Meng Qi

List of Publications by Year
in descending order

Source: https://exaly.com/author-pdf/9413176/publications.pdf

Version: 2024-02-01

65 papers	1,772 citations	218677 26 h-index	276875 41 g-index
65 all docs	65 docs citations	65 times ranked	830 citing authors

#	Article	IF	Citations
1	Molecular detection and genetic characteristics of Cryptosporidium spp. in Chinese racehorses. Equine Veterinary Journal, 2023, 55, 474-480.	1.7	3
2	Molecular characterization of Cryptosporidium spp. and Giardia duodenalis in pet dogs in Xinjiang, China. Parasitology Research, 2022, 121, 1429-1435.	1.6	5
3	Genetic diversity of Blastocystis subtypes in the Alpine musk deer (Moschus) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 6 e12910.	1.7	3
4	Seasonal monitoring of Cryptosporidium species and their genetic diversity in neonatal calves on two largeâ€scale farms in Xinjiang, China. Journal of Eukaryotic Microbiology, 2022, 69, e12878.	1.7	5
5	Molecular detection and genotypes of Enterocytozoon bieneusi in farmed mink (Neovison vison), blue foxes (Alopex lagopus), and raccoon dogs (Nyctereutes procyonoides) in Xinjiang, China. International Journal for Parasitology: Parasites and Wildlife, 2021, 14, 211-215.	1.5	5
6	Common occurrence of Enterocytozoon bieneusi genotypes SHR1 and PL2 in farmed masked palm civet (Paguma larvata) in China. International Journal for Parasitology: Parasites and Wildlife, 2021, 16, 99-102.	1.5	1
7	Molecular Characterization of Blastocystis sp. in Camelus bactrianus in Northwestern China. Animals, 2021, 11, 3016.	2.3	8
8	Genetic diversity of Blastocystis in kindergarten children in southern Xinjiang, China. Parasites and Vectors, 2020, 13, 15.	2.5	21
9	Molecular characterization of Cryptosporidium spp. in minks (Neovison vison), blue foxes (Vulpes) Tj ETQq1 1 0.784314 rgBT /Overlock Parasitology Research, 2020, 119, 3923-3927.	1.6	5
10	Genetic Diversity of Cryptosporidium in Bactrian Camels (Camelus bactrianus) in Xinjiang, Northwestern China. Pathogens, 2020, 9, 946.	2.8	11
11	Longitudinal identification of Enterocytozoon bieneusi in dairy calves on a farm in Southern Xinjiang, China. Comparative Immunology, Microbiology and Infectious Diseases, 2020, 73, 101550.	1.6	9
12	Molecular characterization and novel genotypes of Enterocytozoon bieneusi in pet snakes in Beijing, China. International Journal for Parasitology: Parasites and Wildlife, 2020, 12, 172-175.	1.5	5
13	Molecular identification of Cryptosporidium spp. in pet snakes in Beijing, China. Parasitology Research, 2020, 119, 3119-3123.	1.6	3
14	Genetic Diversity of Cryptosporidium parvum in Neonatal Dairy Calves in Xinjiang, China. Pathogens, 2020, 9, 692.	2.8	11
15	Unusual dominant genotype NIA1 of Enterocytozoon bieneusi in children in Southern Xinjiang, China. PLoS Neglected Tropical Diseases, 2020, 14, e0008293.	3.0	14
16	Molecular detection of Enterocytozoon bieneusi in farm-raised pigs in Hainan Province, China: infection rates, genotype distributions, and zoonotic potential. Parasite, 2020, 27, 12.	2.0	20
17	First Report of Blastocystis Infection in Pigs from Large Farms in Xinjiang, China. Journal of Eukaryotic Microbiology, 2020, 67, 642-647.	1.7	7
18	Molecular identification of Cryptosporidium spp. in alpacas (Vicugna pacos) in China. International Journal for Parasitology: Parasites and Wildlife, 2020, 12, 181-184.	1.5	6

#	Article	IF	Citations
19	Enterocytozoon bieneusi in donkeys from Xinjiang, China: prevalence, molecular characterization and the assessment of zoonotic risk. BMC Veterinary Research, 2020, 16, 196.	1.9	3
20	Molecular identification and genotyping of Enterocytozoon bieneusi in experimental rats in China. Experimental Parasitology, 2020, 210, 107850.	1.2	21
21	Unusual dominant genotype NIA1 of Enterocytozoon bieneusi in children in Southern Xinjiang, China. , 2020, 14, e0008293.		0
22	Unusual dominant genotype NIA1 of Enterocytozoon bieneusi in children in Southern Xinjiang, China. , 2020, 14, e0008293.		0
23	Unusual dominant genotype NIA1 of Enterocytozoon bieneusi in children in Southern Xinjiang, China. , 2020, 14, e0008293.		0
24	Unusual dominant genotype NIA1 of Enterocytozoon bieneusi in children in Southern Xinjiang, China. , 2020, 14, e0008293.		0
25	Unusual dominant genotype NIA1 of Enterocytozoon bieneusi in children in Southern Xinjiang, China. , 2020, 14, e0008293.		0
26	Unusual dominant genotype NIA1 of Enterocytozoon bieneusi in children in Southern Xinjiang, China. , 2020, 14, e0008293.		0
27	Prevalence and Population Genetics Analysis of Enterocytozoon bieneusi in Dairy Cattle in China. Frontiers in Microbiology, 2019, 10, 1399.	3.5	26
28	Genotyping and Zoonotic Potential of Enterocytozoon bieneusi in Pigs in Xinjiang, China. Frontiers in Microbiology, 2019, 10, 2401.	3.5	18
29	Occurrence of a Cryptosporidium xiaoiâ€“like genotype in peafowl (Pavo cristatus) in China. Parasitology Research, 2019, 118, 3555-3559.	1.6	6
30	Molecular characterization of three intestinal protozoans in hospitalized children with different disease backgrounds in Zhengzhou, central China. Parasites and Vectors, 2019, 12, 543.	2.5	32
31	Molecular characterization of Cryptosporidium and Enterocytozoon bieneusi in PÃ¨re David's deer (Elaphurus davidianus) from Shishou, China. International Journal for Parasitology: Parasites and Wildlife, 2019, 10, 184-187.	1.5	18
32	Molecular detection of Enterocytozoon bieneusi in alpacas (Vicugna pacos) in Xinjiang, China. Parasite, 2019, 26, 31.	2.0	7
33	Dominance of zoonotic genotype D of Enterocytozoon bieneusi in bamboo rats (Rhizomys sinensis). Infection, Genetics and Evolution, 2019, 73, 113-118.	2.3	23
34	Distribution and molecular characterization of Cryptosporidium spp., Giardia duodenalis, and Enterocytozoon bieneusi amongst grazing adult sheep in Xinjiang, China. Parasitology International, 2019, 71, 80-86.	1.3	23
35	The Potential Role of Synanthropic Rodents and Flies in the Transmission of Enterocytozoon bieneusi on a Dairy Cattle farm in China. Journal of Eukaryotic Microbiology, 2019, 66, 435-441.	1.7	30
36	Host specificity of Enterocytozoon bieneusi genotypes in Bactrian camels (Camelus bactrianus) in China. Parasites and Vectors, 2018, 11, 219.	2.5	21

#	Article	IF	Citations
37	Revisiting the infectivity and pathogenicity of Cryptosporidium avium provides new information on parasitic sites within the host. Parasites and Vectors, 2018, 11, 514.	2.5	13
38	An investigation of parasitic infections and review of molecular characterization of the intestinal protozoa in nonhuman primates in China from 2009 to 2015. International Journal for Parasitology: Parasites and Wildlife, 2017, 6, 8-15.	1.5	32
39	Prevalence, molecular epidemiology, and zoonotic potential of Entamoeba spp. in nonhuman primates in China. Infection, Genetics and Evolution, 2017, 54, 216-220.	2.3	15
40	Dominance of Enterocytozoon bieneusi genotype J in dairy calves in Xinjiang, Northwest China. Parasitology International, 2017, 66, 960-963.	1.3	31
41	Enterocytozoon bieneusi Genotypes in Grazing Horses in China and their Zoonotic Transmission Potential. Journal of Eukaryotic Microbiology, 2016, 63, 591-597.	1.7	47
42	Prevalence and genotyping of Giardia duodenalis isolated from sheep in Henan Province, central China. Infection, Genetics and Evolution, 2016, 39, 330-335.	2.3	31
43	Prevalence and multilocus genotyping of Cryptosporidium andersoni in dairy cattle and He cattle in Xinjiang, China. Infection, Genetics and Evolution, 2016, 44, 313-317.	2.3	31
44	Prevalence and multilocus genotyping of Giardia duodenalis in dairy calves in Xinjiang, Northwestern China. Parasites and Vectors, 2016, 9, 546.	2.5	29
45	Common occurrence of Cryptosporidium hominis in horses and donkeys. Infection, Genetics and Evolution, 2016, 43, 261-266.	2.3	37
46	Occurrence, molecular characterization and predominant genotypes of Enterocytozoon bieneusi in dairy cattle in Henan and Ningxia, China. Parasites and Vectors, 2016, 9, 142.	2.5	59
47	Infection rate and genetic diversity of Giardia duodenalis in pet and stray dogs in Henan Province, China. Parasitology International, 2016, 65, 159-162.	1.3	21
48	Prevalence of Zoonotic Giardia duodenalis Assemblage B and First Identification of Assemblage E in Rabbit Fecal Samples Isolates from Central China. Journal of Eukaryotic Microbiology, 2015, 62, 810-814.	1.7	26
49	Molecular Characterization of Cryptosporidium spp., Giardia duodenalis, and Enterocytozoon bieneusi in Captive Wildlife at Zhengzhou Zoo, China. Journal of Eukaryotic Microbiology, 2015, 62, 833-839.	1.7	74
50	Molecular identification of Cryptosporidium spp. and Giardia duodenalis in grazing horses from Xinjiang, China. Veterinary Parasitology, 2015, 209, 169-172.	1.8	31
51	Occurrence and molecular identification of Cryptosporidium spp. in dairy calves in Xinjiang, Northwestern China. Veterinary Parasitology, 2015, 212, 404-407.	1.8	39
52	Zoonotic Cryptosporidium spp. and Enterocytozoon bieneusi in pet chinchillas (Chinchilla lanigera) in China. Parasitology International, 2015, 64, 339-341.	1.3	56
53	Multilocus genotyping of potentially zoonotic Giardia duodenalis in pet chinchillas (Chinchilla) Tj ETQq1 1 0.784314 rgBT /Overlock 1	1.8	24
54	Molecular characterization of Cryptosporidium spp. and Giardia duodenalis from yaks in the central western region of China. BMC Microbiology, 2015, 15, 108.	3.3	43

#	Article	IF	Citations
55	<i>Cryptosporidium</i> spp., <i>Giardia intestinalis</i>, and <i>Enterocytozoon bieneusi</i> in Captive Non-Human Primates in Qinling Mountains. Korean Journal of Parasitology, 2015, 53, 395-402.	1.3	53
56	Prevalence and molecular characterization of Cryptosporidium spp. and Giardia duodenalisin dairy cattle in Ningxia, northwestern China. BMC Veterinary Research, 2014, 10, 292.	1.9	88
57	Genetic Polymorphism and Zoonotic Potential of Enterocytozoon bieneusi from Nonhuman Primates in China. Applied and Environmental Microbiology, 2014, 80, 1893-1898.	3.1	128
58	Occurrence and molecular characterization of Cryptosporidium in dogs in Henan Province, China. BMC Veterinary Research, 2014, 10, 26.	1.9	27
59	Natural infection of Cryptosporidium muris in ostriches (Struthio camelus). Veterinary Parasitology, 2014, 205, 518-522.	1.8	22
60	Multilocus typing of Cryptosporidium spp. and Giardia duodenalis from non-human primates in China. International Journal for Parasitology, 2014, 44, 1039-1047.	3.1	51
61	Multilocus sequence typing of Enterocytozoon bieneusi in nonhuman primates in China. Veterinary Parasitology, 2014, 200, 13-23.	1.8	42
62	Cryptosporidium parvum IId family: clonal population and dispersal from Western Asia to other geographical regions. Scientific Reports, 2014, 4, 4208.	3.3	58
63	Cryptosporidium spp. in pet birds: Genetic diversity and potential public health significance. Experimental Parasitology, 2011, 128, 336-340.	1.2	82
64	Characteristics of Cryptosporidium Transmission in Preweaned Dairy Cattle in Henan, China. Journal of Clinical Microbiology, 2011, 49, 1077-1082.	3.9	102
65	Cryptosporidium spp. in Wild, Laboratory, and Pet Rodents in China: Prevalence and Molecular Characterization. Applied and Environmental Microbiology, 2009, 75, 7692-7699.	3.1	110