Marianne Lebbad

List of Publications by Year in descending order

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218677 289244 1,766 38 26 40 citations g-index h-index papers 40 40 40 1702 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	From mouse to moose: Multilocus genotyping of Giardia isolates from various animal species. Veterinary Parasitology, 2010, 168, 231-239.	1.8	188
2	Large Outbreak of <i>Cryptosporidium hominis </i> Infection Transmitted through the Public Water Supply, Sweden. Emerging Infectious Diseases, 2014, 20, 581-589.	4.3	136
3	Multilocus Genotyping of Human Giardia Isolates Suggests Limited Zoonotic Transmission and Association between Assemblage B and Flatulence in Children. PLoS Neglected Tropical Diseases, 2011, 5, e1262.	3.0	118
4	Dominance of Giardia assemblage B in León, Nicaragua. Acta Tropica, 2008, 106, 44-53.	2.0	104
5	Increased Sampling Reveals Novel Lineages of Entamoeba: Consequences of Genetic Diversity and Host Specificity for Taxonomy and Molecular Detection. Protist, 2011, 162, 525-541.	1.5	103
6	High Applicability of a Novel Method for gp60-Based Subtyping of Cryptosporidium meleagridis. Journal of Clinical Microbiology, 2014, 52, 2311-2319.	3.9	74
7	Common Coinfections of Giardia intestinalis and Helicobacter pylori in Non-Symptomatic Ugandan Children. PLoS Neglected Tropical Diseases, 2012, 6, e1780.	3.0	71
8	Zoonotic transmission of Cryptosporidium meleagridis on an organic Swedish farm. International Journal for Parasitology, 2012, 42, 963-967.	3.1	61
9	The impact of genetic diversity in protozoa on molecular diagnostics. Trends in Parasitology, 2011, 27, 53-58.	3.3	57
10	Subtyping Novel Zoonotic Pathogen Cryptosporidium Chipmunk Genotype I. Journal of Clinical Microbiology, 2015, 53, 1648-1654.	3.9	57
11	Comparative genomic analyses of freshly isolated Giardia intestinalis assemblage A isolates. BMC Genomics, 2015, 16, 697.	2.8	55
12	An outbreak of cryptosporidiosis associated with exposure to swimming pool water. Scandinavian Journal of Infectious Diseases, 2005, 37, 354-360.	1.5	49
13	Allelic sequence heterozygosity in single Giardia parasites. BMC Microbiology, 2012, 12, 65.	3.3	47
14	A novel high-resolution multilocus sequence typing of Giardia intestinalis Assemblage A isolates reveals zoonotic transmission, clonal outbreaks and recombination. Infection, Genetics and Evolution, 2018, 60, 7-16.	2.3	42
15	PCR differentiation of Entamoeba histolytica and Entamoeba dispar from patients with amoeba infection initially diagnosed by microscopy. Scandinavian Journal of Infectious Diseases, 2005, 37, 680-685.	1.5	38
16	Overdiagnosis of Entamoeba histolytica and Entamoeba dispar in Nicaragua: A Microscopic, Triage Parasite Panel and PCR Study. Archives of Medical Research, 2006, 37, 529-534.	3.3	38
17	Detection of Giardia duodenalis Assemblages A and B in Human Feces by Simple, Assemblage-Specific PCR Assays. PLoS Neglected Tropical Diseases, 2012, 6, e1776.	3.0	38
18	Genetic characterisation of uninucleated cyst-producing Entamoeba spp. from ruminants. International Journal for Parasitology, 2010, 40, 775-778.	3.1	37

#	Article	IF	Citations
19	Differentiation of Blastocystis and parasitic archamoebids encountered in untreated wastewater samples by amplicon-based next-generation sequencing. Parasite Epidemiology and Control, 2020, 9, e00131.	1.8	37
20	On-Chip Imaging of Schistosoma haematobium Eggs in Urine for Diagnosis by Computer Vision. PLoS Neglected Tropical Diseases, 2013, 7, e2547.	3.0	36
21	High Diversity of Cryptosporidium Species and Subtypes Identified in Cryptosporidiosis Acquired in Sweden and Abroad. Pathogens, 2021, 10, 523.	2.8	34
22	Last of the Human Protists: The Phylogeny and Genetic Diversity of Iodamoeba. Molecular Biology and Evolution, 2012, 29, 39-42.	8.9	33
23	Unusual cryptosporidiosis cases in Swedish patients: extended molecular characterization of <i>Cryptosporidium viatorum </i> and <i>Cryptosporidium </i> chipmunk genotype I. Parasitology, 2013, 140, 1735-1740.	1.5	33
24	Web-Based Virtual Microscopy for Parasitology: A Novel Tool for Education and Quality Assurance. PLoS Neglected Tropical Diseases, 2008, 2, e315.	3.0	31
25	Identification and delineation of members of the Entamoeba complex by pyrosequencing. Molecular and Cellular Probes, 2010, 24, 403-406.	2.1	28
26	Low host-specific Enterocytozoon bieneusi genotype BEB6 is common in Swedish lambs. Veterinary Parasitology, 2014, 205, 371-374.	1.8	27
27	Possible zoonotic transmission of <i>Cryptosporidium felis</i> in a household. Infection Ecology and Epidemiology, 2015, 5, 28463.	0.8	26
28	Evaluation of a PCR Method for Detection of Entamoeba polecki, with an Overview of Its Molecular Epidemiology. Journal of Clinical Microbiology, 2018, 56, .	3.9	22
29	Population-based analyses of Giardia duodenalis is consistent with the clonal assemblage structure. Parasites and Vectors, 2012, 5, 168.	2.5	21
30	Improving the genotyping resolution of Cryptosporidium hominis subtype IbA10G2 using one step PCR-based amplicon sequencing. Infection, Genetics and Evolution, 2017, 55, 297-304.	2.3	20
31	Early outbreak detection by linking health advice line calls to water distribution areas retrospectively demonstrated in a large waterborne outbreak of cryptosporidiosis in Sweden. BMC Public Health, 2017, 17, 328.	2.9	19
32	Subtype distribution of zoonotic pathogen <i>Cryptosporidium felis</i> in humans and animals in several countries. Emerging Microbes and Infections, 2020, 9, 2446-2454.	6.5	19
33	Occurrence of Cryptosporidium spp. and Cystoisospora belli among adult patients with diarrhoea in Maputo, Mozambique. Heliyon, 2018, 4, e00769.	3.2	17
34	From mice to men: Three cases of human infection with Cryptosporidium ditrichi. Infection, Genetics and Evolution, 2020, 78, 104120.	2.3	17
35	Molecular characterization and epidemiological investigation of Cryptosporidium hominis IkA18G1 and C.Âhomini s monkey genotype IiA17, two unusual subtypes diagnosed in Swedish patients. Experimental Parasitology, 2018, 188, 50-57.	1.2	14
36	Cryptosporidium chipmunk genotype I – An emerging cause of human cryptosporidiosis in Sweden. Infection, Genetics and Evolution, 2021, 92, 104895.	2.3	9

#	Article	IF	CITATIONS
37	Comment on article by Eligio-Garcia et al. entitled "Frequency of Giardia intestinalis assemblages isolated from dogs and humans in a community from Culiacan, Sinaloa, Mexico using beta-giardin restriction gene― Veterinary Parasitology, 2008, 158, 159-160.	1.8	4
38	Real-time polymerase chain reaction followed by fast sequencing allows rapid genotyping of microbial pathogens. Scandinavian Journal of Infectious Diseases, 2011, 43, 95-99.	1.5	2