

Patrick Lypaczewski

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5053055/publications.pdf>

Version: 2024-02-01

12
papers

357
citations

933410

10
h-index

1199563

12
g-index

15
all docs

15
docs citations

15
times ranked

390
citing authors

#	ARTICLE	IF	CITATIONS
1	A second generation leishmanization vaccine with a markerless attenuated <i>Leishmania major</i> strain using CRISPR gene editing. <i>Nature Communications</i> , 2020, 11, 3461.	12.8	72
2	Optimized CRISPR-Cas9 Genome Editing for <i>Leishmania</i> and Its Use To Target a Multigene Family, Induce Chromosomal Translocation, and Study DNA Break Repair Mechanisms. <i>MSphere</i> , 2017, 2, .	2.9	66
3	A complete <i>Leishmania donovani</i> reference genome identifies novel genetic variations associated with virulence. <i>Scientific Reports</i> , 2018, 8, 16549.	3.3	41
4	Characterization of a new <i>Leishmania major</i> strain for use in a controlled human infection model. <i>Nature Communications</i> , 2021, 12, 215.	12.8	28
5	<i>Leishmania donovani</i> hybridisation and introgression in nature: a comparative genomic investigation. <i>Lancet Microbe</i> , The, 2021, 2, e250-e258.	7.3	26
6	Membrane Protein Complex ExbB ₄ -ExbD ₁ -TonB ₁ from <i>Escherichia coli</i> Demonstrates Conformational Plasticity. <i>Journal of Bacteriology</i> , 2015, 197, 1873-1885.	2.2	24
7	A review of the leishmanin skin test: A neglected test for a neglected disease. <i>PLoS Neglected Tropical Diseases</i> , 2021, 15, e0009531.	3.0	22
8	Centrin-deficient <i>Leishmania mexicana</i> confers protection against New World cutaneous leishmaniasis. <i>Npj Vaccines</i> , 2022, 7, 32.	6.0	19
9	Application of CRISPR/Cas9-Mediated Genome Editing in <i>Leishmania</i> . <i>Methods in Molecular Biology</i> , 2020, 2116, 199-224.	0.9	18
10	Development of a sandwich ELISA to detect <i>Leishmania</i> 40S ribosomal protein S12 antigen from blood samples of visceral leishmaniasis patients. <i>BMC Infectious Diseases</i> , 2018, 18, 500.	2.9	16
11	An intraspecies <i>Leishmania donovani</i> hybrid from the Indian subcontinent is associated with an atypical phenotype of cutaneous disease. <i>IScience</i> , 2022, 25, 103802.	4.1	12
12	Evidence that a naturally occurring single nucleotide polymorphism in the RagC gene of <i>Leishmania donovani</i> contributes to reduced virulence. <i>PLoS Neglected Tropical Diseases</i> , 2021, 15, e0009079.	3.0	11