

Global distribution maps of the leishmaniases

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Mapping the zoonotic niche of Ebola virus disease in Africa. <i>ELife</i> , 2014, 3, e04395.	2.8	328
2	Measuring progress in global health. <i>Transactions of the Royal Society of Tropical Medicine and Hygiene</i> , 2014, 108, 521-522.	0.7	0
3	Benefit of Insecticide-Treated Nets, Curtains and Screening on Vector Borne Diseases, Excluding Malaria: A Systematic Review and Meta-analysis. <i>PLoS Neglected Tropical Diseases</i> , 2014, 8, e3228.	1.3	60
4	Global database of leishmaniasis occurrence locations, 1960–2012. <i>Scientific Data</i> , 2014, 1, 140036.	2.4	43
5	A National Assessment of the Epidemiology of Severe Fever with Thrombocytopenia Syndrome, China. <i>Scientific Reports</i> , 2015, 5, 9679.	1.6	102
6	Tracking the distribution and impacts of diseases with biological records and distribution modelling. <i>Biological Journal of the Linnean Society</i> , 2015, 115, 664-677.	0.7	36
7	The global distribution of the arbovirus vectors <i>Aedes aegypti</i> and <i>Ae. albopictus</i> . <i>ELife</i> , 2015, 4, e08347.	2.8	1,428
8	Integrating Data and Resources on Neglected Tropical Diseases for Better Planning: The NTD Mapping Tool (NTDmap.org). <i>PLoS Neglected Tropical Diseases</i> , 2015, 9, e0003400.	1.3	13
9	Sources of spatial animal and human health data: Casting the net wide to deal more effectively with increasingly complex disease problems. <i>Spatial and Spatio-temporal Epidemiology</i> , 2015, 13, 15-29.	0.9	25
10	Leishmaniasis in the Americas. <i>Neglected Tropical Diseases</i> , 2015, , 113-128.	0.4	1
11	The many projected futures of dengue. <i>Nature Reviews Microbiology</i> , 2015, 13, 230-239.	13.6	145
12	Characterization of the Biosynthetic Gene Cluster for Benzoxazole Antibiotics A33853 Reveals Unusual Assembly Logic. <i>Chemistry and Biology</i> , 2015, 22, 1313-1324.	6.2	48
13	The global distribution of Crimean-Congo hemorrhagic fever. <i>Transactions of the Royal Society of Tropical Medicine and Hygiene</i> , 2015, 109, 503-513.	0.7	193
14	Visceral leishmaniasis research: operational focus needed. <i>The Lancet Global Health</i> , 2015, 3, e194.	2.9	1
15	Combination of In Silico Methods in the Search for Potential CD4+ and CD8+ T Cell Epitopes in the Proteome of <i>Leishmania braziliensis</i> . <i>Frontiers in Immunology</i> , 2016, 7, 327.	2.2	47
16	Global distribution and environmental suitability for chikungunya virus, 1952 to 2015. <i>Eurosurveillance</i> , 2016, 21, .	3.9	141
17	Mapping global environmental suitability for Zika virus. <i>ELife</i> , 2016, 5, .	2.8	299
18	Spatial Modelling Tools to Integrate Public Health and Environmental Science, Illustrated with Infectious Cryptosporidiosis. <i>International Journal of Environmental Research and Public Health</i> , 2016, 13, 186.	1.2	12

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19	Genetic Diversity and Population Structure of <i>Leishmania infantum</i> from Southeastern France: Evaluation Using Multi-Locus Microsatellite Typing. <i>PLoS Neglected Tropical Diseases</i> , 2016, 10, e0004303.	1.3	10
20	Estimating Geographical Variation in the Risk of Zoonotic <i>Plasmodium knowlesi</i> Infection in Countries Eliminating Malaria. <i>PLoS Neglected Tropical Diseases</i> , 2016, 10, e0004915.	1.3	76
21	<i>Plasmodium knowlesi</i> transmission: integrating quantitative approaches from epidemiology and ecology to understand malaria as a zoonosis. <i>Parasitology</i> , 2016, 143, 389-400.	0.7	42
22	Risk Factors Associated with Human Visceral Leishmaniasis in an Urban Area of Bahia, Brazil. <i>Vector-Borne and Zoonotic Diseases</i> , 2016, 16, 368-376.	0.6	10
23	Trends in the Mechanistic and Dynamic Modeling of Infectious Diseases. <i>Current Epidemiology Reports</i> , 2016, 3, 212-222.	1.1	27
24	Global burden of cutaneous leishmaniasis. <i>Lancet Infectious Diseases</i> , The, 2016, 16, 1004-1005.	4.6	10
25	Tilting the balance between RNA interference and replication eradicates <i>Leishmania</i> RNA virus 1 and mitigates the inflammatory response. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 11998-12005.	3.3	46
26	Defining the targets of antiparasitic compounds. <i>Drug Discovery Today</i> , 2016, 21, 725-739.	3.2	25
27	DNA sequence analysis suggests that <i>cytb-nd1</i> PCR-RFLP may not be applicable to sandfly species identification throughout the Mediterranean region. <i>Parasitology Research</i> , 2016, 115, 1287-1295.	0.6	2
28	Estimations of cutaneous leishmaniasis burden: a constant challenge. <i>Lancet Infectious Diseases</i> , The, 2016, 16, 515-516.	4.6	2
29	Antiviral screening identifies adenosine analogs targeting the endogenous dsRNA <i>Leishmania</i> RNA virus 1 (LRV1) pathogenicity factor. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, E811-E819.	3.3	36
30	Continual renewal and replication of persistent <i>Leishmania major</i> parasites in concomitantly immune hosts. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, E801-E810.	3.3	98
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32	Carbonic anhydrases from <i>Trypanosoma</i> and <i>Leishmania</i> as anti-protozoan drug targets. <i>Bioorganic and Medicinal Chemistry</i> , 2017, 25, 1543-1555.	1.4	52
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34	Global burden of cutaneous leishmaniasis. <i>Lancet Infectious Diseases</i> , The, 2017, 17, 264.	4.6	14
35	Biosynthesized colloidal silver and gold nanoparticles as emerging leishmanicidal agents: an insight. <i>Nanomedicine</i> , 2017, 12, 2807-2819.	1.7	45
36	Spatial and temporal distribution of American cutaneous leishmaniasis in Acre state, Brazil. <i>Infectious Diseases of Poverty</i> , 2017, 6, 99.	1.5	27

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37	Seroprevalence of sandfly fever virus infection in military personnel on the western border of Iran. <i>Journal of Infection and Public Health</i> , 2017, 10, 59-63.	1.9	14
38	Visceral Leishmaniasis and Natural Infection Rates of <i>Leishmania</i> in <i>Lutzomyia longipalpis</i> in Latin America. , 0, , .		2
39	Niche Modeling of Dengue Fever Using Remotely Sensed Environmental Factors and Boosted Regression Trees. <i>Remote Sensing</i> , 2017, 9, 328.	1.8	30
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41	In Silico Identification of B-Cell Epitopes of <i>Leishmania infantum</i> Recombinant Histone Shared with Human Sera Stably Living in Area Where <i>Leishmania</i> Species Does Perpetuate. <i>Journal of Next Generation Sequencing & Applications</i> , 2017, 04, .	0.3	0
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45	Mapping the epidemic changes and risks of hemorrhagic fever with renal syndrome in Shaanxi Province, China, 2005–2016. <i>Scientific Reports</i> , 2018, 8, 749.	1.6	35
46	A Brief Introduction to Leishmaniasis Epidemiology. , 2018, , 1-13.		8
47	<i>Leishmania mexicana</i> can utilize amino acids as major carbon sources in macrophages but not in animal models. <i>Molecular Microbiology</i> , 2018, 108, 143-158.	1.2	31
48	New chalcone compound as a promising antileishmanial drug for an old neglected disease: Biological evaluation using radiolabelled biodistribution. <i>Journal of Global Antimicrobial Resistance</i> , 2018, 13, 139-142.	0.9	9
49	Diagnostic approach to tropical skin infections. <i>Medicine</i> , 2018, 46, 10-15.	0.2	2
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52	Coinfection With <i>Trypanosoma brucei</i> Confers Protection Against Cutaneous Leishmaniasis. <i>Frontiers in Immunology</i> , 2018, 9, 2855.	2.2	4
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56	Visceral leishmaniasis in a Brazilian endemic area: an overview of occurrence, HIV coinfection and lethality. <i>Revista Do Instituto De Medicina Tropical De Sao Paulo</i> , 2018, 60, e12.	0.5	20
57	Histopathological and immunohistochemical characterisation of hepatic granulomas in <i>Leishmania donovani</i> -infected BALB/c mice: a time-course study. <i>Parasites and Vectors</i> , 2018, 11, 73.	1.0	31
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59	Bioinformatics in <i>Leishmania</i> Drug Design. , 2018, , 297-317.		0
60	Pentamidine inhibits the growth of <i>Sporothrix schenckii</i> complex and exhibits synergism with antifungal agents. <i>Future Microbiology</i> , 2018, 13, 1129-1140.	1.0	16
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75	Comments on letter to the editor by Faniyan <i>et al</i> . in response to Imported leishmaniasis in Sweden 1993–2016. <i>Epidemiology and Infection</i> , 2019, 147, e47.	1.0	1
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111	Effects of Visceralising Leishmania on the Spleen, Liver, and Bone Marrow: A Pathophysiological Perspective. Microorganisms, 2021, 9, 759.	1.6	18
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