Aiqin Liu

List of Publications by Year
in descending order

40 papers	1,297 citations	279798 23 h-index	345221 36 g-index
40 all docs	40 docs citations	40 times ranked	881 citing authors

#	Article	IF	Citations
1	Distribution and genetic diversity of Blastocystis subtypes in various mammal and bird species in northeastern China. Parasites and Vectors, 2018, 11, 522.	2.5	77
2	Subtype distribution and genetic characterizations of Blastocystis in pigs, cattle, sheep and goats in northeastern China's Heilongjiang Province. Infection, Genetics and Evolution, 2018, 57, 171-176.	2.3	76
3	Distribution and Genetic Characterizations of Cryptosporidium spp. in Pre-Weaned Dairy Calves in Northeastern Chinaâ€™s Heilongjiang Province. PLoS ONE, 2013, 8, e54857.	2.5	69
4	Enterocytozoon bieneusi in Dairy Cattle in the Northeast of China: Genetic Diversity of <scp>ITS</scp> Gene and Evaluation of Zoonotic Transmission Potential. Journal of Eukaryotic Microbiology, 2015, 62, 553-560.	1.7	58
5	Cryptosporidium parvum IId family: clonal population and dispersal from Western Asia to other geographical regions. Scientific Reports, 2014, 4, 4208.	3.3	58
6	Prevalence of Enterocytozoon bieneusi and genetic diversity of ITS genotypes in sheep and goats in China. Infection, Genetics and Evolution, 2015, 32, 265-270.	2.3	55
7	Cryptosporidium andersoni as a novel predominant Cryptosporidium species in outpatients with diarrhea in Jiangsu Province, China. BMC Infectious Diseases, 2014, 14, 555.	2.9	54
8	Genotyping of Enterocytozoon bieneusi and Subtyping of Blastocystis in Cancer Patients: Relationship to Diarrhea and Assessment of Zoonotic Transmission. Frontiers in Microbiology, 2017, 8, 1835.	3.5	53
9	Prevalence and distribution of Cryptosporidium spp. in dairy cattle in Heilongjiang Province, China. Parasitology Research, 2009, 105, 797-802.	1.6	48
10	A retrospective epidemiological analysis of human Cryptosporidium infection in China during the past three decades (1987-2018). PLoS Neglected Tropical Diseases, 2020, 14, e0008146.	3.0	48
11	Molecular characterizations of Cryptosporidium spp. and Enterocytozoon bieneusi in brown rats (Rattus norvegicus) from Heilongjiang Province, China. Parasites and Vectors, 2018, 11, 313.	2.5	47
12	Occurrence of bovine giardiasis and endemic genetic characterization of Giardia duodenalis isolates in Heilongjiang Province, in the Northeast of China. Parasitology Research, 2012, 111, 655-661.	1.6	45
13	Genotyping of Enterocytozoon bieneusi (Microsporidia) isolated from various birds in China. Infection, Genetics and Evolution, 2016, 40, 151-154.	2.3	44
14	Common occurrence of Cryptosporidium hominis in horses and donkeys. Infection, Genetics and Evolution, 2016, 43, 261-266.	2.3	37
15	Multilocus Sequence Subtyping and Genetic Structure of Cryptosporidium muris and Cryptosporidium andersoni. PLoS ONE, 2012, 7, e43782.	2.5	35
16	Genetic Characterization of Human-Derived Hydatid Cysts of Echinococcus granulosus Sensu Lato in Heilongjiang Province and the First Report of G7 Genotype of E. canadensis in Humans in China. PLoS ONE, 2014, 9, e109059.	2.5	35
17	Molecular identification and distribution of Cryptosporidium and Giardia duodenalis in raw urban wastewater in Harbin, China. Parasitology Research, 2011, 109, 913-918.	1.6	34
18	Genotyping of Enterocytozoon bieneusi in Farmed Blue Foxes (Alopex lagopus) and Raccoon Dogs (Nyctereutes procyonoides) in China. PLoS ONE, 2015, 10, e0142611.	2.5	33

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19	Genetic characterization of three mitochondrial gene sequences of goat/sheep-derived coenurus cerebralis and cysticercus tenuicollis isolates in Inner Mongolia, China. Parasite, 2018, 25, 1.	2.0	33
20	Genetic Analysis of the Gdh and Bg Genes of Animal-Derived Giardia duodenalis Isolates in Northeastern China and Evaluation of Zoonotic Transmission Potential. PLoS ONE, 2014, 9, e95291.	2.5	30
21	Subtyping ofCryptosporidium cuniculusand genotyping ofEnterocytozoon bieneusiin rabbits in two farms in Heilongjiang Province, China. Parasite, 2016, 23, 52.	2.0	29
22	Prevalence and Genetic Characterizations of Cryptosporidium spp. in Pre-Weaned and Post-Weaned Piglets in Heilongjiang Province, China. PLoS ONE, 2013, 8, e67564.	2.5	26
23	The first report of human-derived G10 genotype of Echinococcus canadensis in China and possible sources and routes of transmission. Parasitology International, 2015, 64, 330-333.	1.3	26
24	The first report of Cryptosporidium andersoni in horses with diarrhea and multilocus subtype analysis. Parasites and Vectors, 2015, 8, 483.	2.5	25
25	First survey of Enterocytozoon bieneusi and dominant genotype Peru6 among ethnic minority groups in southwestern Chinaâ€™s Yunnan Province and assessment of risk factors. PLoS Neglected Tropical Diseases, 2019, 13, e0007356.	3.0	23
26	First detection and genotyping of Enterocytozoon bieneusi in reindeers (Rangifer tarandus): a zoonotic potential of ITS genotypes. Parasites and Vectors, 2015, 8, 526.	2.5	22
27	Human cystic echinococcosis in Heilongjiang Province, China: a retrospective study. BMC Gastroenterology, 2015, 15, 29.	2.0	22
28	Dominance of the Enterocytozoon bieneusi genotype BEB6 in red deer (Cervus elaphus) and Siberian roe deer (Capreolus pygargus) in China and a brief literature review. Parasite, 2017, 24, 54.	2.0	22
29	MLST Subtypes and Population Genetic Structure of Cryptosporidium andersoni from Dairy Cattle and Beef Cattle in Northeastern Chinaâ€™s Heilongjiang Province. PLoS ONE, 2014, 9, e102006.	2.5	20
30	First identification and genotyping of Enterocytozoon bieneusi in humans in Myanmar. BMC Microbiology, 2020, 20, 10.	3.3	17
31	Prevalence and subtype distribution of Blastocystis in ethnic minority groups on both sides of the Chinaâ€“Myanmar border, and assessment of risk factors. Parasite, 2019, 26, 46.	2.0	16
32	A human case of Dioctophyma renale (giant kidney worm) accompanied by renal cancer and a retrospective study of dioctophymiasis. Parasite, 2019, 26, 22.	2.0	15
33	Prevalence of Hymenolepis nana and H. diminuta from Brown Rats (Rattus norvegicus) in Heilongjiang Province, China. Korean Journal of Parasitology, 2017, 55, 351-355.	1.3	14
34	Molecular detection and genetic characterizations of Cryptosporidium spp. in farmed foxes, minks, and raccoon dogs in northeastern China. Parasitology Research, 2018, 117, 169-175.	1.6	13
35	Giardia duodenalis in patients with diarrhea and various animals in northeastern China: prevalence and multilocus genetic characterization. Parasites and Vectors, 2022, 15, 165.	2.5	9
36	Multilocus sequence typing and population genetic structure of Cryptosporidium cuniculus in rabbits in Heilongjiang Province, China. Infection, Genetics and Evolution, 2018, 64, 249-253.	2.3	7

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37	Identification of Uncommon Cryptosporidiumviatorum (a Novel Subtype XVcA2G1c) and Cryptosporidium andersoni as Well as Common Giardia duodenalis Assemblages A and B in Humans in Myanmar. Frontiers in Cellular and Infection Microbiology, 2020, 10, 614053.	3.9	7
38	Occurrence and multilocus genotyping of Giardia duodenalis in pets and zoo animals in Shanghai, China. Journal of Infection in Developing Countries, 2017, 11, 479-486.	1.2	7
39	Multilocus Sequence Typing of Enterocytozoon bieneusi Isolates From Various Mammal and Bird Species and Assessment of Population Structure and Substructure. Frontiers in Microbiology, 2020, 11, 1406.	3.5	6
40	Prevalence and Genetic Characterization of Two Mitochondrial Gene Sequences of Strobilocercus Fasciolaris in the Livers of Brown Rats (Rattus norvegicus) in Heilongjiang Province in Northeastern China. Frontiers in Cellular and Infection Microbiology, 2020, 10, 588107.	3.9	2