Patrick Lypaczewski

List of Publications by Year in descending order

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