

Anne-Laure Bañuls

List of Publications by Year in descending order

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120
papers

5,962
citations

101543

36
h-index

79698

73
g-index

121
all docs

121
docs citations

121
times ranked

7575
citing authors

#	ARTICLE	IF	CITATIONS
1	Biodiversity and vector-borne diseases: Host dilution and vector amplification occur simultaneously for Amazonian leishmaniasis. <i>Molecular Ecology</i> , 2023, 32, 1817-1831.	3.9	18
2	Epidemiology and prevalence of extended-spectrum β -lactamase- and carbapenemase-producing Enterobacteriaceae in humans, animals and the environment in West and Central Africa. <i>International Journal of Antimicrobial Agents</i> , 2021, 57, 106203.	2.5	31
3	Lower <i>mgpB</i> diversity in macrolide-resistant <i>Mycoplasma genitalium</i> infecting men visiting two sexually transmitted infection clinics in Montpellier, France. <i>Journal of Antimicrobial Chemotherapy</i> , 2021, 76, 43-47.	3.0	13
4	Resistance to Second-Line Anti-TB Drugs in Cambodia: A Phenotypic and Genetic Study. <i>Infection and Drug Resistance</i> , 2021, Volume 14, 1089-1104.	2.7	6
5	Resistance mechanisms and genetic relatedness among carbapenem-resistant <i>Pseudomonas aeruginosa</i> isolates from three major hospitals in Hanoi, Vietnam (2011-15). <i>JAC-Antimicrobial Resistance</i> , 2021, 3, dlab103.	2.1	4
6	Expansion of KPC-producing Enterobacterales in four large hospitals in Hanoi, Vietnam. <i>Journal of Global Antimicrobial Resistance</i> , 2021, 27, 200-211.	2.2	12
7	Incubation Period for Neuroinvasive Toscana Virus Infections. <i>Emerging Infectious Diseases</i> , 2021, 27, 3147-3150.	4.3	8
8	Taxonomical insights and ecology of sandfly (Diptera, Psychodidae) species in six provinces of Northern Vietnam. <i>Parasite</i> , 2021, 28, 85.	2.0	4
9	Diversity and Ecology of Sand Flies (Diptera: Psychodidae), Potential Vectors of <i>Leishmania</i> in the Quang Ninh Province, Vietnam. <i>Journal of Medical Entomology</i> , 2020, 57, 259-265.	1.8	4
10	Comparative Study of Promastigote- and Amastigote-Initiated Infection of <i>Leishmania infantum</i> (Kinetoplastida: Trypanosomatidae) in <i>Phlebotomus perniciosus</i> (Diptera: Psychodidae) Conducted in Different Biosafety Level Laboratories. <i>Journal of Medical Entomology</i> , 2020, 57, 601-607.	1.8	6
11	Evaluation of the GenoType NTM-DR assay performance for the identification and molecular detection of antibiotic resistance in <i>Mycobacterium abscessus</i> complex. <i>PLoS ONE</i> , 2020, 15, e0239146.	2.5	11
12	Altitude and hillside orientation shapes the population structure of the <i>Leishmania infantum</i> vector <i>Phlebotomus ariasi</i> . <i>Scientific Reports</i> , 2020, 10, 14443.	3.3	3
13	Delamanid Resistance: Update and Clinical Management. <i>Clinical Infectious Diseases</i> , 2020, 71, 3252-3259.	5.8	30
14	Evaluation of Loopamp Assay for the Diagnosis of Pulmonary Tuberculosis in Cambodia. <i>BioMed Research International</i> , 2020, 2020, 1-7.	1.9	2
15	Broader Geographical Distribution of Toscana Virus in the Mediterranean Region Suggests the Existence of Larger Varieties of Sand Fly Vectors. <i>Microorganisms</i> , 2020, 8, 114.	3.6	22
16	Spread of NDM-5 and OXA-181 Carbapenemase-Producing <i>Escherichia coli</i> in Chad. <i>Antimicrobial Agents and Chemotherapy</i> , 2019, 63, .	3.2	12
17	A single amino acid substitution (H451Y) in <i>Leishmania</i> calcium-dependent kinase SCAMK confers high tolerance and resistance to antimony. <i>Journal of Antimicrobial Chemotherapy</i> , 2019, 74, 3231-3239.	3.0	7
18	First insights into the genetic characteristics and drug resistance of <i>Mycobacterium tuberculosis</i> population collected during the first national tuberculosis prevalence survey of Lao PDR (2010-2011). <i>BMC Infectious Diseases</i> , 2019, 19, 851.	2.9	3

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19	Fecal carriage of extended-spectrum β -lactamase-producing Enterobacteriaceae in hospital and community settings in Chad. <i>Antimicrobial Resistance and Infection Control</i> , 2019, 8, 169.	4.1	34
20	A first insight into genetic diversity of <i>Mycobacterium bovis</i> isolated from extrapulmonary tuberculosis patients in South Tunisia assessed by spoligotyping and MIRU VNTR. <i>PLoS Neglected Tropical Diseases</i> , 2019, 13, e0007707.	3.0	7
21	Molecular Diagnosis of Drug-Resistant Tuberculosis; A Literature Review. <i>Frontiers in Microbiology</i> , 2019, 10, 794.	3.5	80
22	High prevalence and characterization of extended-spectrum β -lactamase producing Enterobacteriaceae in Chadian hospitals. <i>BMC Infectious Diseases</i> , 2019, 19, 205.	2.9	38
23	Bedaquiline Resistance: Its Emergence, Mechanism, and Prevention. <i>Clinical Infectious Diseases</i> , 2018, 66, 1625-1630.	5.8	131
24	High-throughput sequencing of kDNA amplicons for the analysis of Leishmanian minicircles and identification of Neotropical species. <i>Parasitology</i> , 2018, 145, 585-594.	1.5	23
25	Insights into the processes that drive the evolution of drug resistance in <i>Mycobacterium tuberculosis</i> . <i>Evolutionary Applications</i> , 2018, 11, 1498-1511.	3.1	51
26	<i>Leishmania major</i> and <i>Trypanosoma lewisi</i> infection in invasive and native rodents in Senegal. <i>PLoS Neglected Tropical Diseases</i> , 2018, 12, e0006615.	3.0	10
27	Vector soup: high-throughput identification of Neotropical phlebotomine sand flies using metabarcoding. <i>Molecular Ecology Resources</i> , 2017, 17, 172-182.	4.8	31
28	Evaluation of short mitochondrial metabarcodes for the identification of Amazonian mammals. <i>Methods in Ecology and Evolution</i> , 2017, 8, 1276-1283.	5.2	33
29	Quadruple-first line drug resistance in <i>Mycobacterium tuberculosis</i> in Vietnam: What can we learn from genes?. <i>Infection, Genetics and Evolution</i> , 2017, 50, 55-61.	2.3	7
30	scp <i>2</i> screening: Disease vectors as vertebrate samplers. <i>Molecular Ecology</i> , 2017, 26, 6478-6486.	3.9	57
31	Reproduction in <i>Leishmania</i> : A focus on genetic exchange. <i>Infection, Genetics and Evolution</i> , 2017, 50, 128-132.	2.3	17
32	<i>Phlebotomus (Paraphlebotomus) chabaudi</i> and <i>Phlebotomus riouxi</i> : closely related species or synonyms?. <i>Parasite</i> , 2017, 24, 47.	2.0	7
33	Molecular analysis of pyrazinamide resistance in <i>Mycobacterium tuberculosis</i> in Vietnam highlights the high rate of pyrazinamide resistance-associated mutations in clinical isolates. <i>Emerging Microbes and Infections</i> , 2017, 6, 1-7.	6.5	17
34	Emergence and spread of antibiotic resistance in West Africa: contributing factors and threat assessment. <i>Medecine Et Sante Tropicales</i> , 2017, 27, 147-154.	0.3	53
35	Transmission of <i>Leishmania infantum</i> in the Canine Leishmaniasis Focus of Mont-Rolland, Senegal: Ecological, Parasitological and Molecular Evidence for a Possible Role of <i>Sergentomyia</i> Sand Flies. <i>PLoS Neglected Tropical Diseases</i> , 2016, 10, e0004940.	3.0	37
36	High Nasal Carriage Rate of <i>Staphylococcus aureus</i> Containing Panton-Valentine leukocidin- and EDIN-Encoding Genes in Community and Hospital Settings in Burkina Faso. <i>Frontiers in Microbiology</i> , 2016, 7, 1406.	3.5	14

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37	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.	3.0	10
38	Ecology and morphological variations in wings of <i>Phlebotomus ariasi</i> (Diptera: Psychodidae) in the region of Roquedur (Gard, France): a geometric morphometrics approach. <i>Parasites and Vectors</i> , 2016, 9, 578.	2.5	26
39	<i>Mycobacterium tuberculosis</i> lineages and anti-tuberculosis drug resistance in reference hospitals across Viet Nam. <i>BMC Microbiology</i> , 2016, 16, 167.	3.3	33
40	Evaluation of the SLOMYCO Sensititre® panel for testing the antimicrobial susceptibility of <i>Mycobacterium marinum</i> isolates. <i>Annals of Clinical Microbiology and Antimicrobials</i> , 2016, 15, 30.	3.8	15
41	Study of <i>Leishmania</i> pathogenesis in mice: experimental considerations. <i>Parasites and Vectors</i> , 2016, 9, 144.	2.5	110
42	Complete mitochondrial genome of <i>Lutzomyia</i> (<i>Nyssomyia</i>) <i>umbratilis</i> (Diptera: Tj ETQq0 0 0 rgBT /Overlock 10 Tff Mapping, Sequencing, and Analysis, 2016, 27, 4219-4221.	0.7	12
43	First insights into the genetic diversity and origin of <i>Leishmania infantum</i> in Mont Rolland (ThiÃ's Tj ETQq1 1 0.784314 rgBT /Overlock 1.9	1.9	7
44	Population structure of colonizing and invasive <i>Staphylococcus aureus</i> strains in northern Vietnam. <i>Journal of Medical Microbiology</i> , 2016, 65, 298-305.	1.8	13
45	Seasonal Dynamics of Phlebotomine Sand Fly Species Proven Vectors of Mediterranean Leishmaniasis Caused by <i>Leishmania infantum</i> . <i>PLoS Neglected Tropical Diseases</i> , 2016, 10, e0004458.	3.0	152
46	Sampling strategies for phlebotomine sand flies (Diptera: Psychodidae) in Europe. <i>Bulletin of Entomological Research</i> , 2015, 105, 664-678.	1.0	52
47	Multilocus microsatellite typing of <i>Leishmania</i> and clinical applications: a review. <i>Parasite</i> , 2015, 22, 16.	2.0	9
48	Multilocus microsatellite typing of <i>Leishmania infantum</i> isolates in monitored <i>Leishmania</i> /HIV coinfecting patients. <i>Parasites and Vectors</i> , 2015, 8, 386.	2.5	1
49	Ecology and spatiotemporal dynamics of sandflies in the Mediterranean Languedoc region (Roquedur) Tj ETQq1 1 0.784314 rgBT /Overlock 2.5 39	2.5	39
50	Spatio-temporal Genetic Structuring of <i>Leishmania major</i> in Tunisia by Microsatellite Analysis. <i>PLoS Neglected Tropical Diseases</i> , 2015, 9, e0004017.	3.0	8
51	Identification of phlebotomine sand flies using one MALDI-TOF MS reference database and two mass spectrometer systems. <i>Parasites and Vectors</i> , 2015, 8, 266.	2.5	66
52	Genetic Evolution of <i>Mycobacterium bovis</i> Causing Tuberculosis in Livestock and Wildlife in France since 1978. <i>PLoS ONE</i> , 2015, 10, e0117103.	2.5	85
53	A primer for <i>Leishmania</i> population genetic studies. <i>Trends in Parasitology</i> , 2015, 31, 52-59.	3.3	30
54	Response to Tibayrenc et al.: can recombination in <i>Leishmania</i> parasites be so rare?. <i>Trends in Parasitology</i> , 2015, 31, 280-281.	3.3	5

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55	Reduced turn-around time for Mycobacterium tuberculosis drug susceptibility testing with a proportional agar microplate assay. <i>Clinical Microbiology and Infection</i> , 2015, 21, 1084-1092.	6.0	9
56	New microsatellite markers for multi-scale genetic studies on <i>Phlebotomus ariasi</i> Tonnoir, vector of <i>Leishmania infantum</i> in the Mediterranean area. <i>Acta Tropica</i> , 2015, 142, 79-85.	2.0	5
57	<i>Mycobacterium tuberculosis</i> : ecology and evolution of a human bacterium. <i>Journal of Medical Microbiology</i> , 2015, 64, 1261-1269.	1.8	97
58	Comparison of <i>Leishmania killicki</i> (syn. <i>L. tropica</i>) and <i>Leishmania tropica</i> Population Structure in Maghreb by Microsatellite Typing. <i>PLoS Neglected Tropical Diseases</i> , 2015, 9, e0004204.	3.0	10
59	<i>Mycobacterium bovis</i> in Burkina Faso: Epidemiologic and Genetic Links between Human and Cattle Isolates. <i>PLoS Neglected Tropical Diseases</i> , 2014, 8, e3142.	3.0	24
60	Methicillin-Sensitive <i>Staphylococcus aureus</i> CC398 in Intensive Care Unit, France. <i>Emerging Infectious Diseases</i> , 2014, 20, 1511-1515.	4.3	22
61	Sacroiliitis secondary to catheter-related bacteremia due to <i>Mycobacterium abscessus</i> (sensu) Tj ETQq1 1 0.784314 rgBT /Overlock 101	3.8	4
62	Parasitic genotypes appear to differ in leishmaniasis patients compared with asymptomatic related carriers. <i>International Journal for Parasitology</i> , 2013, 43, 389-397.	3.1	19
63	Of parasites and men. <i>Infection, Genetics and Evolution</i> , 2013, 20, 61-70.	2.3	24
64	Rifampin-Resistant <i>Mycobacterium bovis</i> BCG-Induced Disease in HIV-Infected Infant, Vietnam. <i>Emerging Infectious Diseases</i> , 2013, 19, 1168-1168.	4.3	5
65	Rifampin-Resistant <i>Mycobacterium bovis</i> BCG-Induced Disease in HIV-Infected Infant, Vietnam. <i>Emerging Infectious Diseases</i> , 2013, 19, 1168-1168.	4.3	2
66	Genetic Diversity and Population Structure of <i>Mycobacterium marinum</i> : New Insights into Host and Environmental Specificities. <i>Journal of Clinical Microbiology</i> , 2012, 50, 3627-3634.	3.9	18
67	DNAGear- a free software for spa type identification in <i>Staphylococcus aureus</i> . <i>BMC Research Notes</i> , 2012, 5, 642.	1.4	24
68	Wing size and shape variation of <i>Phlebotomus papatasi</i> (Diptera: Psychodidae) populations from the south and north slopes of the Atlas Mountains in Morocco. <i>Journal of Vector Ecology</i> , 2012, 37, 137-147.	1.0	26
69	Genetic diversity, population structure and drug resistance of <i>Mycobacterium tuberculosis</i> in Peru. <i>Infection, Genetics and Evolution</i> , 2012, 12, 577-585.	2.3	33
70	High Prevalence of Beijing and EAI4-VNM Genotypes among <i>M. tuberculosis</i> Isolates in Northern Vietnam: Sampling Effect, Rural and Urban Disparities. <i>PLoS ONE</i> , 2012, 7, e45553.	2.5	24
71	Ecology of Phlebotomine Sand Flies in the Rural Community of Mont Rolland (Thiã's Region, Senegal): Area of Transmission of Canine Leishmaniasis. <i>PLoS ONE</i> , 2011, 6, e14773.	2.5	39
72	Reproductive strategies and population structure in <i>Leishmania</i> : substantial amount of sex in <i>Leishmania Viannia guyanensis</i> . <i>Molecular Ecology</i> , 2011, 20, 3116-3127.	3.9	37

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73	Clinical pleiomorphism in human leishmaniasis, with special mention of asymptomatic infection. <i>Clinical Microbiology and Infection</i> , 2011, 17, 1451-1461.	6.0	101
74	Multifaceted Population Structure and Reproductive Strategy in <i>Leishmania donovani</i> Complex in One Sudanese Village. <i>PLoS Neglected Tropical Diseases</i> , 2011, 5, e1448.	3.0	21
75	Canine visceral leishmaniasis caused by <i>Leishmania infantum</i> in Senegal: risk of emergence in humans?. <i>Microbes and Infection</i> , 2010, 12, 1219-1225.	1.9	26
76	Highly structured genetic diversity of the <i>Mycobacterium tuberculosis</i> population in Djibouti. <i>Clinical Microbiology and Infection</i> , 2010, 16, 1023-1026.	6.0	8
77	Pulmonary tuberculosis due to <i>Mycobacterium microti</i> : a study of six recent cases in France. <i>Journal of Medical Microbiology</i> , 2010, 59, 984-989.	1.8	62
78	Intrafamilial Cluster of Pulmonary Tuberculosis Due to <i>Mycobacterium bovis</i> of the African 1 Clonal Complex. <i>Journal of Clinical Microbiology</i> , 2010, 48, 4680-4683.	3.9	12
79	â€œEverything You Always Wanted to Know about Sex (but Were Afraid to Ask)â€•in <i>Leishmania</i> after Two Decades of Laboratory and Field Analyses. <i>PLoS Pathogens</i> , 2010, 6, e1001004.	4.7	60
80	Additional Haplogroups of <i>Toxoplasma gondii</i> out of Africa: Population Structure and Mouse-Virulence of Strains from Gabon. <i>PLoS Neglected Tropical Diseases</i> , 2010, 4, e876.	3.0	117
81	A battery of 12 microsatellite markers for genetic analysis of the <i>Leishmania</i> (<i>Viannia</i>) <i>guyanensis</i> complex. <i>Parasitology</i> , 2010, 137, 1879-1884.	1.5	17
82	Extreme inbreeding in <i>Leishmania braziliensis</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 10224-10229.	7.1	158
83	Identification of Old World <i>Leishmania</i> spp. by specific polymerase chain reaction amplification of cysteine proteinase B genes and rapid dipstick detection. <i>Diagnostic Microbiology and Infectious Disease</i> , 2009, 63, 173-181.	1.8	24
84	PERMANENT GENETIC RESOURCES: A set of 12 microsatellite loci for genetic studies of <i>Leishmania braziliensis</i> . <i>Molecular Ecology Resources</i> , 2008, 8, 351-353.	4.8	16
85	The promastigote surface antigen gene family of the <i>Leishmania</i> parasite: differential evolution by positive selection and recombination. <i>BMC Evolutionary Biology</i> , 2008, 8, 292.	3.2	27
86	First Molecular Epidemiology Study of <i>Mycobacterium tuberculosis</i> in Burkina Faso. <i>Journal of Clinical Microbiology</i> , 2007, 45, 921-927.	3.9	71
87	Specific <i>cpb</i> copies within the <i>Leishmania donovani</i> complex: evolutionary interpretations and potential clinical implications in humans. <i>Parasitology</i> , 2007, 134, 379.	1.5	29
88	Genetic diversity and population structure of <i>Mycobacterium tuberculosis</i> in HIV-1-infected compared with uninfected individuals in Burkina Faso. <i>Aids</i> , 2007, 21, 248-250.	2.2	8
89	A microculture technique for isolating live <i>Leishmania</i> parasites from peripheral blood of visceral leishmaniasis patients. <i>Acta Tropica</i> , 2007, 102, 197-200.	2.0	17
90	<i>Leishmania</i> and the Leishmaniasis: A Parasite Genetic Update and Advances in Taxonomy, Epidemiology and Pathogenicity in Humans. <i>Advances in Parasitology</i> , 2007, 64, 1-458.	3.2	245

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91	Clonal propagation and the fast generation of karyotype diversity: an in vitro Leishmania model. <i>Parasitology</i> , 2007, 134, 33-39.	1.5	15
92	Tuberculosis transmission in a high incidence area: A retrospective molecular epidemiological study of <i>Mycobacterium tuberculosis</i> in Casablanca, Morocco. <i>Infection, Genetics and Evolution</i> , 2007, 7, 636-644.	2.3	17
93	Species-specific PCR assay for <i>L. infantum</i> / <i>L. donovani</i> discrimination. <i>Acta Tropica</i> , 2006, 100, 241-245.	2.0	55
94	Cryptosporidium Population Genetics: Evidence of Clonality in Isolates from France and Haiti. <i>Journal of Eukaryotic Microbiology</i> , 2006, 53, S33-S36.	1.7	38
95	TreeDyn: towards dynamic graphics and annotations for analyses of trees. <i>BMC Bioinformatics</i> , 2006, 7, 439.	2.6	910
96	Un cas inhabituel de spondylodiscite à <i>Staphylococcus saccharolyticus</i> . <i>Revue Du Rhumatisme (Edition) Tj ETQq0 0.0 rgBT /Overlock 10</i>	0.0	0
97	American tegumentary leishmaniasis: antigen-gene polymorphism, taxonomy and clinical pleomorphism. <i>Infection, Genetics and Evolution</i> , 2005, 5, 109-116.	2.3	75
98	Unusual case of spondylodiscitis due to <i>Staphylococcus saccharolyticus</i> . <i>Joint Bone Spine</i> , 2005, 72, 91-93.	1.6	19
99	Fluorogenic Assay for Molecular Typing of the <i>Leishmania donovani</i> Complex: Taxonomic and Clinical Applications. <i>Journal of Infectious Diseases</i> , 2005, 192, 685-692.	4.0	25
100	Molecular Evidence that Nasal Carriage of <i>Staphylococcus aureus</i> Plays a Role in Respiratory Tract Infections of Critically Ill Patients. <i>Journal of Clinical Microbiology</i> , 2005, 43, 3491-3493.	3.9	35
101	Genetic Diversity and Population Structure of <i>Mycobacterium tuberculosis</i> in Casablanca, a Moroccan City with High Incidence of Tuberculosis. <i>Journal of Clinical Microbiology</i> , 2004, 42, 461-466.	3.9	22
102	Predominant clonal evolution leads to a close parity between gene expression profiles and subspecific phylogeny in <i>Trypanosoma cruzi</i> . <i>Molecular and Biochemical Parasitology</i> , 2004, 137, 133-141.	1.1	18
103	Genetic diversity, clonality and sexuality in <i>Toxoplasma gondii</i> . <i>International Journal for Parasitology</i> , 2004, 34, 1185-1196.	3.1	312
104	Rocking the curve. <i>Trends in Microbiology</i> , 2004, 12, 534-536.	7.7	4
105	Linkage disequilibrium between minisatellite loci supports clonal evolution of <i>Mycobacterium tuberculosis</i> in a high tuberculosis incidence area. <i>Molecular Microbiology</i> , 2003, 47, 529-538.	2.5	171
106	Microsatellite analysis of <i>Toxoplasma gondii</i> shows considerable polymorphism structured into two main clonal groups. <i>International Journal for Parasitology</i> , 2002, 32, 27-38.	3.1	236
107	Evolutionary genetics and molecular diagnosis of <i>Leishmania</i> species. <i>Transactions of the Royal Society of Tropical Medicine and Hygiene</i> , 2002, 96, S9-S13.	1.8	22
108	Genetic heterogeneity and phylogenetic status of <i>Leishmania (Leishmania) infantum</i> zymodeme MON-1: epidemiological implications. <i>Parasitology</i> , 2001, 123, 425-432.	1.5	47

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109	Visceral Leishmaniasis in a German Child Who Had Never Entered a Known Endemic Area: Case Report and Review of the Literature. <i>Clinical Infectious Diseases</i> , 2001, 32, 302-306.	5.8	125
110	High-resolution minisatellite-based typing as a portable approach to global analysis of <i>Mycobacterium tuberculosis</i> molecular epidemiology. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2001, 98, 1901-1906.	7.1	393
111	Is <i>Leishmania (Viannia) peruviana</i> a Distinct Species? A MLEE/RAPD Evolutionary Genetics Answer. <i>Journal of Eukaryotic Microbiology</i> , 2000, 47, 197-207.	1.7	37
112	Molecular epidemiology and evolutionary genetics of <i>Leishmania</i> parasites. <i>International Journal for Parasitology</i> , 1999, 29, 1137-1147.	3.1	51
113	Unusual cutaneous lesions in two patients with visceral leishmaniasis and HIV infection. <i>Journal of the American Academy of Dermatology</i> , 1999, 41, 847-850.	1.2	90
114	Polymerase chain reaction-based identification of New World <i>Leishmania</i> species complexes by specific kDNA probes. <i>Acta Tropica</i> , 1999, 73, 283-293.	2.0	20
115	A phylogenetic analysis of the <i>Trypanosoma cruzi</i> genome project CL Brener reference strain by multilocus enzyme electrophoresis and multiprimer random amplified polymorphic DNA fingerprinting. <i>Molecular and Biochemical Parasitology</i> , 1998, 92, 253-263.	1.1	72
116	Population Structure of <i>Trypanosoma brucei</i> s. l. in Cote D'Ivoire Assayed by Multilocus Enzyme Electrophoresis: Epidemiological and Taxonomical Considerations. <i>Journal of Parasitology</i> , 1997, 83, 19.	0.7	6
117	Evidence for Hybridization by Multilocus Enzyme Electrophoresis and Random Amplified Polymorphic DNA Between <i>Leishmania braziliensis</i> and <i>Leishmania panamensis/guyanensis</i> in Ecuador. <i>Journal of Eukaryotic Microbiology</i> , 1997, 44, 408-411.	1.7	81
118	Putative <i>Leishmania</i> hybrids in the Eastern Andean valley of Huanuco, Peru. <i>Acta Tropica</i> , 1995, 59, 293-307.	2.0	96
119	Pulmonary Tuberculosis and <i>Mycobacterium Tuberculosis</i> : Modern Molecular Epidemiology and Perspectives. , 0, , 1-29.		6
120	Metabarcoding: A Powerful Yet Still Underestimated Approach for the Comprehensive Study of Vector-Borne Pathogen Transmission Cycles and Their Dynamics. , 0, , .		7