

Yvonne Qvarnstrom

List of Publications by Year in descending order

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#	ARTICLE	IF	CITATIONS
1	Multilocus Sequence Typing Tool for <i>Cyclospora cayetanensis</i> . Emerging Infectious Diseases, 2016, 22, 1464-1467.	4.3	38
2	Genotyping genetically heterogeneous <i>Cyclospora cayetanensis</i> infections to complement epidemiological case linkage. Parasitology, 2019, 146, 1275-1283.	1.5	36
3	Draft Genome Sequences from <i>Cyclospora cayetanensis</i> Oocysts Purified from a Human Stool Sample. Genome Announcements, 2015, 3, .	0.8	31
4	Mitochondrial Junction Region as Genotyping Marker for <i>Cyclospora cayetanensis</i> . Emerging Infectious Diseases, 2019, 25, 1314-1319.	4.3	28
5	A linear mitochondrial genome of <i>Cyclospora cayetanensis</i> (Eimeriidae, Eucoccidiorida, Coccidiasina,) Tj ETQq1 1 0.784314 rgBT /Ove coccidia. International Journal for Parasitology, 2015, 45, 361-365.	3.1	27
6	Evaluation of an ensemble-based distance statistic for clustering MLST datasets using epidemiologically defined clusters of cyclosporiasis. Epidemiology and Infection, 2020, 148, e172.	2.1	27
7	Molecular detection of <i>Cyclospora cayetanensis</i> in human stool specimens using UNEX-based DNA extraction and real-time PCR. Parasitology, 2018, 145, 865-870.	1.5	26
8	Comparative sequence analysis of <i>Cyclospora cayetanensis</i> apicoplast genomes originating from diverse geographical regions. Parasites and Vectors, 2016, 9, 611.	2.5	24
9	Molecular detection and genotyping of pathogenic protozoan parasites in raw and treated water samples from southwest Colombia. Parasites and Vectors, 2018, 11, 563.	2.5	23
10	Evaluation of Multilocus Sequence Typing of <i>Cyclospora cayetanensis</i> based on microsatellite markers. Parasite, 2019, 26, 3.	2.0	21
11	Purification of <i>Cyclospora cayetanensis</i> oocysts obtained from human stool specimens for whole genome sequencing. Gut Pathogens, 2018, 10, 45.	3.4	18
12	Molecular typing of <i>Cyclospora cayetanensis</i> in produce and clinical samples using targeted enrichment of complete mitochondrial genomes and next-generation sequencing. Parasites and Vectors, 2020, 13, 122.	2.5	18
13	Investigation of US <i>Cyclospora cayetanensis</i> outbreaks in 2019 and evaluation of an improved <i>Cyclospora</i> genotyping system against 2019 cyclosporiasis outbreak clusters. Epidemiology and Infection, 2021, 149, e214.	2.1	18
14	Genotyping <i>Cyclospora cayetanensis</i> From Multiple Outbreak Clusters With An Emphasis on a Cluster Linked to Bagged Salad Mixâ€™United States, 2020. Journal of Infectious Diseases, 2022, 225, 2176-2180.	4.0	16
15	Evaluation of library preparation methods for Illumina next generation sequencing of small amounts of DNA from foodborne parasites. Journal of Microbiological Methods, 2016, 130, 23-26.	1.6	15
16	Development of a workflow for identification of nuclear genotyping markers for <i>Cyclospora cayetanensis</i> . Parasite, 2020, 27, 24.	2.0	13