

Hiroto Kato

List of Publications by Year in descending order

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142
papers

3,412
citations

136740

32
h-index

189595

50
g-index

144
all docs

144
docs citations

144
times ranked

3420
citing authors

#	ARTICLE	IF	CITATIONS
1	Induction of Bcl-x _L Expression by Human T-Cell Leukemia Virus Type 1 Tax through NF- κ B in Apoptosis-Resistant T-Cell Transfectants with Tax. <i>Journal of Virology</i> , 1999, 73, 7981-7987.	1.5	201
2	Venezuela's humanitarian crisis, resurgence of vector-borne diseases, and implications for spillover in the region. <i>Lancet Infectious Diseases</i> , The, 2019, 19, e149-e161.	4.6	138
3	Evidence For Early Aging in the Mucosal Immune System. <i>Journal of Immunology</i> , 2000, 165, 5352-5359.	0.4	97
4	Development of a loop-mediated isothermal amplification method for rapid mass-screening of sand flies for Leishmania infection. <i>Acta Tropica</i> , 2014, 132, 1-6.	0.9	94
5	High degree of conservancy among secreted salivary gland proteins from two geographically distant <i>Phlebotomus duboscqi</i> sandflies populations (Mali and Kenya). <i>BMC Genomics</i> , 2006, 7, 226.	1.2	93
6	DETECTION AND IDENTIFICATION OF LEISHMANIA SPECIES WITHIN NATURALLY INFECTED SAND FLIES IN THE ANDEAN AREAS OF ECUADOR BY A POLYMERASE CHAIN REACTION. <i>American Journal of Tropical Medicine and Hygiene</i> , 2005, 72, 87-93.	0.6	92
7	Loop-mediated isothermal amplification (LAMP): An advanced molecular point-of-care technique for the detection of Leishmania infection. <i>PLoS Neglected Tropical Diseases</i> , 2019, 13, e0007698.	1.3	86
8	Phylogenic analysis of the genus Leishmania by cytochrome b gene sequencing. <i>Experimental Parasitology</i> , 2009, 121, 352-361.	0.5	79
9	Prevention of Adult T-Cell Leukemia-Like Lymphoproliferative Disease in Rats by Adoptively Transferred T Cells from a Donor Immunized with Human T-Cell Leukemia Virus Type 1 Tax-Coding DNA Vaccine. <i>Journal of Virology</i> , 2000, 74, 9610-9616.	1.5	73
10	Genotyping and Quantitation of Noroviruses in Oysters from Two Distinct Sea Areas in Japan. <i>Microbiology and Immunology</i> , 2007, 51, 177-184.	0.7	68
11	Regression of Human T-cell Leukemia Virus Type I (HTLV-I)-Associated Lymphomas in a Rat Model: Peptide-Induced T-Cell Immunity. <i>Journal of the National Cancer Institute</i> , 2001, 93, 1775-1783.	3.0	63
12	Establishment of a Mass Screening Method of Sand Fly Vectors for Leishmania Infection by Molecular Biological Methods. <i>American Journal of Tropical Medicine and Hygiene</i> , 2007, 77, 324-329.	0.6	59
13	Lack of oral tolerance in aging is due to sequential loss of Peyer's patch cell interactions. <i>International Immunology</i> , 2003, 15, 145-158.	1.8	57
14	Expansion of Human T-Cell Leukemia Virus Type 1 (HTLV-1) Reservoir in Orally Infected Rats: Inverse Correlation with HTLV-1-Specific Cellular Immune Response. <i>Journal of Virology</i> , 2003, 77, 2956-2963.	1.5	52
15	DNA barcoding for identification of sand fly species (Diptera: Psychodidae) from leishmaniasis-endemic areas of Peru. <i>Acta Tropica</i> , 2015, 145, 45-51.	0.9	52
16	Molecular Epidemiology for Vector Research on Leishmaniasis. <i>International Journal of Environmental Research and Public Health</i> , 2010, 7, 814-826.	1.2	51
17	Diffuse and disseminated cutaneous leishmaniasis: clinical cases experienced in Ecuador and a brief review. <i>Tropical Medicine and Health</i> , 2016, 44, 2.	1.0	50
18	Use of FTA Cards for Direct Sampling of Patients' Lesions in the Ecological Study of Cutaneous Leishmaniasis. <i>Journal of Clinical Microbiology</i> , 2010, 48, 3661-3665.	1.8	49

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19	A rapid molecular diagnosis of cutaneous leishmaniasis by colorimetric malachite green-loop-mediated isothermal amplification (LAMP) combined with an FTA card as a direct sampling tool. <i>Acta Tropica</i> , 2016, 153, 116-119.	0.9	49
20	Oral Tolerance Revisited: Prior Oral Tolerization Abrogates Cholera Toxin-Induced Mucosal IgA Responses. <i>Journal of Immunology</i> , 2001, 166, 3114-3121.	0.4	45
21	First Detection of <i>Leishmania tropica</i> DNA and <i>Trypanosoma</i> Species in <i>Sergentomyia</i> Sand Flies (Diptera: Psychodidae) from an Outbreak Area of Cutaneous Leishmaniasis in Ghana. <i>PLoS Neglected Tropical Diseases</i> , 2014, 8, e2630.	1.3	44
22	Analysis of salivary gland transcripts of the sand fly <i>Lutzomyia ayacuchensis</i> , a vector of Andean-type cutaneous leishmaniasis. <i>Infection, Genetics and Evolution</i> , 2013, 13, 56-66.	1.0	43
23	Molecular Mass Screening to Incriminate Sand Fly Vectors of Andean-type Cutaneous Leishmaniasis in Ecuador and Peru. <i>American Journal of Tropical Medicine and Hygiene</i> , 2008, 79, 719-721.	0.6	42
24	Differentiation of feline coronavirus type I and II infections by virus neutralization test. <i>Veterinary Microbiology</i> , 2007, 124, 348-352.	0.8	41
25	A repertoire of the dominant transcripts from the salivary glands of the blood-sucking bug, <i>Triatoma dimidiata</i> , a vector of Chagas disease. <i>Infection, Genetics and Evolution</i> , 2010, 10, 184-191.	1.0	40
26	ATYPICAL CLINICAL VARIANTS IN NEW WORLD CUTANEOUS LEISHMANIASIS: DISSEMINATED, ERYSIPELOID, AND RECIDIVA CUTIS DUE TO <i>LEISHMANIA</i> (V.) <i>PANAMENSIS</i> . <i>American Journal of Tropical Medicine and Hygiene</i> , 2005, 73, 281-284.	0.6	39
27	A revisit of mucosal IgA immunity and oral tolerance. <i>Acta Odontologica Scandinavica</i> , 2001, 59, 301-308.	0.9	38
28	Detection of multiple sapovirus genotypes and genogroups in oyster-associated outbreaks. <i>Japanese Journal of Infectious Diseases</i> , 2009, 62, 63-6.	0.5	38
29	MULTILOCUS ENZYME ELECTROPHORESIS AND CYTOCHROME B GENE SEQUENCING-BASED IDENTIFICATION OF <i>LEISHMANIA</i> ISOLATES FROM DIFFERENT FOCI OF CUTANEOUS LEISHMANIASIS IN PAKISTAN. <i>American Journal of Tropical Medicine and Hygiene</i> , 2006, 75, 261-266.	0.6	37
30	Induction of Adult T-Cell Leukemia-Like Lymphoproliferative Disease and Its Inhibition by Adoptive Immunotherapy in T-Cell-Deficient Nude Rats Inoculated with Syngeneic Human T-Cell Leukemia Virus Type 1-Immortalized Cells. <i>Journal of Virology</i> , 1999, 73, 6031-6040.	1.5	36
31	Multilocus characterization and phylogenetic analysis of <i>Leishmania siamensis</i> isolated from autochthonous visceral leishmaniasis cases, southern Thailand. <i>BMC Microbiology</i> , 2013, 13, 60.	1.3	36
32	Detection and identification of <i>Leishmania</i> species within naturally infected sand flies in the andean areas of ecuador by a polymerase chain reaction. <i>American Journal of Tropical Medicine and Hygiene</i> , 2005, 72, 87-93.	0.6	36
33	Polymorphisms of cytochrome <i>b</i> gene in <i>Leishmania</i> parasites and their relation to types of cutaneous leishmaniasis lesions in Pakistan. <i>Journal of Dermatology</i> , 2008, 35, 76-85.	0.6	32
34	Involvement of CD4 ⁺ Foxp3 ⁺ Regulatory T Cells in Persistence of <i>Leishmania donovani</i> in the Liver of Alymphoplastic aly/aly Mice. <i>PLoS Neglected Tropical Diseases</i> , 2012, 6, e1798.	1.3	31
35	Leishmaniases in Ecuador: Comprehensive review and current status. <i>Acta Tropica</i> , 2017, 166, 299-315.	0.9	31
36	Oral Administration of Human T-Cell Leukemia Virus Type 1 Induces Immune Unresponsiveness with Persistent Infection in Adult Rats. <i>Journal of Virology</i> , 1998, 72, 7289-7293.	1.5	31

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37	A trial of immunotherapy against <i>Leishmania amazonensis</i> infection in vitro and in vivo with Z-100, a polysaccharide obtained from <i>Mycobacterium tuberculosis</i> , alone or combined with meglumine antimoniate. <i>Journal of Antimicrobial Chemotherapy</i> , 2007, 59, 1123-1129.	1.3	29
38	Functional characterization of a salivary apyrase from the sand fly, <i>Phlebotomus duboscqi</i> , a vector of <i>Leishmania major</i> . <i>Journal of Insect Physiology</i> , 2009, 55, 1044-1049.	0.9	29
39	First human cases of <i>Leishmania (Viannia) naiffi</i> infection in Ecuador and identification of its suspected vector species. <i>Acta Tropica</i> , 2013, 128, 710-713.	0.9	29
40	PCR-RFLP analyses of <i>Leishmania</i> species causing cutaneous and mucocutaneous leishmaniasis revealed distribution of genetically complex strains with hybrid and mito-nuclear discordance in Ecuador. <i>PLoS Neglected Tropical Diseases</i> , 2019, 13, e0007403.	1.3	29
41	Development of Human T-Cell Leukemia Virus Type 1-Transformed Tumors in Rats following Suppression of T-Cell Immunity by CD80 and CD86 Blockade. <i>Journal of Virology</i> , 2000, 74, 428-435.	1.5	28
42	Identification and characterization of a salivary adenosine deaminase from the sand fly <i>Phlebotomus duboscqi</i> , the vector of <i>Leishmania major</i> in sub-Saharan Africa. <i>Journal of Experimental Biology</i> , 2007, 210, 733-740.	0.8	27
43	Natural Infection of <i>Lutzomyia tortura</i> with <i>Leishmania (Viannia) naiffi</i> in an Amazonian Area of Ecuador. <i>American Journal of Tropical Medicine and Hygiene</i> , 2008, 79, 438-440.	0.6	27
44	Immunological Aspects of Rat Models of HTLV Type 1-Infected T Lymphoproliferative Disease. <i>AIDS Research and Human Retroviruses</i> , 2000, 16, 1737-1740.	0.5	26
45	Molecular Typing of Sand Fly Species (Diptera, Psychodidae, Phlebotominae) from Areas Endemic for Leishmaniasis in Ecuador by PCR-RFLP of 18S Ribosomal RNA Gene. <i>Journal of Veterinary Medical Science</i> , 2008, 70, 907-913.	0.3	26
46	Natural infection of the sand fly <i>Phlebotomus kazeruni</i> by <i>Trypanosoma</i> species in Pakistan. <i>Parasites and Vectors</i> , 2010, 3, 10.	1.0	26
47	Establishment of a mass screening method of sand fly vectors for <i>Leishmania</i> infection by molecular biological methods. <i>American Journal of Tropical Medicine and Hygiene</i> , 2007, 77, 324-9.	0.6	26
48	Molecular cloning of equine interleukin-1 α and -1 β cDNAs. <i>Veterinary Immunology and Immunopathology</i> , 1995, 48, 221-231.	0.5	24
49	<i>Leishmania</i> isoenzyme polymorphisms in Ecuador: Relationships with geographic distribution and clinical presentation. <i>BMC Infectious Diseases</i> , 2006, 6, 139.	1.3	24
50	Man-biting sand fly species and natural infection with the <i>Leishmania</i> promastigote in leishmaniasis-endemic areas of Ecuador. <i>Acta Tropica</i> , 2014, 140, 41-49.	0.9	24
51	Establishment of a Seronegative Human T-Cell Leukemia Virus Type 1 (HTLV-1) Carrier State in Rats Inoculated with a Syngeneic HTLV-1-Immortalized T-Cell Line Preferentially Expressing Tax. <i>Journal of Virology</i> , 1999, 73, 6436-6443.	1.5	24
52	First Evidence of a Hybrid of <i>Leishmania (Viannia) braziliensis/L. (V.) peruviana</i> DNA Detected from the Phlebotomine Sand Fly <i>Lutzomyia tejadai</i> in Peru. <i>PLoS Neglected Tropical Diseases</i> , 2016, 10, e0004336.	1.3	24
53	Natural Infections of Man-Biting Sand Flies by <i>Leishmania</i> and <i>Trypanosoma</i> Species in the Northern Peruvian Andes. <i>Vector-Borne and Zoonotic Diseases</i> , 2011, 11, 515-521.	0.6	23
54	Molecular prevalence and genetic diversity of bovine <i>Theileria orientalis</i> in Myanmar. <i>Parasitology International</i> , 2014, 63, 640-645.	0.6	22

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55	Geographic Distribution of Leishmania Species in Ecuador Based on the Cytochrome B Gene Sequence Analysis. PLoS Neglected Tropical Diseases, 2016, 10, e0004844.	1.3	22
56	Leishmania species identification using FTA card sampling directly from patients's cutaneous lesions in the state of Lara, Venezuela. Transactions of the Royal Society of Tropical Medicine and Hygiene, 2011, 105, 561-567.	0.7	21
57	The isolation and molecular characterization of Leishmania spp. from patients with American tegumentary leishmaniasis in northwest Argentina. Acta Tropica, 2014, 131, 16-21.	0.9	21
58	Molecular survey of Babesia infections in cattle from different areas of Myanmar. Ticks and Tick-borne Diseases, 2016, 7, 204-207.	1.1	21
59	Molecular analysis of multidrug resistance in feline lymphoma cells. American Journal of Veterinary Research, 2000, 61, 1122-1127.	0.3	20
60	Molecular detection and genetic diversity of Babesia gibsoni in dogs in Bangladesh. Infection, Genetics and Evolution, 2015, 31, 53-60.	1.0	20
61	Multilocus enzyme electrophoresis and cytochrome B gene sequencing-based identification of Leishmania isolates from different foci of cutaneous leishmaniasis in Pakistan. American Journal of Tropical Medicine and Hygiene, 2006, 75, 261-6.	0.6	20
62	Molecular characterization of feline immunodeficiency virus genome obtained directly from organs of a naturally infected cat with marked neurological symptoms and encephalitis. Archives of Virology, 1996, 141, 1933-1948.	0.9	19
63	Correlation of Major Histocompatibility Complex Class I Downregulation with Resistance of Human T-Cell Leukemia Virus Type 1-Infected T Cells to Cytotoxic T-Lymphocyte Killing in a Rat Model. Journal of Virology, 2002, 76, 7010-7019.	1.5	19
64	PCR-Based Detection of <i>Leishmania donovani</i> DNA in a Stray Dog from a Visceral Leishmaniasis Endemic Focus in Bangladesh. Journal of Veterinary Medical Science, 2013, 75, 75-78.	0.3	19
65	An outbreak of Leishmania major from an endemic to a non-endemic region posed a public health threat in Iraq from 2014-2017: Epidemiological, molecular and phylogenetic studies. PLoS Neglected Tropical Diseases, 2018, 12, e0006255.	1.3	19
66	Genetic diversity of ribosomal RNA internal transcribed spacer sequences in Lutzomyia species from areas endemic for New World cutaneous leishmaniasis. Acta Tropica, 2009, 112, 131-136.	0.9	18
67	Genetic diversity of Leishmania donovani/infantum complex in China through microsatellite analysis. Infection, Genetics and Evolution, 2014, 22, 112-119.	1.0	18
68	Distribution of Lutzomyia ayacuchensis, the vector of Andean-type cutaneous leishmaniasis, at different altitudes on the Andean slope of Ecuador. Acta Tropica, 2014, 137, 118-122.	0.9	18
69	Identification of causative Leishmania species in Giemsa-stained smears prepared from patients with cutaneous leishmaniasis in Peru using PCR-RFLP. Acta Tropica, 2016, 158, 83-87.	0.9	18
70	The identification of sandfly species, from an area of Argentina with endemic leishmaniasis, by the PCR-based analysis of the gene coding for 18S ribosomal RNA. Annals of Tropical Medicine and Parasitology, 2007, 101, 247-253.	1.6	17
71	Ayadualin, a novel RGD peptide with dual antihemostatic activities from the sand fly Lutzomyia ayacuchensis, a vector of Andean-type cutaneous leishmaniasis. Biochimie, 2015, 112, 49-56.	1.3	17
72	Molecular cloning and expression of canine interleukin 8 cDNA. Cytokine, 1994, 6, 455-461.	1.4	16

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73	Andean cutaneous leishmaniasis (Andean-CL, uta) in Peru and Ecuador: the vector <i>Lutzomyia</i> sand flies and reservoir mammals. <i>Acta Tropica</i> , 2018, 178, 264-275.	0.9	16
74	Molecular mass screening to incriminate sand fly vectors of Andean-type cutaneous leishmaniasis in Ecuador and Peru. <i>American Journal of Tropical Medicine and Hygiene</i> , 2008, 79, 719-21.	0.6	16
75	Molecular cloning and functional expression of equine interleukin-1 receptor antagonist. <i>Veterinary Immunology and Immunopathology</i> , 1997, 56, 221-231.	0.5	15
76	High SDF-1 Expression in HIV-1 Carriers Does Not Correlate with CD8+T-Cell-Mediated Suppression of Viral Replication. <i>Virology</i> , 1998, 244, 467-472.	1.1	14
77	Molecular and Serological Evidence of <i>Leishmania</i> Infection in Stray Dogs from Visceral Leishmaniasis Endemic Areas of Bangladesh. <i>American Journal of Tropical Medicine and Hygiene</i> , 2016, 95, 795-799.	0.6	14
78	Atypical clinical variants in New World cutaneous leishmaniasis: disseminated, erysipeloid, and recidiva cutis due to <i>Leishmania</i> (<i>V.</i>) <i>panamensis</i> . <i>American Journal of Tropical Medicine and Hygiene</i> , 2005, 73, 281-4.	0.6	14
79	Gammopathy with two M-components in a dog with IgA-type multiple myeloma. <i>Veterinary Immunology and Immunopathology</i> , 1995, 49, 161-168.	0.5	13
80	Genetic divergence in populations of <i>Lutzomyia ayacuchensis</i> , a vector of Andean-type cutaneous leishmaniasis, in Ecuador and Peru. <i>Acta Tropica</i> , 2015, 141, 79-87.	0.9	13
81	Andean cutaneous leishmaniasis (Andean-CL, uta) in Peru and Ecuador: the causative <i>Leishmania</i> parasites and clinico-epidemiological features. <i>Acta Tropica</i> , 2018, 177, 135-145.	0.9	13
82	Prevalence of Genetically Complex <i>Leishmania</i> Strains With Hybrid and Mito-Nuclear Discordance. <i>Frontiers in Cellular and Infection Microbiology</i> , 2021, 11, 625001.	1.8	13
83	Dimiconin, a novel coagulation inhibitor from the kissing bug, <i>Triatoma dimidiata</i> , a vector of Chagas disease. <i>Journal of Experimental Biology</i> , 2012, 215, 3597-602.	0.8	12
84	Salivary gland transcripts of the kissing bug, <i>Panstrongylus chinai</i> , a vector of Chagas disease. <i>Acta Tropica</i> , 2017, 174, 122-129.	0.9	12
85	Comparison of LAMP and PCR for molecular mass screening of sand flies for <i>Leishmania martiniquensis</i> infection. <i>Memorias Do Instituto Oswaldo Cruz</i> , 2017, 112, 100-107.	0.8	12
86	Genotyping of sand fly species in Peruvian Andes where leishmaniasis is endemic. <i>Acta Tropica</i> , 2012, 121, 93-98.	0.9	11
87	Autochthonous cutaneous leishmaniasis in urban domestic animals (<i>Felis catus</i> / <i>Canis lupus</i>) Tj ETQq1 1 0.784314 rgBT / Overlock 10	0.9	11
88	Genetic diversity of the mitochondrial cytochrome b gene in <i>Lutzomyia</i> spp., with special reference to <i>Lutzomyia peruensis</i> , a main vector of <i>Leishmania</i> (<i>Viannia</i>) <i>peruviana</i> in the Peruvian Andes. <i>Acta Tropica</i> , 2013, 126, 156-163.	0.9	10
89	Seroepidemiological Study of Chagas Disease in the Southern Amazon Region of Ecuador. <i>Tropical Medicine and Health</i> , 2013, 41, 21-25.	1.0	10
90	Sand Fly Fauna (Diptera, Pycnodidae, Phlebotominae) in Different Leishmaniasis-Endemic Areas of Ecuador, Surveyed Using a Newly Named Mini-Shannon Trap. <i>Tropical Medicine and Health</i> , 2014, 42, 163-170.	1.0	10

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91	Intermediate cutaneous leishmaniasis caused by <i>Leishmania (Viannia) braziliensis</i> successfully treated with fluconazole. <i>Clinical and Experimental Dermatology</i> , 2014, 39, 708-712.	0.6	10
92	First Human Cases of <i>Leishmania (Viannia) lainsoni</i> Infection and a Search for the Vector Sand Flies in Ecuador. <i>PLoS Neglected Tropical Diseases</i> , 2016, 10, e0004728.	1.3	10
93	Palm-size and one-inch gel electrophoretic device for reliable and field-applicable analysis of recombinase polymerase amplification. <i>Analytical Methods</i> , 2019, 11, 4969-4976.	1.3	10
94	Further insight into the geographic distribution of <i>Leishmania</i> species in Peru by cytochrome b and mannose phosphate isomerase gene analyses. <i>PLoS Neglected Tropical Diseases</i> , 2019, 13, e0007496.	1.3	10
95	A prospective mechanism and source of cholesterol uptake by <i>Plasmodium falciparum</i> -infected erythrocytes co-cultured with HepG2 cells. <i>Parasitology International</i> , 2021, 80, 102179.	0.6	10
96	Natural infection of <i>Lutzomyia tortura</i> with <i>Leishmania (Viannia) naiffi</i> in an Amazonian area of Ecuador. <i>American Journal of Tropical Medicine and Hygiene</i> , 2008, 79, 438-40.	0.6	10
97	An analysis of reported cases of leishmaniasis in the southern Ecuadorian Amazon region, 1986–2012. <i>Acta Tropica</i> , 2015, 146, 119-126.	0.9	9
98	First identification of <i>L. major</i> in a dog in an endemic area of human cutaneous leishmaniasis in Iraq: molecular and phylogenetic studies. <i>Parasitology Research</i> , 2018, 117, 585-590.	0.6	8
99	A synthetic male-specific sterilization system using the mammalian pro-apoptotic factor in a malaria vector mosquito. <i>Scientific Reports</i> , 2019, 9, 8160.	1.6	8
100	Liver-Directed AAV8 Booster Vaccine Expressing <i>Plasmodium falciparum</i> Antigen Following Adenovirus Vaccine Priming Elicits Sterile Protection in a Murine Model. <i>Frontiers in Immunology</i> , 2021, 12, 612910.	2.2	8
101	Identification of an alternatively spliced transcript of equine interleukin-1 β . <i>Gene</i> , 1996, 177, 11-16.	1.0	7
102	Population genetics of <i>Leishmania (Leishmania) major</i> DNA isolated from cutaneous leishmaniasis patients in Pakistan based on multilocus microsatellite typing. <i>Parasites and Vectors</i> , 2014, 7, 332.	1.0	7
103	Comparison of Recombinant Proteins of Kinesin 39, Heat Shock Protein 70, Heat Shock Protein 83, and Glycoprotein 63 for Antibody Detection of <i>Leishmania martiniquensis</i> Infection. <i>Journal of Eukaryotic Microbiology</i> , 2017, 64, 820-828.	0.8	7
104	Leucine-Rich Repeat Kinase 2 Controls Inflammatory Cytokines Production through NF- κ B Phosphorylation and Antigen Presentation in Bone Marrow-Derived Dendritic Cells. <i>International Journal of Molecular Sciences</i> , 2020, 21, 1890.	1.8	7
105	Molecular cloning of bovine mb-1 cDNA. <i>Veterinary Immunology and Immunopathology</i> , 1996, 52, 191-200.	0.5	6
106	Application of RFLP-PCR-Based Identification for Sand Fly Surveillance in an Area Endemic for Kala-Azar in Mymensingh, Bangladesh. <i>Journal of Parasitology Research</i> , 2012, 2012, 1-4.	0.5	6
107	Nuclear and kinetoplast DNA analyses reveal genetically complex <i>Leishmania</i> strains with hybrid and mito-nuclear discordance in Peru. <i>PLoS Neglected Tropical Diseases</i> , 2020, 14, e0008797.	1.3	6
108	Role of class I major histocompatibility complex-restricted and -unrestricted suppression of human immunodeficiency virus type 1 replication by CD8+ T lymphocytes. <i>Journal of General Virology</i> , 1999, 80, 209-216.	1.3	6

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109	Development of a Monoclonal Antibody-Based Sandwich ELISA for Detection of Guinea Pig Interleukin-2. <i>Journal of Veterinary Medical Science</i> , 2006, 68, 1281-1287.	0.3	5
110	Leishmaniasis caused by <i>Leishmania (Viannia) guyanensis</i> in north-central Pacific region of Ecuador: A clinico-epidemiological feature. <i>Acta Tropica</i> , 2018, 185, 204-211.	0.9	5
111	Comparative Analysis of Bacterial Communities in <i>Lutzomyia ayacuchensis</i> Populations with Different Vector Competence to <i>Leishmania</i> Parasites in Ecuador and Peru. <i>Microorganisms</i> , 2021, 9, 68.	1.6	5
112	Development of a Highly Sensitive Nested PCR and Its Application for the Diagnosis of Cutaneous Leishmaniasis in Sri Lanka. <i>Microorganisms</i> , 2022, 10, 990.	1.6	5
113	Molecular cloning and functional expression of feline thrombopoietin. <i>Veterinary Immunology and Immunopathology</i> , 1998, 66, 225-236.	0.5	4
114	The Attachment and Entry of <i>Leishmania (Leishmania) Major</i> into Macrophages: Observation by Scanning Electron Microscope. <i>Journal of Dermatology</i> , 2005, 32, 534-540.	0.6	4
115	Induction of IL-10- and IFN- γ -producing T-cell responses by autoreactive T-cells expressing human T-cell leukemia virus type I Tax. <i>International Immunology</i> , 2009, 21, 1089-1100.	1.8	4
116	Leishmaniasis Recidiva Cutis and Its Topical Treatment in Ecuador. <i>Tropical Medicine and Health</i> , 2013, 41, 93-94.	1.0	4
117	Salivary gland transcriptome of the Asiatic <i>Triatoma rubrofasciata</i> . <i>Acta Tropica</i> , 2020, 210, 105473.	0.9	4
118	Validation of an In-House ELISA Method in the Diagnosis of Cutaneous Leishmaniasis Caused by <i>Leishmania donovani</i> in Hambantota District, Sri Lanka. <i>Microorganisms</i> , 2022, 10, 921.	1.6	4
119	Salivary lipocalin family proteins from <i>Panstrongylus chinai</i> , a vector of Chagas disease. <i>Data in Brief</i> , 2017, 15, 272-280.	0.5	3
120	Natural <i>Leishmania (Leishmania) mexicana</i> infection and biting activity of anthropophilic sand fly <i>Lutzomyia ayacuchensis</i> in the Ecuadorian Andes. <i>Acta Tropica</i> , 2020, 203, 105321.	0.9	3
121	Anthropophilic phlebotomine sand fly <i>Lutzomyia</i> species and search for the natural <i>Leishmania</i> infections in an area endemic for cutaneous leishmaniasis in Ecuador. <i>Acta Tropica</i> , 2020, 203, 105287.	0.9	3
122	First molecular identification of <i>Leishmania major</i> in <i>Phlebotomus papatasi</i> in an outbreak cutaneous leishmaniasis area in Iraq. <i>Acta Tropica</i> , 2021, 215, 105807.	0.9	3
123	Natural infections of <i>Pintomyia verrucarum</i> and <i>Pintomyia maranonensis</i> by <i>Leishmania (Viannia) peruviana</i> in the Eastern Andes of northern Peru. <i>PLoS Neglected Tropical Diseases</i> , 2021, 15, e0009352.	1.3	3
124	Sand Flies and Their Microbiota. <i>Parasitologia</i> , 2022, 2, 71-87.	0.6	3
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