

Nariaki Nonaka

List of Publications by Year in descending order

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Version: 2024-02-01

48
papers

389
citations

840776

11
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16
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48
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48
docs citations

48
times ranked

391
citing authors

#	ARTICLE	IF	CITATIONS
1	Reconstruction of mitochondrial genomes from raw sequencing data provides insights on the phylogeny of Ixodes ticks and cautions for species misidentification. <i>Ticks and Tick-borne Diseases</i> , 2022, 13, 101832.	2.7	4
2	<i>Eimeria pragensis</i> infection alters the gut microenvironment to favor extrinsic shiga toxin-producing <i>Escherichia coli</i> O157:H7 colonization in mice. <i>Parasitology International</i> , 2022, 87, 102521.	1.3	0
3	Sensitivity comparison between Mini-FLOTAC and conventional techniques for the detection of <i>Echinococcus multilocularis</i> eggs. <i>Parasitology International</i> , 2022, 87, 102522.	1.3	2
4	The strong influence of management factors on coccidian infections in smallholder pig farms and the first molecular identification of <i>Cystoisospora suis</i> in Myanmar. <i>Parasite</i> , 2022, 29, 1.	2.0	0
5	Morphological and molecular identification of trematode cercariae related with humans and animal health in freshwater snails from a lake and a dam in Myanmar. <i>Parasitology Research</i> , 2022, 121, 653-665.	1.6	6
6	High infection rate of tick-borne protozoan and rickettsial pathogens of cattle in Malawi and the development of a multiplex PCR for <i>Babesia</i> and <i>Theileria</i> species identification. <i>Acta Tropica</i> , 2022, 231, 106413.	2.0	5
7	Scanning electron microscopy of <i>Quilonia renniei</i> from Asian elephants revealing variation in coronal leaflet number. <i>Parasitology</i> , 2022, 149, 529-533.	1.5	0
8	Genotyping of <i>Theileria parva</i> populations in vaccinated and non-vaccinated cattle in Malawi. <i>Parasitology</i> , 2022, , 1-28.	1.5	0
9	Novel symbionts and potential human pathogens excavated from argasid tick microbiomes that are shaped by dual or single symbiosis. <i>Computational and Structural Biotechnology Journal</i> , 2022, 20, 1979-1992.	4.1	4
10	Molecular detection and characterization of tick-borne hemoparasites and Anaplasmataceae in dogs in major cities of Malawi. <i>Parasitology Research</i> , 2021, 120, 267-276.	1.6	6
11	Metagenomic identification, sequencing, and genome analysis of porcine hepe-astroviruses (bastroviruses) in porcine feces in Japan. <i>Infection, Genetics and Evolution</i> , 2021, 88, 104664.	2.3	2
12	Molecular identification and genetic characterization of tick-borne pathogens in sheep and goats at two farms in the central and southern regions of Malawi. <i>Ticks and Tick-borne Diseases</i> , 2021, 12, 101629.	2.7	10
13	<i>Amblyomma testudinarium</i> infestation on a brown bear (<i>Ursus arctos yesoensis</i>) captured in Hokkaido, a northern island of Japan. <i>Parasitology International</i> , 2021, 80, 102209.	1.3	9
14	Anthropogenic interferences lead to gut microbiome dysbiosis in Asian elephants and may alter adaptation processes to surrounding environments. <i>Scientific Reports</i> , 2021, 11, 741.	3.3	24
15	Genetic diversity and population structure analyses based on microsatellite DNA of parthenogenetic <i>Fasciola</i> flukes obtained from cattle and sika deer in Japan. <i>Parasitology Research</i> , 2021, 120, 1341-1350.	1.6	6
16	<i>Spiroplasma</i> Infection among Ixodid Ticks Exhibits Species Dependence and Suggests a Vertical Pattern of Transmission. <i>Microorganisms</i> , 2021, 9, 333.	3.6	5
17	Exploring Prokaryotic and Eukaryotic Microbiomes Helps in Detecting Tick-Borne Infectious Agents in the Blood of Camels. <i>Pathogens</i> , 2021, 10, 351.	2.8	16
18	Applications of Blocker Nucleic Acids and Non-Metazoan PCR Improves the Discovery of the Eukaryotic Microbiome in Ticks. <i>Microorganisms</i> , 2021, 9, 1051.	3.6	2

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19	Molecular Detection and Genotyping of Coxiella-Like Endosymbionts in Ticks Collected from Animals and Vegetation in Zambia. <i>Pathogens</i> , 2021, 10, 779.	2.8	6
20	Identification, genetic variation, and structural analysis of 18S rRNA of <i>Theileria orientalis</i> and <i>Theileria velifera</i> -like isolates from Myanmar. <i>Parasitology International</i> , 2021, 82, 102299.	1.3	5
21	PCR detection and genetic characterization of piroplasms from dogs in Myanmar, and a possible role of dogs as reservoirs for <i>Theileria</i> parasites infecting cattle, water buffaloes, and goats. <i>Ticks and Tick-borne Diseases</i> , 2021, 12, 101729.	2.7	7
22	Molecular Survey of Babesia and Anaplasma Infection in Cattle in Bolivia. <i>Veterinary Sciences</i> , 2021, 8, 188.	1.7	4
23	<i>Ancylostoma ceylanicum</i> infections in humans in Vietnam. <i>Parasitology International</i> , 2021, 84, 102405.	1.3	8
24	Early-phase migration dynamics of <i>Echinococcus multilocularis</i> in two mouse strains showing different infection susceptibilities. <i>International Journal for Parasitology</i> , 2021, 51, 893-898.	3.1	1
25	Application of a real-time PCR assay for the detection of <i>Ascaris suum</i> DNA in the liver of experimentally infected chickens. <i>Journal of Veterinary Medical Science</i> , 2021, 83, 671-674.	0.9	0
26	Cystic echinococcosis in humans and animals in Egypt: An epidemiological overview. <i>Current Research in Parasitology and Vector-borne Diseases</i> , 2021, 1, 100061.	1.9	9
27	Establishment of a serodiagnosis system for the detection of <i>Toxocara</i> spp. and <i>Ascaris suum</i> infection in chickens. <i>Parasitology International</i> , 2020, 75, 102022.	1.3	2
28	First record and analysis of the COI gene of <i>Cobboldia elephantis</i> obtained from a captive Asian elephant from Myanmar. <i>Parasitology International</i> , 2020, 75, 102035.	1.3	2
29	Detection and molecular identification of <i>Leucocytozoon</i> and <i>Plasmodium</i> species from village chickens in different areas of Myanmar. <i>Acta Tropica</i> , 2020, 212, 105719.	2.0	17
30	Genetic Diversity and Sequence Polymorphism of Two Genes Encoding <i>Theileria parva</i> Antigens Recognized by CD8+ T Cells among Vaccinated and Unvaccinated Cattle in Malawi. <i>Pathogens</i> , 2020, 9, 334.	2.8	9
31	Morphological and molecular identification of cyathostomine gastrointestinal nematodes of <i>Murshidia</i> and <i>Quilonia</i> species from Asian elephants in Myanmar. <i>International Journal for Parasitology: Parasites and Wildlife</i> , 2020, 11, 294-301.	1.5	4
32	Adult worm exclusion and histological data of dogs repeatedly infected with the cestode <i>Echinococcus multilocularis</i> . <i>Data in Brief</i> , 2020, 29, 105353.	1.0	1
33	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, .	0.6	3
34	First molecular identification of <i>Strongyloides vituli</i> in cattle in Japan and insights into the evolutionary history of <i>Strongyloides</i> parasites of ruminants. <i>Parasitology International</i> , 2019, 72, 101937.	1.3	9
35	Evaluation of the natural vertical transmission of <i>Theileria orientalis</i> . <i>Veterinary Parasitology</i> , 2018, 263, 1-4.	1.8	22
36	Avian Filariasis in Backyard Chickens in Japan. <i>Avian Diseases</i> , 2018, 62, 326-329.	1.0	6

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37	Development of nested multiplex polymerase chain reaction (PCR) assay for the detection of <i>Toxocara canis</i> , <i>Toxocara cati</i> and <i>Ascaris suum</i> contamination in meat and organ meats. <i>Parasitology International</i> , 2018, 67, 622-626.	1.3	14
38	Diagnosis of canine <i>Echinococcus multilocularis</i> infections by copro-DNA tests: comparison of DNA extraction techniques and evaluation of diagnostic deworming. <i>Parasitology Research</i> , 2017, 116, 2139-2144.	1.6	9
39	Molecular and morphological variation of <i>Paragonimus westermani</i> in Vietnam with records of new second intermediate crab hosts and a new locality in a northern province. <i>Parasitology</i> , 2016, 143, 1639-1646.	1.5	14
40	<i>Echinococcus multilocularis</i> Infection in Pet Dogs in Japan. <i>Vector-Borne and Zoonotic Diseases</i> , 2009, 9, 201-206.	1.5	23
41	Multiplex PCR system for identifying the carnivore origins of faeces for an epidemiological study on <i>Echinococcus multilocularis</i> in Hokkaido, Japan. <i>Parasitology Research</i> , 2009, 106, 75-83.	1.6	31
42	A Vague Understanding of the Biology and Epidemiology of Echinococcosis by Dog Owners in Hokkaido, an Endemic Island for <i>Echinococcus multilocularis</i> in Japan. <i>Journal of Veterinary Medical Science</i> , 2009, 71, 105-107.	0.9	2
43	A latex agglutination test for the detection of <i>Echinococcus multilocularis</i> coproantigen in the definitive hosts. <i>Veterinary Parasitology</i> , 2008, 152, 278-283.	1.8	8
44	The first instance of a cat excreting <i>Echinococcus multilocularis</i> eggs in Japan. <i>Parasitology International</i> , 2008, 57, 519-520.	1.3	18
45	Towards the control of <i>Echinococcus multilocularis</i> in the definitive host in Japan. <i>Parasitology International</i> , 2006, 55, S263-S266.	1.3	19
46	Dose dependency of prednisolone tertiary-butylacetate (PTBA) treatment on the establishment and site predilection of <i>Echinococcus multilocularis</i> in an alternative definitive host model using Mongolian gerbil (<i>Meriones unguiculatus</i>). <i>Parasitology Research</i> , 2000, 86, 521-523.	1.6	2
47	Hyperplasia of gastric mucosa in donor rats orally infected with <i>Taenia taeniaeformis</i> eggs and in recipient rats surgically implanted with the larvae in the abdominal cavity. <i>Parasitology Research</i> , 1999, 85, 431-436.	1.6	10
48	Coproantigen Detection in a Survey of <i>Echinococcus multilocularis</i> Infection among Red Foxes, <i>Vulpes vulpes schrencki</i> , in Hokkaido, Japan.. <i>Journal of Veterinary Medical Science</i> , 1998, 60, 639-641.	0.9	23