

Chihiro Sugimoto

List of Publications by Year in descending order

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Version: 2024-02-01

259
papers

6,972
citations

66234

42
h-index

114278

63
g-index

261
all docs

261
docs citations

261
times ranked

5744
citing authors

#	ARTICLE	IF	CITATIONS
1	High infection rate of tick-borne protozoan and rickettsial pathogens of cattle in Malawi and the development of a multiplex PCR for Babesia and Theileria species identification. Acta Tropica, 2022, 231, 106413.	0.9	5
2	Genotyping of <i>Theileria parva</i> populations in vaccinated and non-vaccinated cattle in Malawi. Parasitology, 2022, , 1-28.	0.7	0
3	Molecular detection and characterization of tick-borne hemoparasites and Anaplasmatataceae in dogs in major cities of Malawi. Parasitology Research, 2021, 120, 267-276.	0.6	6
4	Molecular identification and genetic characterization of tick-borne pathogens in sheep and goats at two farms in the central and southern regions of Malawi. Ticks and Tick-borne Diseases, 2021, 12, 101629.	1.1	10
5	Genetic Diversity of African Trypanosomes in Tsetse Flies and Cattle From the Kafue Ecosystem. Frontiers in Veterinary Science, 2021, 8, 599815.	0.9	6
6	Domestic dog demographics and estimates of canine vaccination coverage in a rural area of Zambia for the elimination of rabies. PLoS Neglected Tropical Diseases, 2021, 15, e0009222.	1.3	6
7	Immunization Coverage and Antibody Retention against Rabies in Domestic Dogs in Lusaka District, Zambia. Pathogens, 2021, 10, 738.	1.2	2
8	Screening of tick-borne pathogens in argasid ticks in Zambia: Expansion of the geographic distribution of Rickettsia lusitaniae and Rickettsia hoogstraalii and detection of putative novel Anaplasma species. Ticks and Tick-borne Diseases, 2021, 12, 101720.	1.1	20
9	Complete genome and bimodal genomic structure of the amoebal symbiont Neochlamydia strain S13 revealed by ultra-long reads obtained from MinION. Journal of Human Genetics, 2020, 65, 41-48.	1.1	1
10	The Lethal(2)-Essential-for-Life [L(2)EFL] Gene Family Modulates Dengue Virus Infection in Aedes aegypti. International Journal of Molecular Sciences, 2020, 21, 7520.	1.8	9
11	Molecular identification of trypanosomes in cattle in Malawi using PCR methods and nanopore sequencing: epidemiological implications for the control of human and animal trypanosomiasis. Parasite, 2020, 27, 46.	0.8	5
12	Molecular characterization of <i>Cryptosporidium</i> spp. from patients with diarrhoea in Lusaka, Zambia. Parasite, 2020, 27, 53.	0.8	4
13	Investigation of the piroplasm diversity circulating in wildlife and cattle of the greater Kafue ecosystem, Zambia. Parasites and Vectors, 2020, 13, 599.	1.0	17
14	Development of a Multiplex Loop-Mediated Isothermal Amplification (LAMP) Method for Simultaneous Detection of Spotted Fever Group Rickettsiae and Malaria Parasites by Dipstick DNA Chromatography. Diagnostics, 2020, 10, 897.	1.3	10
15	Diversity of trypanosomes in wildlife of the Kafue ecosystem, Zambia. International Journal for Parasitology: Parasites and Wildlife, 2020, 12, 34-41.	0.6	12
16	Genetic Diversity and Sequence Polymorphism of Two Genes Encoding Theileria parva Antigens Recognized by CD8+ T Cells among Vaccinated and Unvaccinated Cattle in Malawi. Pathogens, 2020, 9, 334.	1.2	9
17	Blood meal sources and bacterial microbiome diversity in wild-caught tsetse flies. Scientific Reports, 2020, 10, 5005.	1.6	23
18	Isolation of Candidatus Bartonella rousetti and Other Bat-associated Bartonellae from Bats and Their Flies in Zambia. Pathogens, 2020, 9, 469.	1.2	20

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19	Development of a bio-inkjet printed LAMP test kit for detecting human African trypanosomiasis. <i>PLoS Neglected Tropical Diseases</i> , 2020, 14, e0008753.	1.3	7
20	Molecular characterization and phylogenetic analysis of <i>Trypanosoma</i> spp. detected from striped leaf-nosed bats (<i>Hipposideros vittatus</i>) in Zambia. <i>International Journal for Parasitology: Parasites and Wildlife</i> , 2019, 9, 234-238.	0.6	3
21	Genetic diversity and population structure of <i>Glossina morsitans morsitans</i> in the active foci of human African trypanosomiasis in Zambia and Malawi. <i>PLoS Neglected Tropical Diseases</i> , 2019, 13, e0007568.	1.3	3
22	Complete Genome Sequence of <i>Rickettsia asiatica</i> Strain Maytaro1284, a Member of Spotted Fever Group <i>Rickettsiae</i> Isolated from an <i>Ixodes ovatus</i> Tick in Japan. <i>Microbiology Resource Announcements</i> , 2019, 8, .	0.3	3
23	Prevalence of hemoprotezoan parasites in small ruminants along a human-livestock-wildlife interface in western Uganda. <i>Veterinary Parasitology: Regional Studies and Reports</i> , 2019, 17, 100309.	0.3	11
24	Field diagnosis and genotyping of chikungunya virus using a dried reverse transcription loop-mediated isothermal amplification (LAMP) assay and MinION sequencing. <i>PLoS Neglected Tropical Diseases</i> , 2019, 13, e0007480.	1.3	19
25	Viral population analysis of the taiga tick, <i>Ixodes persulcatus</i> , by using Batch Learning Self-Organizing Maps and BLAST search. <i>Journal of Veterinary Medical Science</i> , 2019, 81, 401-410.	0.3	30
26	Molecular detection of <i>Rickettsia felis</i> in dogs, rodents and cat fleas in Zambia. <i>Parasites and Vectors</i> , 2019, 12, 168.	1.0	32
27	Diversity of spotted fever group rickettsiae and their association with host ticks in Japan. <i>Scientific Reports</i> , 2019, 9, 1500.	1.6	43
28	A single test approach for accurate and sensitive detection and taxonomic characterization of Trypanosomes by comprehensive analysis of internal transcribed spacer 1 amplicons. <i>PLoS Neglected Tropical Diseases</i> , 2019, 13, e0006842.	1.3	26
29	Isolation of <i>Rickettsia</i> , <i>Rickettsiella</i> , and <i>Spiroplasma</i> from Questing Ticks in Japan Using Arthropod Cells. <i>Vector-Borne and Zoonotic Diseases</i> , 2019, 19, 474-485.	0.6	11
30	Human Borreliosis Caused by a New World Relapsing Fever <i>Borrelia</i> "like Organism in the Old World. <i>Clinical Infectious Diseases</i> , 2019, 69, 107-112.	2.9	36
31	Recombination and purifying and balancing selection determine the evolution of major antigenic protein 1 (map 1) family genes in <i>Ehrlichia ruminantium</i> . <i>Gene</i> , 2019, 683, 216-224.	1.0	8
32	Comparative evaluation of dry and liquid RIME LAMP in detecting trypanosomes in dead tsetse flies. <i>Onderstepoort Journal of Veterinary Research</i> , 2018, 85, e1-e6.	0.6	4
33	The Unique Phylogenetic Position of a Novel Tick-Borne Phlebovirus Ensures an Ixodid Origin of the Genus <i>Phlebovirus</i> . <i>MSphere</i> , 2018, 3, .	1.3	36
34	Establishment of a mouse-tick infection model for <i>Theileria orientalis</i> and analysis of its transcriptome. <i>International Journal for Parasitology</i> , 2018, 48, 915-924.	1.3	9
35	Comparison of Database Search Methods for the Detection of <i>Legionella pneumophila</i> in Water Samples Using Metagenomic Analysis. <i>Frontiers in Microbiology</i> , 2018, 9, 1272.	1.5	13
36	Development and validation of direct dry loop mediated isothermal amplification for diagnosis of <i>Trypanosoma evansi</i> . <i>Veterinary Parasitology</i> , 2018, 260, 53-57.	0.7	13

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37	Direct detection of falciparum and non-falciparum malaria DNA from a drop of blood with high sensitivity by the dried-LAMP system. <i>Parasites and Vectors</i> , 2017, 10, 26.	1.0	35
38	PEGylation of a TLR2-agonist-based vaccine delivery system improves antigen trafficking and the magnitude of ensuing antibody and CD8+ T cell responses. <i>Biomaterials</i> , 2017, 137, 61-72.	5.7	29
39	Putative RNA viral sequences detected in an Ixodes scapularis-derived cell line. <i>Ticks and Tick-borne Diseases</i> , 2017, 8, 103-111.	1.1	23
40	Occurrence of <i>Coxiella burnetii</i> , <i>Ehrlichia canis</i> , <i>Rickettsia</i> species and <i>Anaplasma phagocytophilum</i> -like bacterium in ticks collected from dogs and cats in South Africa. <i>Journal of the South African Veterinary Association</i> , 2017, 88, e1-e6.	0.2	27
41	Horizontally Transferred Genetic Elements in the Tsetse Fly Genome: An Alignment-Free Clustering Approach Using Batch Learning Self-Organising Map (BLSOM). <i>BioMed Research International</i> , 2016, 2016, 1-8.	0.9	23
42	Ungulate malaria parasites. <i>Scientific Reports</i> , 2016, 6, 23230.	1.6	37
43	Molecular detection of spotted fever group rickettsiae in <i>Amblyomma variegatum</i> ticks from Benin. <i>Ticks and Tick-borne Diseases</i> , 2016, 7, 828-833.	1.1	20
44	A PCR-based survey of animal African trypanosomiasis and selected piroplasm parasites of cattle and goats in Zambia. <i>Journal of Veterinary Medical Science</i> , 2016, 78, 1819-1824.	0.3	19
45	Draft Genome Sequences of Three Strains of <i>Ehrlichia ruminantium</i> , a Tick-Borne Pathogen of Ruminants, Isolated from Zimbabwe, The Gambia, and Ghana. <i>Genome Announcements</i> , 2016, 4, .	0.8	4
46	Molecular identification of the <i>Cryptosporidium</i> deer genotype in the Hokkaido sika deer (<i>Cervus</i>) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50	0.6	12
47	Molecular evidence of spotted fever group rickettsiae and Anaplasmataceae from ticks and stray dogs in Bangladesh. <i>Parasitology Research</i> , 2016, 115, 949-955.	0.6	11
48	Human African Trypanosomiasis in the Kafue National Park, Zambia. <i>PLoS Neglected Tropical Diseases</i> , 2016, 10, e0004567.	1.3	11
49	Molecular detection of zoonotic tick-borne pathogens from ticks collected from ruminants in four South African provinces. <i>Journal of Veterinary Medical Science</i> , 2015, 77, 1573-1579.	0.3	44
50	Studies of trypanosomiasis in the Luangwa valley, north-eastern Zambia. <i>Parasites and Vectors</i> , 2015, 8, 497.	1.0	25
51	Molecular epidemiology of pathogenic <i>Leptospira</i> spp. in the straw-colored fruit bat (<i>Eidolon helvum</i>) migrating to Zambia from the Democratic Republic of Congo. <i>Infection, Genetics and Evolution</i> , 2015, 32, 143-147.	1.0	25
52	Direct Blood Dry LAMP: A Rapid, Stable, and Easy Diagnostic Tool for Human African Trypanosomiasis. <i>PLoS Neglected Tropical Diseases</i> , 2015, 9, e0003578.	1.3	104
53	Molecular Detection of <i>Rickettsia africae</i> in <i>Amblyomma variegatum</i> Collected from Sudan. <i>Vector-Borne and Zoonotic Diseases</i> , 2015, 15, 323-325.	0.6	16
54	Determination of the prevalence of trypanosome species in cattle from Monduli district, northern Tanzania, by loop mediated isothermal amplification. <i>Tropical Animal Health and Production</i> , 2015, 47, 1139-1143.	0.5	18

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55	Seroepidemiological Prevalence of Multiple Species of Filoviruses in Fruit Bats (<i>Eidolon</i>) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50	1.9	94
56	Detection and characterization of zoonotic pathogens of free-ranging non-human primates from Zambia. <i>Parasites and Vectors</i> , 2014, 7, 490.	1.0	29
57	Comparative genomic analysis of eight <i>Leptospira</i> strains from Japan and the Philippines revealing the existence of four putative novel genomic islands/islets in <i>L. interrogans</i> serovar Lai strain 56601. <i>Comparative Immunology, Microbiology and Infectious Diseases</i> , 2014, 37, 289-297.	0.7	2
58	Prevalence and characterization of <i>Staphylococcus aureus</i> and <i>Staphylococcus pseudintermedius</i> isolated from companion animals and environment in the veterinary teaching hospital in Zambia, Africa. <i>Comparative Immunology, Microbiology and Infectious Diseases</i> , 2014, 37, 123-130.	0.7	27
59	Determination of the prevalence of African trypanosome species in indigenous dogs of Mambwe district, eastern Zambia, by loop-mediated isothermal amplification. <i>Parasites and Vectors</i> , 2014, 7, 19.	1.0	33
60	Molecular detection of equine trypanosomes in the Sudan. <i>Veterinary Parasitology</i> , 2014, 200, 246-250.	0.7	15
61	Evolution and genetic diversity of <i>Theileria</i> . <i>Infection, Genetics and Evolution</i> , 2014, 27, 250-263.	1.0	182
62	Interactive transcriptome analysis of malaria patients and infecting <i>Plasmodium falciparum</i> . <i>Genome Research</i> , 2014, 24, 1433-1444.	2.4	76
63	Amoebal Endosymbiont <i>Neochlamydia</i> Genome Sequence Illuminates the Bacterial Role in the Defense of the Host Amoebae against <i>Legionella pneumophila</i> . <i>PLoS ONE</i> , 2014, 9, e95166.	1.1	46
64	Microbial Population Analysis of the Salivary Glands of Ticks; A Possible Strategy for the Surveillance of Bacterial Pathogens. <i>PLoS ONE</i> , 2014, 9, e103961.	1.1	95
65	Rapid detection and identification of <i>Theileria equi</i> and <i>Babesia caballi</i> by high-resolution melting (HRM) analysis. <i>Parasitology Research</i> , 2013, 112, 3883-3886.	0.6	14
66	A novel approach, based on BLSOMs (Batch Learning Self-Organizing Maps), to the microbiome analysis of ticks. <i>ISME Journal</i> , 2013, 7, 1003-1015.	4.4	132
67	High prevalence of spotted fever group rickettsiae in <i>Amblyomma variegatum</i> from Uganda and their identification using sizes of intergenic spacers. <i>Ticks and Tick-borne Diseases</i> , 2013, 4, 506-512.	1.1	34
68	Current status of equine piroplasmiasis in the Sudan. <i>Infection, Genetics and Evolution</i> , 2013, 16, 191-199.	1.0	20
69	MDM2 regulates a novel form of incomplete neoplastic transformation of <i>Theileria parva</i> infected lymphocytes. <i>Experimental and Molecular Pathology</i> , 2013, 94, 228-238.	0.9	20
70	Booster responses by oral vaccination with transgenic plants against chicken leucocytozoonosis. <i>Avian Pathology</i> , 2013, 42, 215-220.	0.8	3
71	Proposed vector candidate: <i>Leptotrombidium palpale</i> for Shimokoshi type <i>Orientia tsutsugamushi</i> . <i>Microbiology and Immunology</i> , 2013, 57, 111-117.	0.7	12
72	Whole-Genome Sequencing of <i>Theileria parva</i> Strains Provides Insight into Parasite Migration and Diversification in the African Continent. <i>DNA Research</i> , 2013, 20, 209-220.	1.5	33

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73	Preliminary Investigation of Trypanosomosis in Exotic Dog Breeds from Zambia's Luangwa and Zambezi Valleys Using LAMP. <i>American Journal of Tropical Medicine and Hygiene</i> , 2013, 89, 116-118.	0.6	20
74	First Genetic Detection of <i>Coxiella burnetii</i> in Zambian Livestock. <i>American Journal of Tropical Medicine and Hygiene</i> , 2013, 89, 518-519.	0.6	11
75	Genome Sequence of Methicillin-Resistant <i>Staphylococcus pseudintermedius</i> Sequence Type 233 (ST233) Strain K7, of Human Origin. <i>Genome Announcements</i> , 2013, 1, .	0.8	5
76	Cross-Reactivity of Secondary Antibodies against African Rodents and Application for Sero-Surveillance. <i>Journal of Veterinary Medical Science</i> , 2013, 75, 819-825.	0.3	7
77	Genetic diversity among <i>Trypanosoma (Duttonella) vivax</i> strains from Zambia and Ghana, based on cathepsin L-like gene. <i>Parasite</i> , 2013, 20, 24.	0.8	12
78	Protochlamydia Induces Apoptosis of Human HEp-2 Cells through Mitochondrial Dysfunction Mediated by Chlamydial Protease-Like Activity Factor. <i>PLoS ONE</i> , 2013, 8, e56005.	1.1	13
79	Emulsified Phosphatidylserine, Simple and Effective Peptide Carrier for Induction of Potent Epitope-Specific T Cell Responses. <i>PLoS ONE</i> , 2013, 8, e60068.	1.1	24
80	Mucosal Adjuvanticity of Fibronectin-Binding Peptide (FBP) Fused with <i>Echinococcus multilocularis</i> Tetraspanin 3: Systemic and Local Antibody Responses. <i>PLoS Neglected Tropical Diseases</i> , 2012, 6, e1842.	1.3	6
81	Comparative Genome Analysis of Three Eukaryotic Parasites with Differing Abilities To Transform Leukocytes Reveals Key Mediators of <i>Theileria</i> -Induced Leukocyte Transformation. <i>MBio</i> , 2012, 3, e00204-12.	1.8	64
82	Development of multiple-locus variable-number tandem-repeat analysis for rapid genotyping of <i>Ehrlichia ruminantium</i> and its application to infected <i>Amblyomma variegatum</i> collected in heartwater endemic areas in Uganda. <i>Parasitology</i> , 2012, 139, 69-82.	0.7	5
83	<i>Theileria equi</i> merozoite antigen-2 interacts with actin molecule of equine erythrocyte during their asexual development. <i>Experimental Parasitology</i> , 2012, 132, 508-512.	0.5	5
84	Molecular epidemiological studies on animal trypanosomiasis in Ghana. <i>Parasites and Vectors</i> , 2012, 5, 217.	1.0	32
85	Population genetic analysis and sub-structuring of <i>Theileria parva</i> in the northern and eastern parts of Zambia. <i>Parasites and Vectors</i> , 2012, 5, 255.	1.0	30
86	The use of Loop-mediated Isothermal Amplification (LAMP) to detect the re-emerging Human African Trypanosomiasis (HAT) in the Luangwa and Zambezi valleys. <i>Parasites and Vectors</i> , 2012, 5, 282.	1.0	36
87	Evidence of <i>Yersinia pestis</i> DNA from fleas in an endemic plague area of Zambia. <i>BMC Research Notes</i> , 2012, 5, 72.	0.6	20
88	Characterisation of full-length cDNA sequences provides insights into the <i>Eimeria tenella</i> transcriptome. <i>BMC Genomics</i> , 2012, 13, 21.	1.2	20
89	A Pilot Study on Developing Mucosal Vaccine against Alveolar Echinococcosis (AE) Using Recombinant Tetraspanin 3: Vaccine Efficacy and Immunology. <i>PLoS Neglected Tropical Diseases</i> , 2012, 6, e1570.	1.3	38
90	Construction and analysis of full-length cDNA library of <i>Cryptosporidium parvum</i> . <i>Parasitology International</i> , 2011, 60, 199-202.	0.6	18

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91	Complete coding sequence, sequence analysis and transmembrane topology modelling of <i>Trypanosoma brucei rhodesiense</i> Putative Oligosaccharyl transferase (TbOST II). <i>International Journal of Data Mining and Bioinformatics</i> , 2011, 5, 574.	0.1	0
92	Genotypic Diversity of <i>Theileria orientalis</i> Detected from Cattle Grazing in Kumamoto and Okinawa Prefectures of Japan. <i>Journal of Veterinary Medical Science</i> , 2011, 73, 305-312.	0.3	43
93	Emergence of new types of <i>Theileria orientalis</i> in Australian cattle and possible cause of theileriosis outbreaks. <i>Parasites and Vectors</i> , 2011, 4, 22.	1.0	138
94	Molecular epidemiology of camel trypanosomiasis based on ITS1 rDNA and RoTat 1.2 VSG gene in the Sudan. <i>Parasites and Vectors</i> , 2011, 4, 31.	1.0	65
95	An outbreak of bovine trypanosomiasis in the Blue Nile State, Sudan. <i>Parasites and Vectors</i> , 2011, 4, 74.	1.0	15
96	Identification of genetic loci affecting the establishment and development of <i>Echinococcus multilocularis</i> larvae in mice. <i>International Journal for Parasitology</i> , 2011, 41, 1121-1128.	1.3	10
97	Prevalence of <i>Trypanosoma</i> sp. in cattle from Tanzania estimated by conventional PCR and loop-mediated isothermal amplification (LAMP). <i>Parasitology Research</i> , 2011, 109, 1735-1739.	0.6	35
98	Multi-locus sequence typing of <i>Ehrlichia ruminantium</i> strains from geographically diverse origins and collected in <i>Amblyomma variegatum</i> from Uganda. <i>Parasites and Vectors</i> , 2011, 4, 137.	1.0	17
99	Rapid discrimination and quantification of <i>Theileria orientalis</i> types using ribosomal DNA internal transcribed spacers. <i>Infection, Genetics and Evolution</i> , 2011, 11, 407-414.	1.0	19
100	Full-parasites: database of full-length cDNAs of apicomplexa parasites, 2010 update. <i>Nucleic Acids Research</i> , 2011, 39, D625-D631.	6.5	15
101	Characterization of influenza A viruses isolated from wild waterfowl in Zambia. <i>Journal of General Virology</i> , 2011, 92, 1416-1427.	1.3	23
102	Population Genetics of <i>Trypanosoma evansi</i> from Camel in the Sudan. <i>PLoS Neglected Tropical Diseases</i> , 2011, 5, e1196.	1.3	14
103	Cross-Protective Peptide Vaccine against Influenza A Viruses Developed in HLA-A*2402 Human Immunity Model. <i>PLoS ONE</i> , 2011, 6, e24626.	1.1	42
104	Nucleotide sequence heterogeneity in the small subunit ribosomal RNA gene within <i>Theileria equi</i> from horses in Sudan. <i>Parasitology Research</i> , 2010, 106, 493-498.	0.6	50
105	Characterization of a leucine aminopeptidase from <i>Toxoplasma gondii</i> . <i>Molecular and Biochemical Parasitology</i> , 2010, 170, 1-6.	0.5	41
106	A schizont-derived protein, TpSCOP, is involved in the activation of NF- κ B in <i>Theileria parva</i> -infected lymphocytes. <i>Molecular and Biochemical Parasitology</i> , 2010, 174, 8-17.	0.5	14
107	Development of Loop-Mediated Isothermal Amplification (LAMP) Assays for Rapid Detection of <i>Ehrlichia ruminantium</i> . <i>BMC Microbiology</i> , 2010, 10, 296.	1.3	61
108	Loop-mediated isothermal amplification (LAMP) assays for detection of <i>Theileria parva</i> infections targeting the PIM and p150 genes. <i>International Journal for Parasitology</i> , 2010, 40, 55-61.	1.3	27

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109	Detection of <i>Trypanosoma cruzi</i> and <i>T. rangeli</i> Infections from <i>Rhodnius pallescens</i> Bugs by Loop-Mediated Isothermal Amplification (LAMP). <i>American Journal of Tropical Medicine and Hygiene</i> , 2010, 82, 855-860.	0.6	33
110	High-Resolution Characterization of <i>Toxoplasma gondii</i> Transcriptome with a Massive Parallel Sequencing Method. <i>DNA Research</i> , 2010, 17, 233-243.	1.5	16
111	Serodiagnosis of ovine toxoplasmosis in Mongolia by an enzyme-linked immunosorbent assay with recombinant <i>Toxoplasma gondii</i> matrix antigen 1. <i>Japanese Journal of Veterinary Research</i> , 2010, 58, 111-9.	0.7	10
112	Sequence analysis of <i>Theileria annulata</i> surface protein in Chinese isolates. <i>Zhongguo Ji Sheng Chong Xue Yu Ji Sheng Chong Bing Za Zhi = Chinese Journal of Parasitology & Parasitic Diseases</i> , 2010, 28, 205-9.	0.7	0
113	Molecular cloning and characterization of a T24-like protein in <i>Echinococcus multilocularis</i> . <i>Molecular and Biochemical Parasitology</i> , 2009, 168, 117-119.	0.5	9
114	Characterization of H3N6 avian influenza virus isolated from a wild white pelican in Zambia. <i>Archives of Virology</i> , 2009, 154, 1517-1522.	0.9	12
115	Characterization of a concealed antigen Hq05 from the hard tick <i>Haemaphysalis qinghaiensis</i> and its effect as a vaccine against tick infestation in sheep. <i>Vaccine</i> , 2009, 27, 483-490.	1.7	12
116	Evaluation of <i>Echinococcus multilocularis</i> tetraspanins as vaccine candidates against primary alveolar echinococcosis. <i>Vaccine</i> , 2009, 27, 7339-7345.	1.7	40
117	Overproduction of the Pro-Apoptotic Molecule, Programmed Cell Death 5, in <i>Toxoplasma gondii</i> Leads to Increased Apoptosis of Host Macrophages. <i>Journal of Veterinary Medical Science</i> , 2009, 71, 1183-1189.	0.3	6
118	Stability of Loop-Mediated Isothermal Amplification (LAMP) Reagents and its Amplification Efficiency on Crude <i>Trypanosome</i> DNA Templates. <i>Journal of Veterinary Medical Science</i> , 2009, 71, 471-475.	0.3	69
119	A Field study to Estimate the Prevalence of Bovine African Trypanosomosis in Butaleja District, Uganda. <i>Journal of Veterinary Medical Science</i> , 2009, 71, 525-527.	0.3	6
120	<i>Theileria</i> . , 2009, , 191-231.		10
121	A perspective on <i>Theileria equi</i> infections in donkeys. <i>Japanese Journal of Veterinary Research</i> , 2009, 56, 171-80.	0.7	32
122	Preliminary studies on the effects of orally-administered Transforming Growth Factor-beta on protozoan diseases in mice. <i>Japanese Journal of Veterinary Research</i> , 2009, 57, 101-8.	0.7	3
123	Cloning and characterization of a cDNA clone encoding calreticulin from <i>Haemaphysalis qinghaiensis</i> (Acari: Ixodidae). <i>Parasitology Research</i> , 2008, 102, 737-746.	0.6	37
124	The development and evaluation of a loop-mediated isothermal amplification (LAMP) method for detection of <i>Babesia</i> spp. infective to sheep and goats in China. <i>Experimental Parasitology</i> , 2008, 120, 39-44.	0.5	46
125	Programmed Cell Death 5 from <i>Toxoplasma gondii</i> : A secreted molecule that exerts a pro-apoptotic effect on host cells. <i>Molecular and Biochemical Parasitology</i> , 2008, 159, 112-120.	0.5	10
126	Identification and molecular characterization of a novel stage-specific surface protein of <i>Trypanosoma congolense</i> epimastigotes. <i>Molecular and Biochemical Parasitology</i> , 2008, 161, 1-11.	0.5	27

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127	Cloning and characterization of a cDNA clone encoding troponin T from tick <i>Haemaphysalis qinghaiensis</i> (Acari: Ixodidae). <i>Comparative Biochemistry and Physiology - B Biochemistry and Molecular Biology</i> , 2008, 151, 323-329.	0.7	5
128	Effect of CD4+CD25+ T Cell-Depletion on Acute Lethal Infection of Mice with <i>Trypanosoma congolense</i> . <i>Journal of Veterinary Medical Science</i> , 2008, 70, 751-759.	0.3	5
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