 2016, 111, 277-279.	1.6	3
69	Deleterious effects of interruption followed by reintroduction of enzyme replacement therapy on a lysosomal storage disorder. Translational Research, 2016, 176, 29-37.e1.	5.0	4
70	Genome Sequence of Mycoplasma hyorhinis Isolated from Cell Cultures. Genome Announcements, 2016, 4, .	0.8	2
71	How many papillomavirus species can go undetected in papilloma lesions?. Scientific Reports, 2016, 6, 36480.	3.3	28
72	Immunomodulator plasmid projected by systems biology as a candidate for the development of adjunctive therapy for respiratory syncytial virus infection. Medical Hypotheses, 2016, 88, 86-90.	1.5	3

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73	Draft Genome Sequence of a Salmonella enterica subsp. enterica Serovar Gallinarum bv. Gallinarum Isolate Associated with Fowl Typhoid Outbreaks in Brazil. Genome Announcements, 2016, 4, .	0.8	2
74	Genotypic and antimicrobial characterization of pathogenic bacteria at different stages of cattle slaughtering in southern Brazil. Meat Science, 2016, 116, 193-200.	5.5	27
75	Novel Bovine Papillomavirus Type Discovered by Rolling-Circle Amplification Coupled with Next-Generation Sequencing. PLoS ONE, 2016, 11, e0162345.	2.5	24
76	A Novel Chiropteran Circovirus Genome Recovered from a Brazilian Insectivorous Bat Species. Genome Announcements, 2015, 3, .	0.8	17
77	Shotgun proteomics reveals possible mechanisms for cognitive impairment in Mucopolysaccharidosis I mice. Molecular Genetics and Metabolism, 2015, 114, 138-145.	1.1	19
78	Detection of Mycobacterium tuberculosis and Mycobacterium avium Complexes by Real-Time PCR in Bovine Milk from Brazilian Dairy Farms. Journal of Food Protection, 2015, 78, 1037-1042.	1.7	9
79	Laronidase-Functionalized Multiple-Wall Lipid-Core Nanocapsules: Promising Formulation for a More Effective Treatment of Mucopolysaccharidosis Type I. Pharmaceutical Research, 2015, 32, 941-954.	3.5	31
80	Full-Genome Sequence of a Reassortant H1N2 Influenza A Virus Isolated from Pigs in Brazil. Genome Announcements, 2014, 2, .	0.8	8
81	Lessons from molecular modeling human \hat{l}_{\pm} -l-iduronidase. Journal of Molecular Graphics and Modelling, 2014, 54, 107-113.	2.4	3
82	Food safety in raw milk production: risk factors associated to bacterial DNA contamination. Tropical Animal Health and Production, 2014, 46, 877-882.	1.4	21
83	Natural occurrence of White spot syndrome virus and Infectious hypodermal and hematopoietic necrosis virus in Neohelice granulata crab. Journal of Invertebrate Pathology, 2013, 114, 86-88.	3.2	21
84	Enzyme replacement therapy started at birth improves outcome in difficult-to-treat organs in mucopolysaccharidosis I mice. Molecular Genetics and Metabolism, 2013, 109, 33-40.	1.1	57
85	Genotypic and phenotypic characterization of Brazilian patients with GM1 gangliosidosis. Gene, 2013, 512, 113-116.	2.2	28
86	Chloramphenicol Enhances IDUA Activity on Fibroblasts from Mucopolysaccharidosis I Patients. Current Pharmaceutical Biotechnology, 2013, 14, 194-198.	1.6	1
87	Genetic Diagnosis in Recently Transfused Patients. Diagnostic Molecular Pathology, 2013, 22, 123-126.	2.1	1
88	Characterization of joint disease in mucopolysaccharidosis type I mice. International Journal of Experimental Pathology, 2013, 94, 305-311.	1.3	19
89	Chloramphenicol Enhances IDUA Activity on Fibroblasts from Mucopolysaccharidosis I Patients. Current Pharmaceutical Biotechnology, 2013, 14, 194-198.	1.6	15
90	Analysis of cDNA Molecules is Not Suitable for the Molecular Diagnosis of Mucopolysaccharidosis Type I. Diagnostic Molecular Pathology, 2012, 21, 53-55.	2.1	3

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91	Recombinant Encapsulated Cells Overexpressing Alpha- <i>L</i> -Iduronidase Correct Enzyme Deficiency in Human Mucopolysaccharidosis Type I Cells. Cells Tissues Organs, 2012, 195, 323-329.	2.3	24
92	Mycobacterium bovis infection in a collared peccary (Tayassu tajacu): Insights on tuberculosis wild reservoirs. Veterinary Microbiology, 2012, 160, 549-551.	1.9	12
93	Evidence of a progressive motor dysfunction in Mucopolysaccharidosis type I mice. Behavioural Brain Research, 2012, 233, 169-175.	2.2	30
94	Intraperitoneal implant of recombinant encapsulated cells overexpressing alpha-l-iduronidase partially corrects visceral pathology in mucopolysaccharidosis type I mice. Cytotherapy, 2012, 14, 860-867.	0.7	17
95	Emerging treatment options for the mucopolysaccharidoses. Research and Reports in Endocrine Disorders, 2012, , 53.	0.4	6
96	Cell microencapsulation: a potential tool for the treatment of neuronopathic lysosomal storage diseases. Journal of Inherited Metabolic Disease, 2011, 34, 983-990.	3.6	16
97	Effects of Cryopreservation and Hypothermic Storage on Cell Viability and Enzyme Activity in Recombinant Encapsulated Cells Overexpressing Alpha‣″duronidase. Artificial Organs, 2010, 34, 434-439.	1.9	17
98	New GLB1 mutation in siblings with Morquio type B disease presenting with mental regression. Molecular Genetics and Metabolism, 2009, 96, 148.	1.1	16