Lihua Xiao

List of Publications by Year in descending order

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464 papers	33,387 citations	2975 93 h-index	6471 157 g-index
482 all docs	482 docs citations	482 times ranked	9727 citing authors

#	Article	IF	Citations
1	Cryptosporidiosis outbreak caused by Cryptosporidium parvum subtype IIdA20G1 in neonatal calves. Transboundary and Emerging Diseases, 2022, 69, 278-285.	3.0	11
2	Enterocytozoon bieneusi. Trends in Parasitology, 2022, 38, 95-96.	3.3	16
3	Comparative Characterization of CpCDPK1 and CpCDPK9, Two Potential Drug Targets Against Cryptosporidiosis. Microorganisms, 2022, 10, 333.	3.6	5
4	Emergence of zoonotic Cryptosporidium parvum in China. Trends in Parasitology, 2022, 38, 335-343.	3.3	24
5	A productive immunocompetent mouse model of cryptosporidiosis with long oocyst shedding duration for immunological studies. Journal of Infection, 2022, 84, 710-721.	3.3	7
6	High zoonotic potential of Cryptosporidium spp., Giardia duodenalis, and Enterocytozoon bieneusi in wild nonhuman primates from Yunnan Province, China. Parasites and Vectors, 2022, 15, 85.	2.5	5
7	Age and episodeâ€associated occurrence of Cryptosporidium species and subtypes in a birthâ€cohort of dairy calves. Transboundary and Emerging Diseases, 2022, 69, .	3.0	3
8	Diarrhoea outbreak caused by coinfections of Cryptosporidium parvum subtype IIdA20G1 and rotavirus in preâ€weaned dairy calves. Transboundary and Emerging Diseases, 2022, 69, .	3.0	8
9	Characterization of Calcium-Dependent Protein Kinase 2A, a Potential Drug Target Against Cryptosporidiosis. Frontiers in Microbiology, 2022, 13, 883674.	3.5	2
10	Prevalence and genetic characterization of Enterocytozoon bieneusi in children in Northeast Egypt. Parasitology Research, 2022, 121, 2087-2092.	1.6	3
11	Characterization of Dense Granule Metalloproteinase INS-16 in Cryptosporidium parvum. International Journal of Molecular Sciences, 2022, 23, 7617.	4.1	3
12	Sympatric Recombination in Zoonotic Cryptosporidium Leads to Emergence of Populations with Modified Host Preference. Molecular Biology and Evolution, 2022, 39, .	8.9	10
13	Cryptosporidium ratti n. sp. (Apicomplexa: Cryptosporidiidae) and genetic diversity ofCryptosporidiumspp. in brown rats (Rattus norvegicus) in the Czech Republic. Parasitology, 2021, 148, 84-97.	1.5	24
14	Characterizations of Enterocytozoon bieneusi at new genetic loci reveal a lack of strict host specificity among common genotypes and the existence of a canine-adapted Enterocytozoon species. International Journal for Parasitology, 2021, 51, 215-223.	3.1	9
15	Development of a Subtyping Tool for Zoonotic Pathogen Cryptosporidium canis. Journal of Clinical Microbiology, 2021, 59, .	3.9	20
16	Cryptosporidial Infection Suppresses Intestinal Epithelial Cell MAPK Signaling Impairing Host Anti-Parasitic Defense. Microorganisms, 2021, 9, 151.	3.6	11
17	Subtype Characterization and Zoonotic Potential of Cryptosporidium felis in Cats in Guangdong and Shanghai, China. Pathogens, 2021, 10, 89.	2.8	8
18	Molecular Epidemiology of Human Cryptosporidiosis in Low- and Middle-Income Countries. Clinical Microbiology Reviews, 2021, 34, .	13.6	56

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19	Small ruminants and zoonotic cryptosporidiosis. Parasitology Research, 2021, 120, 4189-4198.	1.6	28
20	Genetic Manipulation of Cryptosporidium. , 2021, , 489-498.		0
21	Zoonotic parasites in farmed exotic animals in China: Implications to public health. International Journal for Parasitology: Parasites and Wildlife, 2021, 14, 241-247.	1.5	9
22	Codon usage analysis of zoonotic coronaviruses reveals lower adaptation to humans by SARS-CoV-2. Infection, Genetics and Evolution, 2021, 89, 104736.	2.3	13
23	Comparative Study of Two Insulinlike Proteases in Cryptosporidium parvum. Microorganisms, 2021, 9, 861.	3.6	3
24	Insulinase-like Protease 1 Contributes to Macrogamont Formation in Cryptosporidium parvum. MBio, 2021, 12, .	4.1	10
25	Prevalence and molecular characterization of novel species of the Diplomonad genus Octomitus (Diplomonadida: Giardiinae) from wildlife in a New York watershed. International Journal for Parasitology: Parasites and Wildlife, 2021, 14, 267-272.	1.5	0
26	Preliminary Characterization of Two Small Insulinase-Like Proteases in Cryptosporidium parvum. Frontiers in Microbiology, 2021, 12, 651512.	3.5	3
27	Genetic Characterization of Cryptosporidium cuniculus from Rabbits in Egypt. Pathogens, 2021, 10, 775.	2.8	9
28	Ecological and public health significance of Enterocytozoon bieneusi. One Health, 2021, 12, 100209.	3.4	46
29	Subtyping Cryptosporidium xiaoi, a Common Pathogen in Sheep and Goats. Pathogens, 2021, 10, 800.	2.8	11
30	Zoonotic parasites: the One Health challenge. Parasitology Research, 2021, 120, 4073-4074.	1.6	4
31	Molecular detection of Cryptosporidium spp., Giardia duodenalis, and Enterocytozoon bieneusi in school children at the Thai-Myanmar border. Parasitology Research, 2021, 120, 2887-2895.	1.6	4
32	Genus-level evolutionary relationships of FAR proteins reflect the diversity of lifestyles of free-living and parasitic nematodes. BMC Biology, 2021, 19, 178.	3.8	4
33	Molecular analysis of cryptosporidiosis cases in Western Australia in 2019 and 2020 supports the occurrence of two swimming pool associated outbreaks and reveals the emergence of a rare C. hominis IbA12G3 subtype. Infection, Genetics and Evolution, 2021, 92, 104859.	2.3	12
34	Editorial: Recent Advances in the Controversial Human Pathogens Pneumocystis, Microsporidia and Blastocystis. Frontiers in Microbiology, 2021, 12, 701879.	3.5	0
35	Advances in molecular epidemiology of cryptosporidiosis in dogs and cats. International Journal for Parasitology, 2021, 51, 787-795.	3.1	13
36	Genetic characterizations of Cryptosporidium spp. from pet rodents indicate high zoonotic potential of pathogens from chinchillas. One Health, 2021, 13, 100269.	3.4	5

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37	Molecular characterization of the waterborne pathogens Cryptosporidium spp., Giardia duodenalis, Enterocytozoon bieneusi, Cyclospora cayetanensis and Eimeria spp. in wastewater and sewage in Guangzhou, China. Parasites and Vectors, 2021, 14, 66.	2.5	17
38	Development and Application of a gp60-Based Subtyping Tool for Cryptosporidium bovis. Microorganisms, 2021, 9, 2067.	3.6	8
39	Taxonomy and molecular epidemiology of Cryptosporidium and Giardia â€“ a 50Âyear perspective (1971â€“2021). International Journal for Parasitology, 2021, 51, 1099-1119.	3.1	128
40	Zoonotic giardiasis: an update. Parasitology Research, 2021, 120, 4199-4218.	1.6	71
41	An Update on Zoonotic Cryptosporidium Species and Genotypes in Humans. Animals, 2021, 11, 3307.	2.3	84
42	Association of Common Zoonotic Pathogens With Concentrated Animal Feeding Operations. Frontiers in Microbiology, 2021, 12, 810142.	3.5	6
43	Cryptosporidium felis differs from other Cryptosporidium spp. in codon usage. Microbial Genomics, 2021, 7, .	2.0	3
44	Detection of SARS-CoV-2 RNA with a Simple Concentration Method in Wastewater in Turkey: A Pilot Study in Ã‡orum. Flora: the Journal of Infectious Diseses and Clinical Microbiology = Infeksiyon HastalÄ±klarÄ± Ve Klinik Mikrobiyoloji Dergisi, 2021, 26, 620-627.	0.1	1
45	Cryptosporidiosis. , 2020, , 712-718.		4
46	Cyclospora cayetanensis infection in humans: biological characteristics, clinical features, epidemiology, detection method and treatment. Parasitology, 2020, 147, 160-170.	1.5	38
47	Cryptosporidium parvum as a risk factor of diarrhea occurrence in neonatal alpacas in Peru. Parasitology Research, 2020, 119, 243-248.	1.6	5
48	Population genetic analysis suggests genetic recombination is responsible for increased zoonotic potential of Enterocytozoon bieneusi from ruminants in China. One Health, 2020, 11, 100184.	3.4	7
49	Subtype distribution of zoonotic pathogen Cryptosporidium felis in humans and animals in several countries. Emerging Microbes and Infections, 2020, 9, 2446-2454.	6.5	19
50	Diagnosis and molecular typing of Enterocytozoon bieneusi: the significant role of domestic animals in transmission of human microsporidiosis. Research in Veterinary Science, 2020, 133, 251-261.	1.9	29
51	Occurrence and molecular characterization of Giardia duodenalis in lambs in Djelfa, the central steppe of Algeria. Parasitology Research, 2020, 119, 2965-2973.	1.6	4
52	Cryptosporidium Species and C. parvum Subtypes in Farmed Bamboo Rats. Pathogens, 2020, 9, 1018.	2.8	8
53	Contribution of hospitals to the occurrence of enteric protists in urban wastewater. Parasitology Research, 2020, 119, 3033-3040.	1.6	12
54	Molecular characterization and zoonotic potential of Enterocytozoon bieneusi, Giardia duodenalis and Cryptosporidium sp. in farmed masked palm civets (Paguma larvata) in southern China. Parasites and Vectors, 2020, 13, 403.	2.5	19

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55	Subtyping Cryptosporidium ryanae: A Common Pathogen in Bovine Animals. Microorganisms, 2020, 8, 1107.	3.6	18
56	Population structure and geographical segregation of Cryptosporidium parvum IId subtypes in cattle in China. Parasites and Vectors, 2020, 13, 425.	2.5	15
57	Characterization of Calcium-Dependent Protein Kinases 3, a Protein Involved in Growth of Cryptosporidium parvum. Frontiers in Microbiology, 2020, 11, 907.	3.5	8
58	Expression and Functional Studies of INS-5, an Insulinase-Like Protein in Cryptosporidium parvum. Frontiers in Microbiology, 2020, 11, 719.	3.5	7
59	Update on Cryptosporidium spp.: highlights from the Seventh International Giardia and Cryptosporidium Conference. Parasite, 2020, 27, 14.	2.0	40
60	Common occurrence of divergent Cryptosporidium species and Cryptosporidium parvum subtypes in farmed bamboo rats (Rhizomys sinensis). Parasites and Vectors, 2020, 13, 149.	2.5	19
61	Isolation of SARS-CoV-2-related coronavirus from Malayan pangolins. Nature, 2020, 583, 286-289.	27.8	599
62	Zoonotic potential of Enterocytozoon bieneusi and Giardia duodenalis in horses and donkeys in northern China. Parasitology Research, 2020, 119, 1101-1108.	1.6	20
63	Cryptosporidiosis in HIV-positive patients and related risk factors: A systematic review and meta-analysis. Parasite, 2020, 27, 27.	2.0	33
64	Multilocus sequence typing of Enterocytozoon bieneusi in crab-eating macaques (Macaca) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 382 Td	2.5	2
65	Characterization of Three Calcium-Dependent Protein Kinases of Cryptosporidium parvum. Frontiers in Microbiology, 2020, 11, 622203.	3.5	6
66	Cryptosporidium Genotyping for Epidemiology Tracking. Methods in Molecular Biology, 2020, 2052, 103-116.	0.9	8
67	Comparative genomic analysis of three intestinal species reveals reductions in secreted pathogenesis determinants in bovine-specific and non-pathogenic Cryptosporidium species. Microbial Genomics, 2020, 6, .	2.0	13
68	Infection patterns, clinical significance, and genetic characteristics of Enterocytozoon bieneusi and Giardia duodenalis in dairy cattle in Jiangsu, China. Parasitology Research, 2019, 118, 3053-3060.	1.6	30
69	Cryptosporidium parvum and Cryptosporidium hominis subtypes in crab-eating macaques. Parasites and Vectors, 2019, 12, 350.	2.5	26
70	Different distribution of Cryptosporidium species between horses and donkeys. Infection, Genetics and Evolution, 2019, 75, 103954.	2.3	21
71	Characterization of INS-15, A Metalloprotease Potentially Involved in the Invasion of Cryptosporidium parvum. Microorganisms, 2019, 7, 452.	3.6	16
72	Trichostatin A, a Histone Deacetylase Inhibitor, Alleviates Eosinophilic Meningitis Induced by Angiostrongylus cantonensis Infection in Mice. Frontiers in Microbiology, 2019, 10, 2280.	3.5	7

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73	Divergent Copies of a Cryptosporidium parvum-Specific Subtelomeric Gene. Microorganisms, 2019, 7, 366.	3.6	4
74	Comparative genomics: how has it advanced our knowledge of cryptosporidiosis epidemiology?. Parasitology Research, 2019, 118, 3195-3204.	1.6	17
75	Epidemiological distribution of genotypes of Giardia duodenalis in humans in Spain. Parasites and Vectors, 2019, 12, 432.	2.5	29
76	Cryptosporidium infections in terrestrial ungulates with focus on livestock: a systematic review and meta-analysis. Parasites and Vectors, 2019, 12, 453.	2.5	59
77	Potential impacts of host specificity on zoonotic or interspecies transmission of Enterocytozoon bieneusi. Infection, Genetics and Evolution, 2019, 75, 104033.	2.3	47
78	Prevalence and genotypic identification of Cryptosporidium spp., Giardia duodenalis and Enterocytozoon bieneusi in pre-weaned dairy calves in Guangdong, China. Parasites and Vectors, 2019, 12, 41.	2.5	55
79	Genotypes and public health potential of Enterocytozoon bieneusi and Giardia duodenalis in crab-eating macaques. Parasites and Vectors, 2019, 12, 254.	2.5	22
80	Comparative analysis reveals conservation in genome organization among intestinal Cryptosporidium species and sequence divergence in potential secreted pathogenesis determinants among major human-infecting species. BMC Genomics, 2019, 20, 406.	2.8	37
81	Isolation, genotyping and subtyping of single Cryptosporidium oocysts from calves with special reference to zoonotic significance. Veterinary Parasitology, 2019, 271, 80-86.	1.8	12
82	Differential Expression of Three Cryptosporidium Species-Specific MEDLE Proteins. Frontiers in Microbiology, 2019, 10, 1177.	3.5	11
83	Retrospective analysis of Cryptosporidium species in Western Australian human populations (2015â€“2018), and emergence of the C. hominis IfA12G1R5 subtype. Infection, Genetics and Evolution, 2019, 73, 306-313.	2.3	28
84	Outbreak of cryptosporidiosis due to Cryptosporidium parvum subtype IIdA19G1 in neonatal calves on a dairy farm in China. International Journal for Parasitology, 2019, 49, 569-577.	3.1	39
85	Multilocus Sequence Typing and Population Genetic Analysis of Enterocytozoon bieneusi: Host Specificity and Its Impacts on Public Health. Frontiers in Genetics, 2019, 10, 307.	2.3	41
86	Characterization of a Species-Specific Insulinase-Like Protease in Cryptosporidium parvum. Frontiers in Microbiology, 2019, 10, 354.	3.5	18
87	Genetic characterization of Cryptosporidium spp. and Giardia duodenalis in dogs and cats in Guangdong, China. Parasites and Vectors, 2019, 12, 571.	2.5	28
88	Host-adapted Cryptosporidium and Enterocytozoon bieneusi genotypes in straw-colored fruit bats in Nigeria. International Journal for Parasitology: Parasites and Wildlife, 2019, 8, 19-24.	1.5	17
89	Giardia: an under-reported foodborne parasite. International Journal for Parasitology, 2019, 49, 1-11.	3.1	131
90	Divergent Cryptosporidium parvum subtype and Enterocytozoon bieneusi genotypes in dromedary camels in Algeria. Parasitology Research, 2018, 117, 905-910.	1.6	21

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91	Population genetic characterization of Cyclospora cayetanensis from discrete geographical regions. Experimental Parasitology, 2018, 184, 121-127.	1.2	11
92	Enterocytozoon bieneusi genotypes in Tibetan sheep and yaks. Parasitology Research, 2018, 117, 721-727.	1.6	37
93	Epidemiological observations on cryptosporidiosis and molecular characterization of Cryptosporidium spp. in sheep and goats in Kuwait. Parasitology Research, 2018, 117, 1631-1636.	1.6	26
94	Genotypes and subtypes of Cryptosporidium spp. in diarrheic lambs and goat kids in northern Greece. Parasitology International, 2018, 67, 472-475.	1.3	25
95	Clinical Manifestations of Cryptosporidiosis and Identification of a New Cryptosporidium Subtype in Patients From Sonora, Mexico. Pediatric Infectious Disease Journal, 2018, 37, e136-e138.	2.0	15
96	Cryptosporidiuminfecting wild cricetid rodents from the subfamilies Arvicolinae and Neotominae. Parasitology, 2018, 145, 326-334.	1.5	14
97	Foodborne cryptosporidiosis. International Journal for Parasitology, 2018, 48, 1-12.	3.1	143
98	Water quality, availability, and acute gastroenteritis on the Navajo Nation â€“ a pilot case-control study. Journal of Water and Health, 2018, 16, 1018-1028.	2.6	4
99	Zoonotic Cryptosporidium species and subtypes in lambs and goat kids in Algeria. Parasites and Vectors, 2018, 11, 582.	2.5	30
100	Outbreaks Associated with Treated Recreational Water â€” United States, 2000â€“2014. Morbidity and Mortality Weekly Report, 2018, 67, 547-551.	15.1	51
101	Persistent Occurrence of Cryptosporidium hominis and Giardia duodenalis Subtypes in a Welfare Institute. Frontiers in Microbiology, 2018, 9, 2830.	3.5	13
102	Age patterns of Cryptosporidium species and Giardia duodenalis in dairy calves in Egypt. Parasitology International, 2018, 67, 736-741.	1.3	32
103	Outbreaks associated with treated recreational water - United States, 2000-2014. American Journal of Transplantation, 2018, 18, 1815-1819.	4.7	8
104	Molecular characterization of Cryptosporidium spp. and Giardia duodenalis in children in Egypt. Parasites and Vectors, 2018, 11, 403.	2.5	40
105	Clinical, environmental, and behavioral characteristics associated with Cryptosporidium infection among children with moderate-to-severe diarrhea in rural western Kenya, 2008â€“2012: The Global Enteric Multicenter Study (GEMS). PLoS Neglected Tropical Diseases, 2018, 12, e0006640.	3.0	25
106	Genetic diversity within dominant Enterocytozoon bieneusi genotypes in pre-weaned calves. Parasites and Vectors, 2018, 11, 170.	2.5	32
107	Widespread occurrence of Cryptosporidium infections in patients with HIV/AIDS: Epidemiology, clinical feature, diagnosis, and therapy. Acta Tropica, 2018, 187, 257-263.	2.0	76
108	Characterization of MEDLE-1, a protein in early development of Cryptosporidium parvum. Parasites and Vectors, 2018, 11, 312.	2.5	14

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109	Genetic Diversity and Population Structure of Cryptosporidium. Trends in Parasitology, 2018, 34, 997-1011.	3.3	365
110	Cryptosporidium and Cryptosporidiosis. , 2018, , 73-117.		8
111	A Randomized Controlled Trial to Assess the Impact of Ceramic Water Filters on Prevention of Diarrhea and Cryptosporidiosis in Infants and Young Childrenâ€"Western Kenya, 2013. American Journal of Tropical Medicine and Hygiene, 2018, 98, 1260-1268.	1.4	22
112	Cryptosporidium. , 2018, , 551-563.		0
113	Comparative genomic analysis of the IId subtype family of Cryptosporidium parvum. International Journal for Parasitology, 2017, 47, 281-290.	3.1	58
114	Molecular characterization of zoonotic pathogens Cryptosporidium spp., Giardia duodenalis and Enterocytozoon bieneusi in calves in Algeria. Veterinary Parasitology: Regional Studies and Reports, 2017, 8, 66-69.	0.5	10
115	Longitudinal monitoring of Cryptosporidium species in pre-weaned dairy calves on five farms in Shanghai, China. Veterinary Parasitology, 2017, 241, 14-19.	1.8	51
116	High genetic diversity of Giardia duodenalis assemblage E in pre-weaned dairy calves in Shanghai, China, revealed by multilocus genotyping. Parasitology Research, 2017, 116, 2101-2110.	1.6	31
117	Environmental Transport of Emerging Human-Pathogenic Cryptosporidium Species and Subtypes through Combined Sewer Overflow and Wastewater. Applied and Environmental Microbiology, 2017, 83, .	3.1	50
118	Molecular epidemiologic tools for waterborne pathogens Cryptosporidium spp. and Giardia duodenalis. Food and Waterborne Parasitology, 2017, 8-9, 14-32.	2.7	162
119	Multilocus genotyping of Giardia duodenalis in Tibetan sheep and yaks in Qinghai, China. Veterinary Parasitology, 2017, 247, 70-76.	1.8	32
120	Differences in staining intensities affect reported occurrences and concentrations ofGiardiaspp. in surface drinking water sources. Journal of Applied Microbiology, 2017, 123, 1607-1613.	3.1	8
121	Subtype analysis of zoonotic pathogen Cryptosporidium skunk genotype. Infection, Genetics and Evolution, 2017, 55, 20-25.	2.3	22
122	Molecular Epidemiology of Giardia, Blastocystis and Cryptosporidium among Indigenous Children from the Colombian Amazon Basin. Frontiers in Microbiology, 2017, 8, 248.	3.5	99
123	Preliminary Characterization of MEDLE-2, a Protein Potentially Involved in the Invasion of Cryptosporidium parvum. Frontiers in Microbiology, 2017, 8, 1647.	3.5	16
124	Molecular Epidemiology of Cryptosporidiosis in China. Frontiers in Microbiology, 2017, 8, 1701.	3.5	103
125	Community Laboratory Testing for Cryptosporidium: Multicenter Study Retesting Public Health Surveillance Stool Samples Positive for Cryptosporidium by Rapid Cartridge Assay with Direct Fluorescent Antibody Testing. PLoS ONE, 2017, 12, e0169915.	2.5	20
126	Animal-related factors associated with moderate-to-severe diarrhea in children younger than five years in western Kenya: A matched case-control study. PLoS Neglected Tropical Diseases, 2017, 11, e0005795.	3.0	40

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127	Using Molecular Characterization to Support Investigations of Aquatic Facilityâ€“Associated Outbreaks of Cryptosporidiosis â€" Alabama, Arizona, and Ohio, 2016. Morbidity and Mortality Weekly Report, 2017, 66, 493-497.	15.1	26
128	Cryptosporidium species and subtypes in diarrheal children and HIV-infected persons in Ebonyi and Nsukka, Nigeria. Journal of Infection in Developing Countries, 2017, 11, 173-179.	1.2	33
129	Prevalence, Clinical Manifestations and Genotyping of Spp. in Patients with Gastrointestinal Illnesses in Western Iran. Iranian Journal of Parasitology, 2017, 12, 169-176.	0.6	10
130	Multilocus Sequence Typing Tool forCyclospora cayetanensis. Emerging Infectious Diseases, 2016, 22, 1464-1467.	4.3	38
131	Clonal Evolution of Enterocytozoon bieneusi Populations in Swine and Genetic Differentiation in Subpopulations between Isolates from Swine and Humans. PLoS Neglected Tropical Diseases, 2016, 10, e0004966.	3.0	26
132	Genotypes of Cryptosporidium spp. and Enterocytozoon bieneusi in Human Immunodeficiency Virusâ€Infected Patients in Lagos, Nigeria. Journal of Eukaryotic Microbiology, 2016, 63, 414-418.	1.7	17
133	Fast Technology Analysis Enables Identification of Species and Genotypes of Latent Microsporidia Infections in Healthy Native Cameroonians. Journal of Eukaryotic Microbiology, 2016, 63, 146-152.	1.7	9
134	Cryptosporidium species and Cryptosporidium parvum subtypes in dairy calves and goat kids reared under traditional farming systems in Turkey. Experimental Parasitology, 2016, 170, 16-20.	1.2	34
135	Identity of Fasciola spp. in sheep in Egypt. Parasites and Vectors, 2016, 9, 623.	2.5	42
136	Human infective potential of Cryptosporidium spp., Giardia duodenalis and Enterocytozoon bieneusi in urban wastewater treatment plant effluents. Journal of Water and Health, 2016, 14, 411-423.	2.6	56
137	Evolution of mitosome metabolism and invasion-related proteins in Cryptosporidium. BMC Genomics, 2016, 17, 1006.	2.8	63
138	Genetic variation of mini- and microsatellites and a clonal structure in Enterocytozoon bieneusi population in foxes and raccoon dogs and population differentiation of the parasite between fur animals and humans. Parasitology Research, 2016, 115, 2899-2904.	1.6	26
139	Communitywide cryptosporidiosis outbreak associated with a surface water-supplied municipal water system â€" Baker City, Oregon, 2013. Epidemiology and Infection, 2016, 144, 274-284.	2.1	29
140	Development of a multilocus sequence typing tool for high-resolution subtyping and genetic structure characterization of Cryptosporidium ubiquitum. Infection, Genetics and Evolution, 2016, 45, 256-261.	2.3	14
141	Comparative genomics reveals adaptive evolution of Asian tapeworm in switching to a new intermediate host. Nature Communications, 2016, 7, 12845.	12.8	43
142	Annotated draft genome sequences of three species ofCryptosporidium:Cryptosporidium meleagridisisolate UKMEL1,C. baileyiisolate TAMU-09Q1 andC. hominisisolates TU502_2012 and UKH1. Pathogens and Disease, 2016, 74, ftw080.	2.0	33
143	Cryptosporidium canis in Two Mexican Toddlers. Pediatric Infectious Disease Journal, 2016, 35, 1265-1266.	2.0	6
144	Common occurrence of Cryptosporidium hominis in horses and donkeys. Infection, Genetics and Evolution, 2016, 43, 261-266.	2.3	37

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145	Comparative genomics reveals Cyclospora cayetanensis possesses coccidia-like metabolism and invasion components but unique surface antigens. BMC Genomics, 2016, 17, 316.	2.8	42
146	Distribution of Cryptosporidium species in Tibetan sheep and yaks in Qinghai, China. Veterinary Parasitology, 2016, 215, 58-62.	1.8	52
147	Genotypes of Cryptosporidium spp., Enterocytozoon bieneusi and Giardia duodenalis in dogs and cats in Shanghai, China. Parasites and Vectors, 2016, 9, 121.	2.5	84
148	Cryptosporidium proliferans n. sp. (Apicomplexa: Cryptosporidiidae): Molecular and Biological Evidence of Cryptic Species within Gastric Cryptosporidium of Mammals. PLoS ONE, 2016, 11, e0147090.	2.5	68
149	Hypothesis: Cryptosporidium genetic diversity mirrors national disease notification rate. Parasites and Vectors, 2015, 8, 308.	2.5	3
150	Genetic similarities between Cyclospora cayetanensis and cecum-infecting avian Eimeria spp. in apicoplast and mitochondrial genomes. Parasites and Vectors, 2015, 8, 358.	2.5	40
151	Molecular Characterization of Echinococcus granulosus Sensu Lato from Farm Animals in Egypt. PLoS ONE, 2015, 10, e0118509.	2.5	44
152	Morphologic and Genotypic Characterization of Psoroptes Mites from Water Buffaloes in Egypt. PLoS ONE, 2015, 10, e0141554.	2.5	3
153	Epidemiological Observations on Cryptosporidiosis in Diarrheic Goat Kids in Greece. Veterinary Medicine International, 2015, 2015, 1-4.	1.5	10
154	Subtyping Novel Zoonotic Pathogen Cryptosporidium Chipmunk Genotype I. Journal of Clinical Microbiology, 2015, 53, 1648-1654.	3.9	57
155	Comparative genomic analysis reveals occurrence of genetic recombination in virulent Cryptosporidium hominis subtypes and telomeric gene duplications in Cryptosporidium parvum. BMC Genomics, 2015, 16, 320.	2.8	74
156	Cryptosporidium genotypes and subtypes distribution in river water in Iran. Journal of Water and Health, 2015, 13, 600-606.	2.6	16
157	Multi-locus analysis of Giardia duodenalis from nonhuman primates kept in zoos in China: Geographical segregation and host-adaptation of assemblage B isolates. Infection, Genetics and Evolution, 2015, 30, 82-88.	2.3	37
158	Complex epidemiology and zoonotic potential for Cryptosporidium suis in rural Madagascar. Veterinary Parasitology, 2015, 207, 140-143.	1.8	38
159	Isolation and Enrichment of Cryptosporidium DNA and Verification of DNA Purity for Whole-Genome Sequencing. Journal of Clinical Microbiology, 2015, 53, 641-647.	3.9	45
160	Cryptosporidium huwi n. sp. (Apicomplexa: Eimeriidae) from the guppy (Poecilia reticulata). Experimental Parasitology, 2015, 150, 31-35.	1.2	64
161	Prevalence and genetic characteristics of Cryptosporidium, Enterocytozoon bieneusi and Giardia duodenalis in cats and dogs in Heilongjiang province, China. Veterinary Parasitology, 2015, 208, 125-134.	1.8	114
162	The First Association of a Primary Amebic Meningoencephalitis Death With Culturable Naegleria fowleri in Tap Water From a US Treated Public Drinking Water System. Clinical Infectious Diseases, 2015, 60, e36-e42.	5.8	84

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