

Meng Qi

List of Publications by Year in descending order

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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	Genetic Polymorphism and Zoonotic Potential of <i>Enterocytozoon bienersi</i> from Nonhuman Primates in China. <i>Applied and Environmental Microbiology</i> , 2014, 80, 1893-1898.	3.1	128
2	<i>Cryptosporidium</i> spp. in Wild, Laboratory, and Pet Rodents in China: Prevalence and Molecular Characterization. <i>Applied and Environmental Microbiology</i> , 2009, 75, 7692-7699.	3.1	110
3	Characteristics of <i>Cryptosporidium</i> Transmission in Prewaned Dairy Cattle in Henan, China. <i>Journal of Clinical Microbiology</i> , 2011, 49, 1077-1082.	3.9	102
4	Prevalence and molecular characterization of <i>Cryptosporidium</i> spp. and <i>Giardia duodenalis</i> in dairy cattle in Ningxia, northwestern China. <i>BMC Veterinary Research</i> , 2014, 10, 292.	1.9	88
5	<i>Cryptosporidium</i> spp. in pet birds: Genetic diversity and potential public health significance. <i>Experimental Parasitology</i> , 2011, 128, 336-340.	1.2	82
6	Molecular Characterization of <i>Cryptosporidium</i> spp., <i>Giardia duodenalis</i> , and <i>Enterocytozoon bienersi</i> in Captive Wildlife at Zhengzhou Zoo, China. <i>Journal of Eukaryotic Microbiology</i> , 2015, 62, 833-839.	1.7	74
7	Occurrence, molecular characterization and predominant genotypes of <i>Enterocytozoon bienersi</i> in dairy cattle in Henan and Ningxia, China. <i>Parasites and Vectors</i> , 2016, 9, 142.	2.5	59
8	<i>Cryptosporidium parvum</i> IId family: clonal population and dispersal from Western Asia to other geographical regions. <i>Scientific Reports</i> , 2014, 4, 4208.	3.3	58
9	Zoonotic <i>Cryptosporidium</i> spp. and <i>Enterocytozoon bienersi</i> in pet chinchillas (<i>Chinchilla lanigera</i>) in China. <i>Parasitology International</i> , 2015, 64, 339-341.	1.3	56
10	<i>Cryptosporidium</i> spp., <i>Giardia intestinalis</i> , and <i>Enterocytozoon bienersi</i> in Captive Non-Human Primates in Qinling Mountains. <i>Korean Journal of Parasitology</i> , 2015, 53, 395-402.	1.3	53
11	Multilocus typing of <i>Cryptosporidium</i> spp. and <i>Giardia duodenalis</i> from non-human primates in China. <i>International Journal for Parasitology</i> , 2014, 44, 1039-1047.	3.1	51
12	<i>Enterocytozoon bienersi</i> Genotypes in Grazing Horses in China and their Zoonotic Transmission Potential. <i>Journal of Eukaryotic Microbiology</i> , 2016, 63, 591-597.	1.7	47
13	Molecular characterization of <i>Cryptosporidium</i> spp. and <i>Giardia duodenalis</i> from yaks in the central western region of China. <i>BMC Microbiology</i> , 2015, 15, 108.	3.3	43
14	Multilocus sequence typing of <i>Enterocytozoon bienersi</i> in nonhuman primates in China. <i>Veterinary Parasitology</i> , 2014, 200, 13-23.	1.8	42
15	Occurrence and molecular identification of <i>Cryptosporidium</i> spp. in dairy calves in Xinjiang, Northwestern China. <i>Veterinary Parasitology</i> , 2015, 212, 404-407.	1.8	39
16	Common occurrence of <i>Cryptosporidium hominis</i> in horses and donkeys. <i>Infection, Genetics and Evolution</i> , 2016, 43, 261-266.	2.3	37
17	An investigation of parasitic infections and review of molecular characterization of the intestinal protozoa in nonhuman primates in China from 2009 to 2015. <i>International Journal for Parasitology: Parasites and Wildlife</i> , 2017, 6, 8-15.	1.5	32
18	Molecular characterization of three intestinal protozoans in hospitalized children with different disease backgrounds in Zhengzhou, central China. <i>Parasites and Vectors</i> , 2019, 12, 543.	2.5	32

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19	Molecular identification of <i>Cryptosporidium</i> spp. and <i>Giardia duodenalis</i> in grazing horses from Xinjiang, China. <i>Veterinary Parasitology</i> , 2015, 209, 169-172.	1.8	31
20	Prevalence and genotyping of <i>Giardia duodenalis</i> isolated from sheep in Henan Province, central China. <i>Infection, Genetics and Evolution</i> , 2016, 39, 330-335.	2.3	31
21	Prevalence and multilocus genotyping of <i>Cryptosporidium andersoni</i> in dairy cattle and He cattle in Xinjiang, China. <i>Infection, Genetics and Evolution</i> , 2016, 44, 313-317.	2.3	31
22	Dominance of <i>Enterocytozoon bienersi</i> genotype J in dairy calves in Xinjiang, Northwest China. <i>Parasitology International</i> , 2017, 66, 960-963.	1.3	31
23	The Potential Role of Synanthropic Rodents and Flies in the Transmission of <i>Enterocytozoon bienersi</i> on a Dairy Cattle farm in China. <i>Journal of Eukaryotic Microbiology</i> , 2019, 66, 435-441.	1.7	30
24	Prevalence and multilocus genotyping of <i>Giardia duodenalis</i> in dairy calves in Xinjiang, Northwestern China. <i>Parasites and Vectors</i> , 2016, 9, 546.	2.5	29
25	Occurrence and molecular characterization of <i>Cryptosporidium</i> in dogs in Henan Province, China. <i>BMC Veterinary Research</i> , 2014, 10, 26.	1.9	27
26	Prevalence of Zoonotic <i>Giardia duodenalis</i> Assemblage B and First Identification of Assemblage E in Rabbit Fecal Samples Isolates from Central China. <i>Journal of Eukaryotic Microbiology</i> , 2015, 62, 810-814.	1.7	26
27	Prevalence and Population Genetics Analysis of <i>Enterocytozoon bienersi</i> in Dairy Cattle in China. <i>Frontiers in Microbiology</i> , 2019, 10, 1399.	3.5	26
28	Multilocus genotyping of potentially zoonotic <i>Giardia duodenalis</i> in pet chinchillas (<i>Chinchilla</i>) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 382	1.8	24
29	Dominance of zoonotic genotype D of <i>Enterocytozoon bienersi</i> in bamboo rats (<i>Rhizomys sinensis</i>). <i>Infection, Genetics and Evolution</i> , 2019, 73, 113-118.	2.3	23
30	Distribution and molecular characterization of <i>Cryptosporidium</i> spp., <i>Giardia duodenalis</i> , and <i>Enterocytozoon bienersi</i> amongst grazing adult sheep in Xinjiang, China. <i>Parasitology International</i> , 2019, 71, 80-86.	1.3	23
31	Natural infection of <i>Cryptosporidium muris</i> in ostriches (<i>Struthio camelus</i>). <i>Veterinary Parasitology</i> , 2014, 205, 518-522.	1.8	22
32	Infection rate and genetic diversity of <i>Giardia duodenalis</i> in pet and stray dogs in Henan Province, China. <i>Parasitology International</i> , 2016, 65, 159-162.	1.3	21
33	Host specificity of <i>Enterocytozoon bienersi</i> genotypes in Bactrian camels (<i>Camelus bactrianus</i>) in China. <i>Parasites and Vectors</i> , 2018, 11, 219.	2.5	21
34	Genetic diversity of <i>Blastocystis</i> in kindergarten children in southern Xinjiang, China. <i>Parasites and Vectors</i> , 2020, 13, 15.	2.5	21
35	Molecular identification and genotyping of <i>Enterocytozoon bienersi</i> in experimental rats in China. <i>Experimental Parasitology</i> , 2020, 210, 107850.	1.2	21
36	Molecular detection of <i>Enterocytozoon bienersi</i> in farm-raised pigs in Hainan Province, China: infection rates, genotype distributions, and zoonotic potential. <i>Parasite</i> , 2020, 27, 12.	2.0	20

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37	Genotyping and Zoonotic Potential of <i>Enterocytozoon bienersi</i> in Pigs in Xinjiang, China. <i>Frontiers in Microbiology</i> , 2019, 10, 2401.	3.5	18
38	Molecular characterization of <i>Cryptosporidium</i> and <i>Enterocytozoon bienersi</i> in Père David's deer (<i>Elaphurus davidianus</i>) from Shishou, China. <i>International Journal for Parasitology: Parasites and Wildlife</i> , 2019, 10, 184-187.	1.5	18
39	Prevalence, molecular epidemiology, and zoonotic potential of <i>Entamoeba</i> spp. in nonhuman primates in China. <i>Infection, Genetics and Evolution</i> , 2017, 54, 216-220.	2.3	15
40	Unusual dominant genotype NIA1 of <i>Enterocytozoon bienersi</i> in children in Southern Xinjiang, China. <i>PLoS Neglected Tropical Diseases</i> , 2020, 14, e0008293.	3.0	14
41	Revisiting the infectivity and pathogenicity of <i>Cryptosporidium avium</i> provides new information on parasitic sites within the host. <i>Parasites and Vectors</i> , 2018, 11, 514.	2.5	13
42	Genetic Diversity of <i>Cryptosporidium</i> in Bactrian Camels (<i>Camelus bactrianus</i>) in Xinjiang, Northwestern China. <i>Pathogens</i> , 2020, 9, 946.	2.8	11
43	Genetic Diversity of <i>Cryptosporidium parvum</i> in Neonatal Dairy Calves in Xinjiang, China. <i>Pathogens</i> , 2020, 9, 692.	2.8	11
44	Longitudinal identification of <i>Enterocytozoon bienersi</i> in dairy calves on a farm in Southern Xinjiang, China. <i>Comparative Immunology, Microbiology and Infectious Diseases</i> , 2020, 73, 101550.	1.6	9
45	Molecular Characterization of <i>Blastocystis</i> sp. in <i>Camelus bactrianus</i> in Northwestern China. <i>Animals</i> , 2021, 11, 3016.	2.3	8
46	Molecular detection of <i>Enterocytozoon bienersi</i> in alpacas (<i>Vicugna pacos</i>) in Xinjiang, China. <i>Parasite</i> , 2019, 26, 31.	2.0	7
47	First Report of <i>Blastocystis</i> Infection in Pigs from Large Farms in Xinjiang, China. <i>Journal of Eukaryotic Microbiology</i> , 2020, 67, 642-647.	1.7	7
48	Occurrence of a <i>Cryptosporidium</i> xiaoi-like genotype in peafowl (<i>Pavo cristatus</i>) in China. <i>Parasitology Research</i> , 2019, 118, 3555-3559.	1.6	6
49	Molecular identification of <i>Cryptosporidium</i> spp. in alpacas (<i>Vicugna pacos</i>) in China. <i>International Journal for Parasitology: Parasites and Wildlife</i> , 2020, 12, 181-184.	1.5	6
50	Molecular characterization of <i>Cryptosporidium</i> spp. in minks (<i>Neovison vison</i>), blue foxes (<i>Vulpes</i>) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 <i>Parasitology Research</i> , 2020, 119, 3923-3927.	1.6	5
51	Molecular characterization and novel genotypes of <i>Enterocytozoon bienersi</i> in pet snakes in Beijing, China. <i>International Journal for Parasitology: Parasites and Wildlife</i> , 2020, 12, 172-175.	1.5	5
52	Molecular detection and genotypes of <i>Enterocytozoon bienersi</i> in farmed mink (<i>Neovison vison</i>), blue foxes (<i>Alopex lagopus</i>), and raccoon dogs (<i>Nyctereutes procyonoides</i>) in Xinjiang, China. <i>International Journal for Parasitology: Parasites and Wildlife</i> , 2021, 14, 211-215.	1.5	5
53	Molecular characterization of <i>Cryptosporidium</i> spp. and <i>Giardia duodenalis</i> in pet dogs in Xinjiang, China. <i>Parasitology Research</i> , 2022, 121, 1429-1435.	1.6	5
54	Seasonal monitoring of <i>Cryptosporidium</i> species and their genetic diversity in neonatal calves on two large-scale farms in Xinjiang, China. <i>Journal of Eukaryotic Microbiology</i> , 2022, 69, e12878.	1.7	5

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55	Molecular identification of <i>Cryptosporidium</i> spp. in pet snakes in Beijing, China. <i>Parasitology Research</i> , 2020, 119, 3119-3123.	1.6	3
56	<i>Enterocytozoon bieneusi</i> in donkeys from Xinjiang, China: prevalence, molecular characterization and the assessment of zoonotic risk. <i>BMC Veterinary Research</i> , 2020, 16, 196.	1.9	3
57	Genetic diversity of <i>Blastocystis</i> subtypes in the Alpine musk deer (<i>Moschus moschiferus</i>) in Xinjiang, China. <i>Parasitology Research</i> , 2020, 119, e12910.	1.7	3
58	Molecular detection and genetic characteristics of <i>Cryptosporidium</i> spp. in Chinese racehorses. <i>Equine Veterinary Journal</i> , 2023, 55, 474-480.	1.7	3
59	Common occurrence of <i>Enterocytozoon bieneusi</i> genotypes SHR1 and PL2 in farmed masked palm civet (<i>Paguma larvata</i>) in China. <i>International Journal for Parasitology: Parasites and Wildlife</i> , 2021, 16, 99-102.	1.5	1
60	Unusual dominant genotype NIA1 of <i>Enterocytozoon bieneusi</i> in children in Southern Xinjiang, China. <i>Parasitology Research</i> , 2020, 14, e0008293.		0
61	Unusual dominant genotype NIA1 of <i>Enterocytozoon bieneusi</i> in children in Southern Xinjiang, China. <i>Parasitology Research</i> , 2020, 14, e0008293.		0
62	Unusual dominant genotype NIA1 of <i>Enterocytozoon bieneusi</i> in children in Southern Xinjiang, China. <i>Parasitology Research</i> , 2020, 14, e0008293.		0
63	Unusual dominant genotype NIA1 of <i>Enterocytozoon bieneusi</i> in children in Southern Xinjiang, China. <i>Parasitology Research</i> , 2020, 14, e0008293.		0
64	Unusual dominant genotype NIA1 of <i>Enterocytozoon bieneusi</i> in children in Southern Xinjiang, China. <i>Parasitology Research</i> , 2020, 14, e0008293.		0
65	Unusual dominant genotype NIA1 of <i>Enterocytozoon bieneusi</i> in children in Southern Xinjiang, China. <i>Parasitology Research</i> , 2020, 14, e0008293.		0