

Claudia P Herrera

List of Publications by Year in descending order

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Version: 2024-02-01

50
papers

1,221
citations

394421

19
h-index

414414

32
g-index

51
all docs

51
docs citations

51
times ranked

1023
citing authors

#	ARTICLE	IF	CITATIONS
1	Identifying four <i>Trypanosoma cruzi</i> I isolate haplotypes from different geographic regions in Colombia. <i>Infection, Genetics and Evolution</i> , 2007, 7, 535-539.	2.3	127
2	Haplotype identification within <i>Trypanosoma cruzi</i> I in Colombian isolates from several reservoirs, vectors and humans. <i>Acta Tropica</i> , 2009, 110, 15-21.	2.0	108
3	Detailed ecological associations of triatomines revealed by metabarcoding and next-generation sequencing: implications for triatomine behavior and <i>Trypanosoma cruzi</i> transmission cycles. <i>Scientific Reports</i> , 2018, 8, 4140.	3.3	106
4	Genotype diversity of <i>Trypanosoma cruzi</i> in small rodents and <i>Triatoma sanguisuga</i> from a rural area in New Orleans, Louisiana. <i>Parasites and Vectors</i> , 2015, 8, 123.	2.5	58
5	Molecular identification and genotyping of <i>Trypanosoma cruzi</i> DNA in autochthonous Chagas disease patients from Texas, USA. <i>Infection, Genetics and Evolution</i> , 2017, 49, 151-156.	2.3	52
6	Genetic Variability and Phylogenetic Relationships within <i>Trypanosoma cruzi</i> I Isolated in Colombia Based on Miniexon Gene Sequences. <i>Journal of Parasitology Research</i> , 2009, 2009, 1-9.	1.2	48
7	Congenital Transmission of <i>Trypanosoma cruzi</i> in Argentina, Honduras, and Mexico: An Observational Prospective Study. <i>American Journal of Tropical Medicine and Hygiene</i> , 2018, 98, 478-485.	1.4	48
8	Estimating the current burden of Chagas disease in Mexico: A systematic review and meta-analysis of epidemiological surveys from 2006 to 2017. <i>PLoS Neglected Tropical Diseases</i> , 2019, 13, e0006859.	3.0	46
9	Interest and limitations of Spliced Leader Intergenic Region sequences for analyzing <i>Trypanosoma cruzi</i> I phylogenetic diversity in the Argentinean Chaco. <i>Infection, Genetics and Evolution</i> , 2011, 11, 300-307.	2.3	38
10	High prevalence of <i>Trypanosoma cruzi</i> infection in shelter dogs from southern Louisiana, USA. <i>Parasites and Vectors</i> , 2019, 12, 322.	2.5	36
11	Molecular Genotyping of <i>Trypanosoma cruzi</i> by Next-Generation Sequencing of the Mini-Exon Gene Reveals Infections With Multiple Parasite Discrete Typing Units in Chagasic Patients From Yucatan, Mexico. <i>Journal of Infectious Diseases</i> , 2019, 219, 1980-1988.	4.0	31
12	Deep sequencing reveals multiclonality and new discrete typing units of <i>Trypanosoma cruzi</i> in rodents from the southern United States. <i>Journal of Microbiology, Immunology and Infection</i> , 2020, 53, 622-633.	3.1	31
13	Interactions among <i>Triatoma sanguisuga</i> blood feeding sources, gut microbiota and <i>Trypanosoma cruzi</i> diversity in southern Louisiana. <i>Molecular Ecology</i> , 2020, 29, 3747-3761.	3.9	29
14	Chagas Disease Has Not Been Controlled in Ecuador. <i>PLoS ONE</i> , 2016, 11, e0158145.	2.5	27
15	Striking Divergence in <i>Toxoplasma ROP16</i> Nucleotide Sequences From Human and Meat Samples. <i>Journal of Infectious Diseases</i> , 2015, 211, 2006-2013.	4.0	26
16	A therapeutic preconceptional vaccine against Chagas disease: A novel indication that could reduce congenital transmission and accelerate vaccine development. <i>PLoS Neglected Tropical Diseases</i> , 2019, 13, e0006985.	3.0	26
17	Ten years of Chagas disease research: Looking back to achievements, looking ahead to challenges. <i>PLoS Neglected Tropical Diseases</i> , 2017, 11, e0005422.	3.0	24
18	Assessing <i>Trypanosoma cruzi</i> Parasite Diversity through Comparative Genomics: Implications for Disease Epidemiology and Diagnostics. <i>Pathogens</i> , 2021, 10, 212.	2.8	24

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19	Geographic Variations in Test Reactivity for the Serological Diagnosis of <i>Trypanosoma cruzi</i> Infection. <i>Journal of Clinical Microbiology</i> , 2021, 59, e0106221.	3.9	24
20	Toxoplasmosis in military personnel involved in jungle operations. <i>Acta Tropica</i> , 2012, 122, 46-51.	2.0	23
21	Phylogenetic Analysis of <i>Trypanosoma cruzi</i> from Pregnant Women and Newborns from Argentina, Honduras, and Mexico Suggests an Association of Parasite Haplotypes with Congenital Transmission of the Parasite. <i>Journal of Molecular Diagnostics</i> , 2019, 21, 1095-1105.	2.8	21
22	<i>Trypanosoma cruzi</i> diversity in naturally infected nonhuman primates in Louisiana assessed by deep sequencing of the mini-exon gene. <i>Transactions of the Royal Society of Tropical Medicine and Hygiene</i> , 2019, 113, 281-286.	1.8	21
23	Complex evolutionary pathways of the intergenic region of the mini-exon gene in <i>Trypanosoma cruzi</i> TcI: A possible ancient origin in the Gran Chaco and lack of strict genetic structuration. <i>Infection, Genetics and Evolution</i> , 2013, 16, 27-37.	2.3	19
24	An Improved Approach to <i>Trypanosoma cruzi</i> Molecular Genotyping by Next-Generation Sequencing of the Mini-exon Gene. <i>Methods in Molecular Biology</i> , 2019, 1955, 47-60.	0.9	18
25	The Case for the Development of a Chagas Disease Vaccine: Why? How? When?. <i>Tropical Medicine and Infectious Disease</i> , 2021, 6, 16.	2.3	17
26	Polymorphism and Selection Pressure of SARS-CoV-2 Vaccine and Diagnostic Antigens: Implications for Immune Evasion and Serologic Diagnostic Performance. <i>Pathogens</i> , 2020, 9, 584.	2.8	16
27	Short-course Benznidazole treatment to reduce <i>Trypanosoma cruzi</i> parasitic load in women of reproductive age (BETTY): a non-inferiority randomized controlled trial study protocol. <i>Reproductive Health</i> , 2020, 17, 128.	3.1	16
28	Safety and immunogenicity of a recombinant vaccine against <i>Trypanosoma cruzi</i> in Rhesus macaques. <i>Vaccine</i> , 2020, 38, 4584-4591.	3.8	16
29	Genetic diversity of <i>Trypanosoma cruzi</i> parasites infecting dogs in southern Louisiana sheds light on parasite transmission cycles and serological diagnostic performance. <i>PLoS Neglected Tropical Diseases</i> , 2020, 14, e0008932.	3.0	14
30	Molecular epidemiology of <i>Trypanosoma cruzi</i> and <i>Triatoma dimidiata</i> in coastal Ecuador. <i>Infection, Genetics and Evolution</i> , 2016, 41, 207-212.	2.3	13
31	Diversity and interactions among triatomine bugs, their blood feeding sources, gut microbiota and <i>Trypanosoma cruzi</i> in the Sierra Nevada de Santa Marta in Colombia. <i>Scientific Reports</i> , 2021, 11, 12306.	3.3	13
32	Seroprevalence of <i>Trypanosoma cruzi</i> Infection in Schoolchildren and in Pregnant Women from an Amazonian Region in Orellana Province, Ecuador. <i>American Journal of Tropical Medicine and Hygiene</i> , 2015, 93, 774-778.	1.4	12
33	Extent of polymorphism and selection pressure on the <i>Trypanosoma cruzi</i> vaccine candidate antigen Tc24. <i>Evolutionary Applications</i> , 2020, 13, 2663-2672.	3.1	11
34	In the heart of the city: <i>Trypanosoma cruzi</i> infection prevalence in rodents across New Orleans. <i>Parasites and Vectors</i> , 2020, 13, 577.	2.5	10
35	Raccoons As an Important Reservoir for <i>Trypanosoma cruzi</i> : A Prevalence Study from Two Metropolitan Areas in Louisiana. <i>Vector-Borne and Zoonotic Diseases</i> , 2020, 20, 535-540.	1.5	10
36	Shelter cats host infections with multiple <i>Trypanosoma cruzi</i> discrete typing units in southern Louisiana. <i>Veterinary Research</i> , 2021, 52, 53.	3.0	10

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37	Diversity of <i>Trypanosoma cruzi</i> parasites infecting <i>Triatoma dimidiata</i> in Central Veracruz, Mexico, and their One Health ecological interactions. <i>Infection, Genetics and Evolution</i> , 2021, 95, 105050.	2.3	10
38	Molecular ecology of <i>Triatoma dimidiata</i> in southern Belize reveals risk for human infection and the local differentiation of <i>Trypanosoma cruzi</i> parasites. <i>International Journal of Infectious Diseases</i> , 2021, 108, 320-329.	3.3	9
39	Validation of a Poisson-distributed limiting dilution assay (LDA) for a rapid and accurate resolution of multiclonal infections in natural <i>Trypanosoma cruzi</i> populations. <i>Journal of Microbiological Methods</i> , 2013, 92, 220-225.	1.6	8
40	Metabarcoding: A Powerful Yet Still Underestimated Approach for the Comprehensive Study of Vector-Borne Pathogen Transmission Cycles and Their Dynamics. , 0, , .		7
41	Genomic Signatures of SARS-CoV-2 Associated with Patient Mortality. <i>Viruses</i> , 2021, 13, 227.	3.3	7
42	Sequence of <i>Trypanosoma cruzi</i> reference strain SC43 nuclear genome and kinetoplast maxicircle confirms a strong genetic structure among closely related parasite discrete typing units. <i>Genome</i> , 2021, 64, 1-7.	2.0	6
43	Locally Transmitted <i>Trypanosoma cruzi</i> in a Domestic Llama (<i>Lama glama</i>) in a Rural Area of Greater New Orleans, Louisiana, USA. <i>Vector-Borne and Zoonotic Diseases</i> , 2021, 21, 762-768.	1.5	4
44	Active Transmission of <i>Trypanosoma cruzi</i> in Schoolchildren from the Amazon Region in Napo Province, Ecuador. <i>Acta Parasitologica</i> , 2021, 66, 1059-1062.	1.1	0
45	Title is missing!. , 2020, 14, e0008932.		0
46	Title is missing!. , 2020, 14, e0008932.		0
47	Title is missing!. , 2020, 14, e0008932.		0
48	Title is missing!. , 2020, 14, e0008932.		0
49	Title is missing!. , 2020, 14, e0008932.		0
50	Title is missing!. , 2020, 14, e0008932.		0