

Yvonne Qvarnstrom

List of Publications by Year in descending order

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#	ARTICLE	IF	CITATIONS
1	Genotyping <i>Cyclospora cayetanensis</i> From Multiple Outbreak Clusters With An Emphasis on a Cluster Linked to Bagged Salad Mix—United States, 2020. <i>Journal of Infectious Diseases</i> , 2022, 225, 2176-2180.	4.0	16
2	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. <i>Epidemiology and Infection</i> , 2021, 149, e214.	2.1	18
3	Evaluation of an ensemble-based distance statistic for clustering MLST datasets using epidemiologically defined clusters of cyclosporiasis. <i>Epidemiology and Infection</i> , 2020, 148, e172.	2.1	27
4	Molecular typing of <i>Cyclospora cayetanensis</i> in produce and clinical samples using targeted enrichment of complete mitochondrial genomes and next-generation sequencing. <i>Parasites and Vectors</i> , 2020, 13, 122.	2.5	18
5	Development of a workflow for identification of nuclear genotyping markers for <i>Cyclospora cayetanensis</i> . <i>Parasite</i> , 2020, 27, 24.	2.0	13
6	Evaluation of Multilocus Sequence Typing of <i>Cyclospora cayetanensis</i> based on microsatellite markers. <i>Parasite</i> , 2019, 26, 3.	2.0	21
7	Mitochondrial Junction Region as Genotyping Marker for <i>Cyclospora cayetanensis</i> . <i>Emerging Infectious Diseases</i> , 2019, 25, 1314-1319.	4.3	28
8	Genotyping genetically heterogeneous <i>Cyclospora cayetanensis</i> infections to complement epidemiological case linkage. <i>Parasitology</i> , 2019, 146, 1275-1283.	1.5	36
9	Molecular detection of <i>Cyclospora cayetanensis</i> in human stool specimens using UNEX-based DNA extraction and real-time PCR. <i>Parasitology</i> , 2018, 145, 865-870.	1.5	26
10	Molecular detection and genotyping of pathogenic protozoan parasites in raw and treated water samples from southwest Colombia. <i>Parasites and Vectors</i> , 2018, 11, 563.	2.5	23
11	Purification of <i>Cyclospora cayetanensis</i> oocysts obtained from human stool specimens for whole genome sequencing. <i>Gut Pathogens</i> , 2018, 10, 45.	3.4	18
12	Multilocus Sequence Typing Tool for <i>Cyclospora cayetanensis</i> . <i>Emerging Infectious Diseases</i> , 2016, 22, 1464-1467.	4.3	38
13	Comparative sequence analysis of <i>Cyclospora cayetanensis</i> apicoplast genomes originating from diverse geographical regions. <i>Parasites and Vectors</i> , 2016, 9, 611.	2.5	24
14	Evaluation of library preparation methods for Illumina next generation sequencing of small amounts of DNA from foodborne parasites. <i>Journal of Microbiological Methods</i> , 2016, 130, 23-26.	1.6	15
15	Draft Genome Sequences from <i>Cyclospora cayetanensis</i> Oocysts Purified from a Human Stool Sample. <i>Genome Announcements</i> , 2015, 3, .	0.8	31
16	A linear mitochondrial genome of <i>Cyclospora cayetanensis</i> (Eimeriidae, Eucoccidiorida, Coccidiasina), Tj ETQq0 0 0 rgBT /Overlock 10 Tf coccidia. <i>International Journal for Parasitology</i> , 2015, 45, 361-365.	3.1	27