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	Microbiota-derived acetate protects against respiratory syncytial virus infection through a GPR43-type 1 interferon response. Nature Communications, 2019, 10, 3273.	12.8	234
2	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
3	The intestinal virome of malabsorption syndrome-affected and unaffected broilers through shotgun metagenomics. Virus Research, 2019, 261, 9-20.	2.2	64
4	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
5	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
6	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
7	Short-chain fatty acid acetate triggers antiviral response mediated by RIG-I in cells from infants with respiratory syncytial virus bronchiolitis. EBioMedicine, 2022, 77, 103891.	6.1	37
8	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
9	Evidence of a progressive motor dysfunction in Mucopolysaccharidosis type I mice. Behavioural Brain Research, 2012, 233, 169-175.	2.2	30
10	Genotypic and phenotypic characterization of Brazilian patients with GM1 gangliosidosis. Gene, 2013, 512, 113-116.	2.2	28
11	How many papillomavirus species can go undetected in papilloma lesions?. Scientific Reports, 2016, 6, 36480.	3.3	28
12	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
13	Tuberculosis in Southern Brazilian wild boars (<i>Sus scrofa</i>): First epidemiological findings. Transboundary and Emerging Diseases, 2018, 65, 518-526.	3.0	27
14	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
15	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
16	Novel Bovine Papillomavirus Type Discovered by Rolling-Circle Amplification Coupled with Next-Generation Sequencing. PLoS ONE, 2016, 11, e0162345.	2.5	24
17	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
18	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

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19	Characterization of dog serum virome from Northeastern Brazil. Virology, 2018, 525, 192-199.	2.4	21
20	Characterization of joint disease in mucopolysaccharidosis type I mice. International Journal of Experimental Pathology, 2013, 94, 305-311.	1.3	19
21	Shotgun proteomics reveals possible mechanisms for cognitive impairment in Mucopolysaccharidosis I mice. Molecular Genetics and Metabolism, 2015, 114, 138-145.	1.1	19
22	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
23	Pathogenic Leptospira spp. in bats: Molecular investigation in Southern Brazil. Comparative Immunology, Microbiology and Infectious Diseases, 2017, 52, 14-18.	1.6	18
24	Detection of coronavirus in vampire bats (<i>Desmodus rotundus</i>) in southern Brazil. Transboundary and Emerging Diseases, 2022, 69, 2384-2389.	3.0	18
25	Effects of Cryopreservation and Hypothermic Storage on Cell Viability and Enzyme Activity in Recombinant Encapsulated Cells Overexpressing Alpha‣â€Iduronidase. Artificial Organs, 2010, 34, 434-439.	1.9	17
26	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
27	A Novel Chiropteran Circovirus Genome Recovered from a Brazilian Insectivorous Bat Species. Genome Announcements, 2015, 3, .	0.8	17
28	Diversity of cyclic antimicrobial lipopeptides from Bacillus P34 revealed by functional annotation and comparative genome analysis. Microbiological Research, 2020, 238, 126515.	5.3	17
29	Detection of multiple viruses in oropharyngeal samples from Brazilian free-tailed bats (Tadarida) Tj ETQq1 1 0.78	43 <u>1</u> 4 rgB1	7 / Qyerlock (1)
30	New GLB1 mutation in siblings with Morquio type B disease presenting with mental regression. Molecular Genetics and Metabolism, 2009, 96, 148.	1.1	16
31	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
32	Liver virome of healthy pigs reveals diverse small ssDNA viral genomes. Infection, Genetics and Evolution, 2020, 81, 104203.	2.3	16
33	A plate of viruses: Viral metagenomics of supermarket chicken, pork and beef from Brazil. Virology, 2021, 552, 1-9.	2.4	16
34	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
35	Chloramphenicol Enhances IDUA Activity on Fibroblasts from Mucopolysaccharidosis I Patients. Current Pharmaceutical Biotechnology, 2013, 14, 194-198.	1.6	15
36	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

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37	Mycobacterium bovis infection in a collared peccary (Tayassu tajacu): Insights on tuberculosis wild reservoirs. Veterinary Microbiology, 2012, 160, 549-551.	1.9	12
38	Genomic characterization of a bovine viral diarrhea virus subtype 1i in Brazil. Archives of Virology, 2017, 162, 1119-1123.	2.1	12
39	Columbid circoviruses detected in free ranging pigeons from Southern Brazil: insights on PiCV evolution. Archives of Virology, 2018, 163, 3083-3090.	2.1	11
40	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
41	Viral DNA genomes in sera of farrowing sows with or without stillbirths. PLoS ONE, 2020, 15, e0230714.	2.5	11
42	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
43	Genomic and antigenic relationships between two †HoBi†HoBiâ€. It like strains and other members of the Pestivirus genus. Archives of Virology, 2017, 162, 3025-3034.	2.1	10
44	Investigation on porcine circovirus type 3 in serum of farrowing sows with stillbirths. Microbial Pathogenesis, 2020, 149, 104316.	2.9	10
45	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
46	Phylogenetic and evolutionary analysis of HoBiâ€like pestivirus: Insights into origin and dispersal. Transboundary and Emerging Diseases, 2020, 67, 1909.	3.0	10
47	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
48	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
49	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
50	Novel Gyrovirus genomes recovered from free-living pigeons in Southern Brazil. Virology, 2020, 548, 132-135.	2.4	9
51	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
52	Full-Genome Sequence of a Reassortant H1N2 Influenza A Virus Isolated from Pigs in Brazil. Genome Announcements, 2014, 2, .	0.8	8
53	Genome sequence of bubaline alphaherpesvirus 1 (BuHV1) isolated in Australia in 1972. Archives of Virology, 2017, 162, 1169-1176.	2.1	8
54	Emerging treatment options for the mucopolysaccharidoses. Research and Reports in Endocrine Disorders, 2012, , 53.	0.4	6

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55	Genome characterization of a bovine papillomavirus type 5 from cattle in the Amazon region, Brazil. Virus Genes, 2017, 53, 130-133.	1.6	6
56	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
57	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
58	Clinicopathological characteristics and papillomavirus types in cutaneous warts in bovine. Brazilian Journal of Microbiology, 2020, 51, 395-401.	2.0	6
59	Mycobacterium tuberculosis var. tuberculosis infection in two captive black capuchins (Sapajus) Tj ETQq1 1 0.784	1314 rgBT 2.0	 Qverlock 10
60	New polyomavirus species identified in nutria, Myocastor coypus polyomavirus 1. Archives of Virology, 2018, 163, 3203-3206.	2.1	5
61	Highly divergent cattle hepacivirus N in Southern Brazil. Archives of Virology, 2019, 164, 3133-3136.	2.1	5
62	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
63	The genetic diversity of "papillomavirome―in bovine teat papilloma lesions. Animal Microbiome, 2021, 3, 51.	3.8	5
64	Molecular characterization of bacterial communities of two neotropical tick species (Amblyomma) Tj ETQq0 0 0 rg	gBT /Overlo 2.7	ock 10 Tf 50 5
65	Molecular survey of porcine respiratory disease complex pathogens in Brazilian wild boars. Preventive Veterinary Medicine, 2022, 206, 105698.	1.9	5
66	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
67	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
68	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
69	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
70	Lessons from molecular modeling human \hat{l}_{\pm} -l-iduronidase. Journal of Molecular Graphics and Modelling, 2014, 54, 107-113.	2.4	3
71	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
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 the d -Xylose-Fermenting Yeast Spathaspora xylofermentans UFMG-HMD23.3. Genome Announcements, 2017, 5, .	0.8	3
74	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
75	Genomic analysis on Brazilian strains of Anaplasma marginale. Brazilian Journal of Veterinary Parasitology, 2021, 30, e000421.	0.7	3
76	Molecular identification of Mycobacterium spp. isolated from Brazilian wild boars. Molecular Biology Reports, 2021, 48, 1025-1031.	2.3	3
77	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
78	A variety of highly divergent eukaryotic ssDNA viruses in sera of pigs. Journal of General Virology, 2021, 102, .	2.9	3
79	Genome Sequence of Mycoplasma hyorhinis Isolated from Cell Cultures. Genome Announcements, 2016, 4, .	0.8	2
80	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
81	Draft Genome Sequence of Acholeplasma laidlawii, a Common Contaminant of Cell Cultures. Genome Announcements, 2017, 5, .	0.8	2
82	A new highly divergent copiparvovirus in sheep. Archives of Virology, 2021, 166, 1517-1520.	2.1	2
83	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
84	Cows' reproductive performances and parity order influences the cervicovaginal fungal community. Microbial Pathogenesis, 2022, 162, 105351.	2.9	2
85	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
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	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
89	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
90	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

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91	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
92	Genomic mosaicism: A neglected factor that promotes variability in asthma diagnosis. Medical Hypotheses, 2019, 127, 112-115.	1.5	0
93	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	O
94	Penile Tuberculosis in a Bull. Journal of Comparative Pathology, 2020, 180, 5-8.	0.4	0
95	Tuberculosis outbreak in intensive swine farming from southern Brazil. Ciencia Rural, 2021, 51, .	0.5	О
96	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
97	Bovine abortion by a vaccine strain of Bacillus anthracis. Ciencia Rural, 2020, 50, .	0.5	0
98	Valvular endocarditis associated with Helcococcus ovis in a cow in Southern Brazil. Ciencia Rural, 2020, 50, .	0.5	0