

Yao-Yu Feng

List of Publications by Year in descending order

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200
papers

11,399
citations

34493

54
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39744

98
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206
all docs

206
docs citations

206
times ranked

9185
citing authors

#	ARTICLE	IF	CITATIONS
1	Cryptosporidiosis outbreak caused by <i>Cryptosporidium parvum</i> subtype IIdA20G1 in neonatal calves. <i>Transboundary and Emerging Diseases</i> , 2022, 69, 278-285.	1.3	11
2	<i>Enterocytozoon bienersi</i> . <i>Trends in Parasitology</i> , 2022, 38, 95-96.	1.5	16
3	Comparative Characterization of CpCDPK1 and CpCDPK9, Two Potential Drug Targets Against Cryptosporidiosis. <i>Microorganisms</i> , 2022, 10, 333.	1.6	5
4	Emergence of zoonotic <i>Cryptosporidium parvum</i> in China. <i>Trends in Parasitology</i> , 2022, 38, 335-343.	1.5	24
5	A productive immunocompetent mouse model of cryptosporidiosis with long oocyst shedding duration for immunological studies. <i>Journal of Infection</i> , 2022, 84, 710-721.	1.7	7
6	High zoonotic potential of <i>Cryptosporidium</i> spp., <i>Giardia duodenalis</i> , and <i>Enterocytozoon bienersi</i> in wild nonhuman primates from Yunnan Province, China. <i>Parasites and Vectors</i> , 2022, 15, 85.	1.0	5
7	Molecular detection of a novel <i>Ancylostoma</i> sp. by whole mtDNA sequence from pangolin <i>Manis javanica</i> . <i>Parasites and Vectors</i> , 2022, 15, 70.	1.0	3
8	Age and episode-associated occurrence of <i>Cryptosporidium</i> species and subtypes in a birth cohort of dairy calves. <i>Transboundary and Emerging Diseases</i> , 2022, 69, .	1.3	3
9	Diarrhoea outbreak caused by coinfections of <i>Cryptosporidium parvum</i> subtype IIdA20G1 and rotavirus in pre-weaned dairy calves. <i>Transboundary and Emerging Diseases</i> , 2022, 69, .	1.3	8
10	Characterization of Calcium-Dependent Protein Kinase 2A, a Potential Drug Target Against Cryptosporidiosis. <i>Frontiers in Microbiology</i> , 2022, 13, 883674.	1.5	2
11	Characterization of Dense Granule Metalloproteinase INS-16 in <i>Cryptosporidium parvum</i> . <i>International Journal of Molecular Sciences</i> , 2022, 23, 7617.	1.8	3
12	Sympatric Recombination in Zoonotic <i>Cryptosporidium</i> Leads to Emergence of Populations with Modified Host Preference. <i>Molecular Biology and Evolution</i> , 2022, 39, .	3.5	10
13	<i>Cryptosporidium ratti</i> n. sp. (Apicomplexa: Cryptosporidiidae) and genetic diversity of <i>Cryptosporidium</i> spp. in brown rats (<i>Rattus norvegicus</i>) in the Czech Republic. <i>Parasitology</i> , 2021, 148, 84-97.	0.7	24
14	Characterizations of <i>Enterocytozoon bienersi</i> at new genetic loci reveal a lack of strict host specificity among common genotypes and the existence of a canine-adapted <i>Enterocytozoon</i> species. <i>International Journal for Parasitology</i> , 2021, 51, 215-223.	1.3	9
15	Development of a Subtyping Tool for Zoonotic Pathogen <i>Cryptosporidium canis</i> . <i>Journal of Clinical Microbiology</i> , 2021, 59, .	1.8	20
16	Cryptosporidial Infection Suppresses Intestinal Epithelial Cell MAPK Signaling Impairing Host Anti-Parasitic Defense. <i>Microorganisms</i> , 2021, 9, 151.	1.6	11
17	Subtype Characterization and Zoonotic Potential of <i>Cryptosporidium felis</i> in Cats in Guangdong and Shanghai, China. <i>Pathogens</i> , 2021, 10, 89.	1.2	8
18	Molecular Epidemiology of Human Cryptosporidiosis in Low- and Middle-Income Countries. <i>Clinical Microbiology Reviews</i> , 2021, 34, .	5.7	56

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19	Occurrence and genetic diversity of <i>Cryptosporidium</i> spp. in wild foxes, wolves, jackals, and bears in central Europe. <i>Folia Parasitologica</i> , 2021, 68, .	0.7	9
20	Small ruminants and zoonotic cryptosporidiosis. <i>Parasitology Research</i> , 2021, 120, 4189-4198.	0.6	28
21	Zoonotic parasites in farmed exotic animals in China: Implications to public health. <i>International Journal for Parasitology: Parasites and Wildlife</i> , 2021, 14, 241-247.	0.6	9
22	Impact of <i>mcr-1</i> on the Development of High Level Colistin Resistance in <i>Klebsiella pneumoniae</i> and <i>Escherichia coli</i> . <i>Frontiers in Microbiology</i> , 2021, 12, 666782.	1.5	10
23	Codon usage analysis of zoonotic coronaviruses reveals lower adaptation to humans by SARS-CoV-2. <i>Infection, Genetics and Evolution</i> , 2021, 89, 104736.	1.0	13
24	Comparative Study of Two Insulinlike Proteases in <i>Cryptosporidium parvum</i> . <i>Microorganisms</i> , 2021, 9, 861.	1.6	3
25	<i>Cryptosporidium myocastoris</i> n. sp. (Apicomplexa: Cryptosporidiidae), the Species Adapted to the Nutria (<i>Myocastor coypus</i>). <i>Microorganisms</i> , 2021, 9, 813.	1.6	35
26	Insulinase-like Protease 1 Contributes to Macrogamont Formation in <i>Cryptosporidium parvum</i> . <i>MBio</i> , 2021, 12, .	1.8	10
27	Preliminary Characterization of Two Small Insulinase-Like Proteases in <i>Cryptosporidium parvum</i> . <i>Frontiers in Microbiology</i> , 2021, 12, 651512.	1.5	3
28	Subtyping <i>Cryptosporidium xiaoi</i> , a Common Pathogen in Sheep and Goats. <i>Pathogens</i> , 2021, 10, 800.	1.2	11
29	Molecular detection of <i>Cryptosporidium</i> spp., <i>Giardia duodenalis</i> , and <i>Enterocytozoon bienersi</i> in school children at the Thai-Myanmar border. <i>Parasitology Research</i> , 2021, 120, 2887-2895.	0.6	4
30	Genus-level evolutionary relationships of FAR proteins reflect the diversity of lifestyles of free-living and parasitic nematodes. <i>BMC Biology</i> , 2021, 19, 178.	1.7	4
31	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 <i>C. hominis</i> IbA12G3 subtype. <i>Infection, Genetics and Evolution</i> , 2021, 92, 104859.	1.0	12
32	Advances in molecular epidemiology of cryptosporidiosis in dogs and cats. <i>International Journal for Parasitology</i> , 2021, 51, 787-795.	1.3	13
33	Genetic characterizations of <i>Cryptosporidium</i> spp. from pet rodents indicate high zoonotic potential of pathogens from chinchillas. <i>One Health</i> , 2021, 13, 100269.	1.5	5
34	Molecular characterization of the waterborne pathogens <i>Cryptosporidium</i> spp., <i>Giardia duodenalis</i> , <i>Enterocytozoon bienersi</i> , <i>Cyclospora cayatanensis</i> and <i>Eimeria</i> spp. in wastewater and sewage in Guangzhou, China. <i>Parasites and Vectors</i> , 2021, 14, 66.	1.0	17
35	Development and Application of a gp60-Based Subtyping Tool for <i>Cryptosporidium bovis</i> . <i>Microorganisms</i> , 2021, 9, 2067.	1.6	8
36	Taxonomy and molecular epidemiology of <i>Cryptosporidium</i> and <i>Giardia</i> – a 50-year perspective (1971–2021). <i>International Journal for Parasitology</i> , 2021, 51, 1099-1119.	1.3	128

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37	Zoonotic giardiasis: an update. <i>Parasitology Research</i> , 2021, 120, 4199-4218.	0.6	71
38	An Update on Zoonotic <i>Cryptosporidium</i> Species and Genotypes in Humans. <i>Animals</i> , 2021, 11, 3307.	1.0	84
39	Association of Common Zoonotic Pathogens With Concentrated Animal Feeding Operations. <i>Frontiers in Microbiology</i> , 2021, 12, 810142.	1.5	6
40	<i>Cryptosporidium felis</i> differs from other <i>Cryptosporidium</i> spp. in codon usage. <i>Microbial Genomics</i> , 2021, 7, .	1.0	3
41	Population genetic analysis suggests genetic recombination is responsible for increased zoonotic potential of <i>Enterocytozoon bienewsi</i> from ruminants in China. <i>One Health</i> , 2020, 11, 100184.	1.5	7
42	Subtype distribution of zoonotic pathogen <i>Cryptosporidium felis</i> in humans and animals in several countries. <i>Emerging Microbes and Infections</i> , 2020, 9, 2446-2454.	3.0	19
43	Diagnosis and molecular typing of <i>Enterocytozoon bienewsi</i> : the significant role of domestic animals in transmission of human microsporidiosis. <i>Research in Veterinary Science</i> , 2020, 133, 251-261.	0.9	29
44	Occurrence and molecular characterization of <i>Giardia duodenalis</i> in lambs in Djelfa, the central steppe of Algeria. <i>Parasitology Research</i> , 2020, 119, 2965-2973.	0.6	4
45	<i>Cryptosporidium</i> Species and <i>C. parvum</i> Subtypes in Farmed Bamboo Rats. <i>Pathogens</i> , 2020, 9, 1018.	1.2	8
46	Contribution of hospitals to the occurrence of enteric protists in urban wastewater. <i>Parasitology Research</i> , 2020, 119, 3033-3040.	0.6	12
47	Molecular characterization and zoonotic potential of <i>Enterocytozoon bienewsi</i> , <i>Giardia duodenalis</i> and <i>Cryptosporidium</i> sp. in farmed masked palm civets (<i>Paguma larvata</i>) in southern China. <i>Parasites and Vectors</i> , 2020, 13, 403.	1.0	19
48	Subtyping <i>Cryptosporidium ryanae</i> : A Common Pathogen in Bovine Animals. <i>Microorganisms</i> , 2020, 8, 1107.	1.6	18
49	Population structure and geographical segregation of <i>Cryptosporidium parvum</i> IId subtypes in cattle in China. <i>Parasites and Vectors</i> , 2020, 13, 425.	1.0	15
50	Characterization of Calcium-Dependent Protein Kinases 3, a Protein Involved in Growth of <i>Cryptosporidium parvum</i> . <i>Frontiers in Microbiology</i> , 2020, 11, 907.	1.5	8
51	Expression and Functional Studies of INS-5, an Insulinase-Like Protein in <i>Cryptosporidium parvum</i> . <i>Frontiers in Microbiology</i> , 2020, 11, 719.	1.5	7
52	Common occurrence of divergent <i>Cryptosporidium</i> species and <i>Cryptosporidium parvum</i> subtypes in farmed bamboo rats (<i>Rhizomys sinensis</i>). <i>Parasites and Vectors</i> , 2020, 13, 149.	1.0	19
53	Isolation of SARS-CoV-2-related coronavirus from Malayan pangolins. <i>Nature</i> , 2020, 583, 286-289.	13.7	599
54	Zoonotic potential of <i>Enterocytozoon bienewsi</i> and <i>Giardia duodenalis</i> in horses and donkeys in northern China. <i>Parasitology Research</i> , 2020, 119, 1101-1108.	0.6	20

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55	Multilocus sequence typing of <i>Enterocytozoon bieneusi</i> in crab-eating macaques (<i>Macaca</i>) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50	1.0	10
56	Characterization of Three Calcium-Dependent Protein Kinases of <i>Cryptosporidium parvum</i> . <i>Frontiers in Microbiology</i> , 2020, 11, 622203.	1.5	6
57	Comparative genomic analysis of three intestinal species reveals reductions in secreted pathogenesis determinants in bovine-specific and non-pathogenic <i>Cryptosporidium</i> species. <i>Microbial Genomics</i> , 2020, 6, .	1.0	13
58	Infection patterns, clinical significance, and genetic characteristics of <i>Enterocytozoon bieneusi</i> and <i>Giardia duodenalis</i> in dairy cattle in Jiangsu, China. <i>Parasitology Research</i> , 2019, 118, 3053-3060.	0.6	30
59	<i>Cryptosporidium parvum</i> and <i>Cryptosporidium hominis</i> subtypes in crab-eating macaques. <i>Parasites and Vectors</i> , 2019, 12, 350.	1.0	26
60	Different distribution of <i>Cryptosporidium</i> species between horses and donkeys. <i>Infection, Genetics and Evolution</i> , 2019, 75, 103954.	1.0	21
61	Characterization of INS-15, A Metalloprotease Potentially Involved in the Invasion of <i>Cryptosporidium parvum</i> . <i>Microorganisms</i> , 2019, 7, 452.	1.6	16
62	Divergent Copies of a <i>Cryptosporidium parvum</i> -Specific Subtelomeric Gene. <i>Microorganisms</i> , 2019, 7, 366.	1.6	4
63	Comparative genomics: how has it advanced our knowledge of cryptosporidiosis epidemiology?. <i>Parasitology Research</i> , 2019, 118, 3195-3204.	0.6	17
64	Epidemiological distribution of genotypes of <i>Giardia duodenalis</i> in humans in Spain. <i>Parasites and Vectors</i> , 2019, 12, 432.	1.0	29
65	Potential impacts of host specificity on zoonotic or interspecies transmission of <i>Enterocytozoon bieneusi</i> . <i>Infection, Genetics and Evolution</i> , 2019, 75, 104033.	1.0	47
66	Prevalence and genotypic identification of <i>Cryptosporidium</i> spp., <i>Giardia duodenalis</i> and <i>Enterocytozoon bieneusi</i> in pre-weaned dairy calves in Guangdong, China. <i>Parasites and Vectors</i> , 2019, 12, 41.	1.0	55
67	Genotypes and public health potential of <i>Enterocytozoon bieneusi</i> and <i>Giardia duodenalis</i> in crab-eating macaques. <i>Parasites and Vectors</i> , 2019, 12, 254.	1.0	22
68	Comparative analysis reveals conservation in genome organization among intestinal <i>Cryptosporidium</i> species and sequence divergence in potential secreted pathogenesis determinants among major human-infecting species. <i>BMC Genomics</i> , 2019, 20, 406.	1.2	37
69	Differential Expression of Three <i>Cryptosporidium</i> Species-Specific MEDLE Proteins. <i>Frontiers in Microbiology</i> , 2019, 10, 1177.	1.5	11
70	Host Specificity of <i>Enterocytozoon bieneusi</i> and Public Health Implications. <i>Trends in Parasitology</i> , 2019, 35, 436-451.	1.5	196
71	Outbreak of cryptosporidiosis due to <i>Cryptosporidium parvum</i> subtype IIdA19G1 in neonatal calves on a dairy farm in China. <i>International Journal for Parasitology</i> , 2019, 49, 569-577.	1.3	39
72	Characterization of a Species-Specific Insulinase-Like Protease in <i>Cryptosporidium parvum</i> . <i>Frontiers in Microbiology</i> , 2019, 10, 354.	1.5	18

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73	Mitochondrial genome sequence variation as a useful marker for assessing genetic heterogeneity among <i>Cyclospora cayetanensis</i> isolates and source-tracking. <i>Parasites and Vectors</i> , 2019, 12, 47.	1.0	13
74	Genetic characterization of <i>Cryptosporidium</i> spp. and <i>Giardia duodenalis</i> in dogs and cats in Guangdong, China. <i>Parasites and Vectors</i> , 2019, 12, 571.	1.0	28
75	Host-adapted <i>Cryptosporidium</i> and <i>Enterocytozoon bienewsi</i> genotypes in straw-colored fruit bats in Nigeria. <i>International Journal for Parasitology: Parasites and Wildlife</i> , 2019, 8, 19-24.	0.6	17
76	<i>Giardia</i> : an under-reported foodborne parasite. <i>International Journal for Parasitology</i> , 2019, 49, 1-11.	1.3	131
77	Divergent <i>Cryptosporidium parvum</i> subtype and <i>Enterocytozoon bienewsi</i> genotypes in dromedary camels in Algeria. <i>Parasitology Research</i> , 2018, 117, 905-910.	0.6	21
78	Population genetic characterization of <i>Cyclospora cayetanensis</i> from discrete geographical regions. <i>Experimental Parasitology</i> , 2018, 184, 121-127.	0.5	11
79	<i>Enterocytozoon bienewsi</i> genotypes in Tibetan sheep and yaks. <i>Parasitology Research</i> , 2018, 117, 721-727.	0.6	37
80	Epidemiological observations on cryptosporidiosis and molecular characterization of <i>Cryptosporidium</i> spp. in sheep and goats in Kuwait. <i>Parasitology Research</i> , 2018, 117, 1631-1636.	0.6	26
81	Genotypes and subtypes of <i>Cryptosporidium</i> spp. in diarrheic lambs and goat kids in northern Greece. <i>Parasitology International</i> , 2018, 67, 472-475.	0.6	25
82	Zoonotic <i>Cryptosporidium</i> species and subtypes in lambs and goat kids in Algeria. <i>Parasites and Vectors</i> , 2018, 11, 582.	1.0	30
83	Persistent Occurrence of <i>Cryptosporidium hominis</i> and <i>Giardia duodenalis</i> Subtypes in a Welfare Institute. <i>Frontiers in Microbiology</i> , 2018, 9, 2830.	1.5	13
84	Catalytic N ₂ O decomposition over CeMeO _y /Al ₂ O ₃ (Me=Mn, Cu, Zn) catalysts prepared by impregnation method. <i>Asia-Pacific Journal of Chemical Engineering</i> , 2018, 13, e2233.	0.8	4
85	Molecular characterization of <i>Cryptosporidium</i> spp. and <i>Giardia duodenalis</i> in children in Egypt. <i>Parasites and Vectors</i> , 2018, 11, 403.	1.0	40
86	Genetic diversity within dominant <i>Enterocytozoon bienewsi</i> genotypes in pre-weaned calves. <i>Parasites and Vectors</i> , 2018, 11, 170.	1.0	32
87	Characterization of MEDLE-1, a protein in early development of <i>Cryptosporidium parvum</i> . <i>Parasites and Vectors</i> , 2018, 11, 312.	1.0	14
88	Genetic Diversity and Population Structure of <i>Cryptosporidium</i> . <i>Trends in Parasitology</i> , 2018, 34, 997-1011.	1.5	365
89	Ginkgolide B ameliorates oxidized low-density lipoprotein-induced endothelial dysfunction via modulating Lectin-like oxLDL receptor 1 and NADPH oxidase 4 expression and inflammatory cascades. <i>Phytotherapy Research</i> , 2018, 32, 2417-2427.	2.8	27
90	Comparative genomic analysis of the IId subtype family of <i>Cryptosporidium parvum</i> . <i>International Journal for Parasitology</i> , 2017, 47, 281-290.	1.3	58

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91	Longitudinal monitoring of <i>Cryptosporidium</i> species in pre-weaned dairy calves on five farms in Shanghai, China. <i>Veterinary Parasitology</i> , 2017, 241, 14-19.	0.7	51
92	High genetic diversity of <i>Giardia duodenalis</i> assemblage E in pre-weaned dairy calves in Shanghai, China, revealed by multilocus genotyping. <i>Parasitology Research</i> , 2017, 116, 2101-2110.	0.6	31
93	Environmental Transport of Emerging Human-Pathogenic <i>Cryptosporidium</i> Species and Subtypes through Combined Sewer Overflow and Wastewater. <i>Applied and Environmental Microbiology</i> , 2017, 83, .	1.4	50
94	Molecular epidemiologic tools for waterborne pathogens <i>Cryptosporidium</i> spp. and <i>Giardia duodenalis</i> . <i>Food and Waterborne Parasitology</i> , 2017, 8-9, 14-32.	1.1	162
95	Multilocus genotyping of <i>Giardia duodenalis</i> in Tibetan sheep and yaks in Qinghai, China. <i>Veterinary Parasitology</i> , 2017, 247, 70-76.	0.7	32
96	Subtype analysis of zoonotic pathogen <i>Cryptosporidium</i> skunk genotype. <i>Infection, Genetics and Evolution</i> , 2017, 55, 20-25.	1.0	22
97	Preliminary Characterization of MEDLE-2, a Protein Potentially Involved in the Invasion of <i>Cryptosporidium parvum</i> . <i>Frontiers in Microbiology</i> , 2017, 8, 1647.	1.5	16
98	Molecular Epidemiology of Cryptosporidiosis in China. <i>Frontiers in Microbiology</i> , 2017, 8, 1701.	1.5	103
99	Multilocus Sequence Typing Tool for <i>Cyclospora cayetanensis</i> . <i>Emerging Infectious Diseases</i> , 2016, 22, 1464-1467.	2.0	38
100	Dominant genera of cyanobacteria in Lake Taihu and their relationships with environmental factors. <i>Journal of Microbiology</i> , 2016, 54, 468-476.	1.3	17
101	Genotypes of <i>Cryptosporidium</i> spp. and <i>Enterocytozoon bienersi</i> in Human Immunodeficiency Virus-Infected Patients in Lagos, Nigeria. <i>Journal of Eukaryotic Microbiology</i> , 2016, 63, 414-418.	0.8	17
102	<i>Cryptosporidium</i> species and <i>Cryptosporidium parvum</i> subtypes in dairy calves and goat kids reared under traditional farming systems in Turkey. <i>Experimental Parasitology</i> , 2016, 170, 16-20.	0.5	34
103	Identity of <i>Fasciola</i> spp. in sheep in Egypt. <i>Parasites and Vectors</i> , 2016, 9, 623.	1.0	42
104	Human infective potential of <i>Cryptosporidium</i> spp., <i>Giardia duodenalis</i> and <i>Enterocytozoon bienersi</i> in urban wastewater treatment plant effluents. <i>Journal of Water and Health</i> , 2016, 14, 411-423.	1.1	56
105	Evolution of mitosome metabolism and invasion-related proteins in <i>Cryptosporidium</i> . <i>BMC Genomics</i> , 2016, 17, 1006.	1.2	63
106	Development of a multilocus sequence typing tool for high-resolution subtyping and genetic structure characterization of <i>Cryptosporidium ubiquitum</i> . <i>Infection, Genetics and Evolution</i> , 2016, 45, 256-261.	1.0	14
107	Comparative genomics reveals <i>Cyclospora cayetanensis</i> possesses coccidia-like metabolism and invasion components but unique surface antigens. <i>BMC Genomics</i> , 2016, 17, 316.	1.2	42
108	Distribution of <i>Cryptosporidium</i> species in Tibetan sheep and yaks in Qinghai, China. <i>Veterinary Parasitology</i> , 2016, 215, 58-62.	0.7	52

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109	Genotypes of <i>Cryptosporidium</i> spp., <i>Enterocytozoon bienersi</i> and <i>Giardia duodenalis</i> in dogs and cats in Shanghai, China. <i>Parasites and Vectors</i> , 2016, 9, 121.	1.0	84
110	Correlation Between ABCA1 Gene Polymorphism and aopA-I and HDL-C in Abdominal Aortic Aneurysm. <i>Medical Science Monitor</i> , 2016, 22, 172-176.	0.5	9
111	Genetic similarities between <i>Cyclospora cayatanensis</i> and cecum-infecting avian <i>Eimeria</i> spp. in apicoplast and mitochondrial genomes. <i>Parasites and Vectors</i> , 2015, 8, 358.	1.0	40
112	Molecular Characterization of <i>Echinococcus granulosus</i> Sensu Lato from Farm Animals in Egypt. <i>PLoS ONE</i> , 2015, 10, e0118509.	1.1	44
113	Morphologic and Genotypic Characterization of Psoroptes Mites from Water Buffaloes in Egypt. <i>PLoS ONE</i> , 2015, 10, e0141554.	1.1	3
114	Subtyping Novel Zoonotic Pathogen <i>Cryptosporidium</i> Chipmunk Genotype I. <i>Journal of Clinical Microbiology</i> , 2015, 53, 1648-1654.	1.8	57
115	Comparative genomic analysis reveals occurrence of genetic recombination in virulent <i>Cryptosporidium hominis</i> subtypes and telomeric gene duplications in <i>Cryptosporidium parvum</i> . <i>BMC Genomics</i> , 2015, 16, 320.	1.2	74
116	Isolation and Enrichment of <i>Cryptosporidium</i> DNA and Verification of DNA Purity for Whole-Genome Sequencing. <i>Journal of Clinical Microbiology</i> , 2015, 53, 641-647.	1.8	45
117	Occurrence and molecular characterization of <i>Cryptosporidium</i> spp. and <i>Enterocytozoon bienersi</i> in dairy cattle, beef cattle and water buffaloes in China. <i>Veterinary Parasitology</i> , 2015, 207, 220-227.	0.7	108
118	Identification and morphologic and molecular characterization of <i>Cyclospora macacae</i> n. sp. from rhesus monkeys in China. <i>Parasitology Research</i> , 2015, 114, 1811-1816.	0.6	32
119	Dominance of <i>Giardia duodenalis</i> assemblage A and <i>Enterocytozoon bienersi</i> genotype BEB6 in sheep in Inner Mongolia, China. <i>Veterinary Parasitology</i> , 2015, 210, 235-239.	0.7	57
120	Development and Evaluation of Three Real-Time PCR Assays for Genotyping and Source Tracking <i>Cryptosporidium</i> spp. in Water. <i>Applied and Environmental Microbiology</i> , 2015, 81, 5845-5854.	1.4	27
121	<i>Enterocytozoon bienersi</i> Genotypes in Yaks (<i>Bos grunniens</i>) and Their Public Health Potential. <i>Journal of Eukaryotic Microbiology</i> , 2015, 62, 21-25.	0.8	28
122	Preliminary Molecular Characterizations of <i>Sarcoptes scabiei</i> (Acari: Sarcoptidae) from Farm Animals in Egypt. <i>PLoS ONE</i> , 2014, 9, e94705.	1.1	25
123	Distribution and Clinical Manifestations of <i>Cryptosporidium</i> Species and Subtypes in HIV/AIDS Patients in Ethiopia. <i>PLoS Neglected Tropical Diseases</i> , 2014, 8, e2831.	1.3	133
124	Occurrence, Source, and Human Infection Potential of <i>Cryptosporidium</i> and <i>Enterocytozoon bienersi</i> in Drinking Source Water in Shanghai, China, during a Pig Carcass Disposal Incident. <i>Environmental Science & Technology</i> , 2014, 48, 14219-14227.	4.6	88
125	Non-coding RNAs in epithelial immunity to <i>Cryptosporidium</i> infection. <i>Parasitology</i> , 2014, 141, 1233-1243.	0.7	38
126	Host Specificity and Source of <i>Enterocytozoon bienersi</i> Genotypes in a Drinking Source Watershed. <i>Applied and Environmental Microbiology</i> , 2014, 80, 218-225.	1.4	104

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127	Multilocus Sequence Typing of an Emerging <i>Cryptosporidium hominis</i> Subtype in the United States. <i>Journal of Clinical Microbiology</i> , 2014, 52, 524-530.	1.8	47
128	Population genetics of <i>Cryptosporidium meleagridis</i> in humans and birds: evidence for cross-species transmission. <i>International Journal for Parasitology</i> , 2014, 44, 515-521.	1.3	44
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