

# Patricio Diosque

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6776151/publications.pdf>

Version: 2024-02-01

11  
papers

798  
citations

1040056

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1281871

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times ranked

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citing authors

#	ARTICLE	IF	CITATIONS
1	International Study to Evaluate PCR Methods for Detection of <i>Trypanosoma cruzi</i> DNA in Blood Samples from Chagas Disease Patients. <i>PLoS Neglected Tropical Diseases</i> , 2011, 5, e931.	3.0	300
2	Genome-Scale Multilocus Microsatellite Typing of <i>Trypanosoma cruzi</i> Discrete Typing Unit I Reveals Phylogeographic Structure and Specific Genotypes Linked to Human Infection. <i>PLoS Pathogens</i> , 2009, 5, e1000410.	4.7	180
3	Multilocus Sequence Typing (MLST) for Lineage Assignment and High Resolution Diversity Studies in <i>Trypanosoma cruzi</i> . <i>PLoS Neglected Tropical Diseases</i> , 2011, 5, e1049.	3.0	94
4	Candidate targets for Multilocus Sequence Typing of <i>Trypanosoma cruzi</i> : Validation using parasite stocks from the Chaco Region and a set of reference strains. <i>Infection, Genetics and Evolution</i> , 2012, 12, 350-358.	2.3	54
5	<i>Trypanosoma cruzi</i> diversity in the Gran Chaco: Mixed infections and differential host distribution of TcV and TcVI. <i>Infection, Genetics and Evolution</i> , 2015, 29, 53-59.	2.3	54
6	Evolution of <i>Trypanosoma cruzi</i> : clarifying hybridisations, mitochondrial introgressions and phylogenetic relationships between major lineages. <i>Memorias Do Instituto Oswaldo Cruz</i> , 2015, 110, 403-413.	1.6	45
7	Optimized Multilocus Sequence Typing (MLST) Scheme for <i>Trypanosoma cruzi</i> . <i>PLoS Neglected Tropical Diseases</i> , 2014, 8, e3117.	3.0	31
8	Preponderant clonal evolution of <i>Trypanosoma cruzi</i> I from Argentinean Chaco revealed by Multilocus Sequence Typing (MLST). <i>Infection, Genetics and Evolution</i> , 2014, 27, 348-354.	2.3	15
9	Elucidating diversity in the class composition of the minicircle hypervariable region of <i>Trypanosoma cruzi</i> : New perspectives on typing and kDNA inheritance. <i>PLoS Neglected Tropical Diseases</i> , 2019, 13, e0007536.	3.0	13
10	Guide RNA Repertoires in the Main Lineages of <i>Trypanosoma cruzi</i> : High Diversity and Variable Redundancy Among Strains. <i>Frontiers in Cellular and Infection Microbiology</i> , 2021, 11, 663416.	3.9	7
11	Evidence of hybridization, mitochondrial introgression and biparental inheritance of the kDNA minicircles in <i>Trypanosoma cruzi</i> I. <i>PLoS Neglected Tropical Diseases</i> , 2020, 14, e0007770.	3.0	5