

Katrin Kuhls

List of Publications by Year in Descending Order

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Version: 2024-04-28

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

50
papers

2,649
citations

25
h-index

51
g-index

52
ext. papers

2,969
ext. citations

4.9
avg, IF

4.52
L-index

#	Paper	IF	Citations
50	Microsatellite based molecular epidemiology of <i>Leishmania infantum</i> from re-emerging foci of visceral leishmaniasis in Armenia and pilot risk assessment by ecological niche modeling. <i>PLoS Neglected Tropical Diseases</i> , 2021 , 15, e0009288	4.8	1
49	Combined climate and regional mosquito habitat model based on machine learning. <i>Ecological Modelling</i> , 2021 , 452, 109594	3	2
48	Integrative Approach to Grassi, 1908: First Record in Vienna with New Morphological and Molecular Insights. <i>Pathogens</i> , 2020 , 9,	4.5	5
47	History of the E.I. Martsinovskiy Institute of Medical Parasitology and Tropical Medicine: research on malaria and leishmaniasis. <i>Historia, Ciencias, Saude - Manguinhos</i> , 2020 , 27, 1097-1124	0.2	1
46	A novel multilocus sequence typing scheme identifying genetic diversity amongst <i>Leishmania donovani</i> isolates from a genetically homogeneous population in the Indian subcontinent. <i>International Journal for Parasitology</i> , 2019 , 49, 555-567	4.3	6
45	Phylogenetic Studies. <i>Methods in Molecular Biology</i> , 2019 , 1971, 9-68	1.4	2
44	Re-Emerging foci of visceral leishmaniasis in Armenia - first molecular diagnosis of clinical samples. <i>Parasitology</i> , 2019 , 146, 857-864	2.7	1
43	<i>Leishmania</i> infections: Molecular targets and diagnosis. <i>Molecular Aspects of Medicine</i> , 2017 , 57, 1-29	16.7	142
42	A pilot study on fingerprinting <i>Leishmania</i> species from the Old World using Fourier transform infrared spectroscopy. <i>Analytical and Bioanalytical Chemistry</i> , 2017 , 409, 6907-6923	4.4	12
41	Epidemiological analysis of <i>Leishmania tropica</i> strains and giemsa-stained smears from Syrian and Turkish leishmaniasis patients using multilocus microsatellite typing (MLMT). <i>PLoS Neglected Tropical Diseases</i> , 2017 , 11, e0005538	4.8	6
40	Spatiotemporal and molecular epidemiology of cutaneous leishmaniasis in Libya. <i>PLoS Neglected Tropical Diseases</i> , 2017 , 11, e0005873	4.8	9
39	Phylogenetic structure of <i>Leishmania tropica</i> in the new endemic focus Birjand in East Iran in comparison to other Iranian endemic regions. <i>Acta Tropica</i> , 2016 , 158, 68-76	3.2	19
38	A Historical Overview of the Classification, Evolution, and Dispersion of <i>Leishmania</i> Parasites and Sandflies. <i>PLoS Neglected Tropical Diseases</i> , 2016 , 10, e0004349	4.8	403
37	Heterogeneity of the internal transcribed spacer region in <i>Leishmania tropica</i> isolates from southern Iran. <i>Experimental Parasitology</i> , 2014 , 144, 44-51	2.1	22
36	Genetic diversity evaluation on Portuguese <i>Leishmania infantum</i> strains by multilocus microsatellite typing. <i>Infection, Genetics and Evolution</i> , 2014 , 26, 20-31	4.5	11
35	<i>Leishmania donovani</i> populations in Eastern Sudan: temporal structuring and a link between human and canine transmission. <i>Parasites and Vectors</i> , 2014 , 7, 496	4	12
34	Population structure and evidence for both clonality and recombination among Brazilian strains of the subgenus <i>Leishmania</i> (<i>Viannia</i>). <i>PLoS Neglected Tropical Diseases</i> , 2013 , 7, e2490	4.8	28

33	Genetic typing reveals monomorphism between antimony sensitive and resistant <i>Leishmania donovani</i> isolates from visceral leishmaniasis or post kala-azar dermal leishmaniasis cases in India. <i>Parasitology Research</i> , 2012 , 111, 1559-68	2.4	9
32	Multilocus microsatellite typing (MLMT) of strains from Turkey and Cyprus reveals a novel monophyletic <i>L. donovani</i> sensu lato group. <i>PLoS Neglected Tropical Diseases</i> , 2012 , 6, e1507	4.8	40
31	Molecular approaches for a better understanding of the epidemiology and population genetics of <i>Leishmania</i> . <i>Parasitology</i> , 2011 , 138, 405-25	2.7	117
30	Multilocus microsatellite typing shows three different genetic clusters of <i>Leishmania major</i> in Iran. <i>Microbes and Infection</i> , 2011 , 13, 937-42	9.3	24
29	Comparative microsatellite typing of new world <i>Leishmania infantum</i> reveals low heterogeneity among populations and its recent old world origin. <i>PLoS Neglected Tropical Diseases</i> , 2011 , 5, e1155	4.8	128
28	Multilocus genotyping reveals a polyphyletic pattern among naturally antimony-resistant <i>Leishmania braziliensis</i> isolates from Peru. <i>Infection, Genetics and Evolution</i> , 2011 , 11, 1873-80	4.5	14
27	Multilocus microsatellite typing revealed high genetic variability of <i>Leishmania donovani</i> strains isolated during and after a Kala-azar epidemic in Libo Kemkem district, northwest Ethiopia. <i>Microbes and Infection</i> , 2011 , 13, 595-601	9.3	11
26	Guns, germs and dogs: On the origin of <i>Leishmania chagasi</i> . <i>Infection, Genetics and Evolution</i> , 2011 , 11, 1091-5	4.5	46
25	Disseminated cutaneous leishmaniasis resembling post-kala-azar dermal leishmaniasis caused by <i>Leishmania donovani</i> in three patients co-infected with visceral leishmaniasis and human immunodeficiency virus/acquired immunodeficiency syndrome in Ethiopia. <i>American Journal of Tropical Medicine and Hygiene</i> , 2011 , 84, 906-12	3.2	17
24	Atypical lesions as a sign of cutaneous dissemination of visceral leishmaniasis in a human immunodeficiency virus-positive patient simultaneously infected by two viscerotropic <i>Leishmania</i> species. <i>American Journal of Tropical Medicine and Hygiene</i> , 2011 , 85, 55-9	3.2	15
23	A clinical isolate of <i>Leishmania donovani</i> with ITS1 sequence polymorphism as a cause of para-kala-azar dermal leishmaniasis in an Ethiopian human immunodeficiency virus-positive patient on highly active antiretroviral therapy. <i>British Journal of Dermatology</i> , 2010 , 163, 870-4	4	13
22	Inference of population structure of <i>Leishmania donovani</i> strains isolated from different Ethiopian visceral leishmaniasis endemic areas. <i>PLoS Neglected Tropical Diseases</i> , 2010 , 4, e889	4.8	60
21	Identification of the agent causing visceral leishmaniasis in Uzbeki and Tajiki foci by analysing parasite DNA extracted from patients Giemsa-stained tissue preparations. <i>Parasitology</i> , 2009 , 136, 981-6	2.7	14
20	Development of a multilocus microsatellite typing approach for discriminating strains of <i>Leishmania</i> (<i>Viannia</i>) species. <i>Journal of Clinical Microbiology</i> , 2009 , 47, 2818-25	9.7	43
19	Population structure of Tunisian <i>Leishmania infantum</i> and evidence for the existence of hybrids and gene flow between genetically different populations. <i>International Journal for Parasitology</i> , 2009 , 39, 801-11	4.3	60
18	PCR diagnosis of visceral leishmaniasis in an endemic region, Mymensingh district, Bangladesh. <i>Tropical Medicine and International Health</i> , 2009 , 14, 499-503	2.3	16
17	Population genetics of <i>Leishmania infantum</i> in Israel and the Palestinian Authority through microsatellite analysis. <i>Microbes and Infection</i> , 2009 , 11, 484-92	9.3	23
16	Multilocus microsatellite typing (MLMT) reveals genetic homogeneity of <i>Leishmania donovani</i> strains in the Indian subcontinent. <i>Infection, Genetics and Evolution</i> , 2009 , 9, 24-31	4.5	72

15	The paraphyletic composition of <i>Leishmania donovani</i> zymodeme MON-37 revealed by multilocus microsatellite typing. <i>Microbes and Infection</i> , 2009 , 11, 707-15	9.3	39
14	Identification of geographically distributed sub-populations of <i>Leishmania (Leishmania) major</i> by microsatellite analysis. <i>BMC Evolutionary Biology</i> , 2008 , 8, 183	3	56
13	Genetic polymorphism of Algerian <i>Leishmania infantum</i> strains revealed by multilocus microsatellite analysis. <i>Microbes and Infection</i> , 2008 , 10, 1309-15	9.3	43
12	Differentiation and gene flow among European populations of <i>Leishmania infantum</i> MON-1. <i>PLoS Neglected Tropical Diseases</i> , 2008 , 2, e261	4.8	68
11	Epidemiological dynamics of antimonial resistance in <i>Leishmania donovani</i> : genotyping reveals a polyclonal population structure among naturally-resistant clinical isolates from Nepal. <i>Infection, Genetics and Evolution</i> , 2007 , 7, 206-12	4.5	44
10	Multilocus microsatellite typing (MLMT) reveals genetically isolated populations between and within the main endemic regions of visceral leishmaniasis. <i>Microbes and Infection</i> , 2007 , 9, 334-43	9.3	133
9	Evolutionary and geographical history of the <i>Leishmania donovani</i> complex with a revision of current taxonomy. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007 , 104, 9375-80	11.5	309
8	Multilocus microsatellite typing as a new tool for discrimination of <i>Leishmania infantum</i> MON-1 strains. <i>Journal of Clinical Microbiology</i> , 2006 , 44, 495-503	9.7	84
7	<i>Leishmania major</i> : genetic heterogeneity of Iranian isolates by single-strand conformation polymorphism and sequence analysis of ribosomal DNA internal transcribed spacer. <i>Acta Tropica</i> , 2006 , 98, 52-8	3.2	56
6	Comparison of molecular markers for strain typing of <i>Leishmania infantum</i> . <i>Infection, Genetics and Evolution</i> , 2006 , 6, 440-6	4.5	65
5	Analysis of ribosomal DNA internal transcribed spacer sequences of the <i>Leishmania donovani</i> complex. <i>Microbes and Infection</i> , 2005 , 7, 1224-34	9.3	101
4	Revision of <i>Trichoderma</i> sect. <i>Longibrachiatum</i> including related teleomorphs based on analysis of ribosomal DNA internal transcribed spacer sequences. <i>Mycologia</i> , 1997 , 89, 442-460	2.4	100
3	Biogeography and phenotypic variation in <i>Trichoderma</i> sect. <i>Longibrachiatum</i> and associated <i>Hypocrea</i> species. <i>Mycological Research</i> , 1997 , 101, 449-459		87
2	Cellulase formation by species of <i>Trichoderma</i> sect. <i>Longibrachiatum</i> and of <i>Hypocrea</i> spp. with anamorphs referable to <i>Trichoderma</i> sect. <i>Longibrachiatum</i> . <i>Fungal Genetics and Biology</i> , 1996 , 20, 105-149		24
1	PCR-fingerprinting used for comparison of ex type strains of <i>Trichoderma</i> species deposited in different culture collections. <i>Microbiological Research</i> , 1995 , 150, 363-71	5.3	36