

Tadashi Itagaki

List of Publications by Year in descending order

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71
papers

1,686
citations

304743

22
h-index

302126

39
g-index

71
all docs

71
docs citations

71
times ranked

810
citing authors

#	ARTICLE	IF	CITATIONS
1	The causative agents of fascioliasis in animals and humans: Parthenogenetic <i>Fasciola</i> in Asia and other regions. <i>Infection, Genetics and Evolution</i> , 2022, 99, 105248.	2.3	6
2	Molecular analyses confirm the coexistence of <i>Fasciola gigantica</i> and parthenogenetic <i>Fasciola</i> in the Philippines. <i>Parasitology International</i> , 2022, 88, 102562.	1.3	3
3	Molecular characterization of <i>Oxyspirura mansoni</i> and <i>Philophthalmus gralli</i> collected from the eyes of domestic chickens in Bangladesh. <i>Parasitology International</i> , 2021, 80, 102243.	1.3	2
4	Molecular characterization revealed <i>Fasciola</i> specimens in Ecuador are all <i>Fasciola hepatica</i> , none of all of <i>Fasciola gigantica</i> or parthenogenetic <i>Fasciola</i> species. <i>Parasitology International</i> , 2021, 80, 102215.	1.3	9
5	Prevalence of <i>Trichinella</i> T9 in Japanese black bears (<i>Ursus thibetanus japonicus</i>) in Iwate prefecture, Japan. <i>Parasitology International</i> , 2021, 80, 102217.	1.3	4
6	Do aspermic (parthenogenetic) <i>Fasciola</i> forms have the ability to reproduce their progeny via parthenogenesis?. <i>Journal of Helminthology</i> , 2021, 95, e36.	1.0	3
7	Molecular characterization of <i>Ascaridia galli</i> from Bangladesh and development of a PCR method for distinguishing <i>A. galli</i> from <i>Heterakis</i> spp.. <i>Journal of Veterinary Medical Science</i> , 2021, 83, 666-670.	0.9	2
8	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
9	Development of a multiplex PCR method for discriminating between <i>Heterakis gallinarum</i> , <i>H. beramporia</i> , and <i>H. indica</i> parasites of poultry. <i>Veterinary Parasitology</i> , 2021, 295, 109463.	1.8	2
10	Prevalence of <i>Nosema</i> species infections in <i>Apis cerana japonica</i> and <i>Apis mellifera</i> honeybees in the Tohoku region of Japan. <i>Parasitology International</i> , 2021, 83, 102361.	1.3	1
11	Detection and molecular characteristics of <i>Rhytidodoides</i> sp. (Digenea: Rhytidodidae) from the gall bladder of green sea turtles (<i>Chelonia mydas</i>) in the Ogasawara Islands, Japan. <i>Parasitology International</i> , 2021, 83, 102377.	1.3	0
12	Detection and molecular characteristics of <i>Pyelosomum cochlear</i> (Digenea: Pronocephalidae) in the urinary bladder of the green sea turtle (<i>Chelonia mydas</i>) in the Northwest Pacific Ocean. <i>Infection, Genetics and Evolution</i> , 2021, 93, 104962.	2.3	0
13	Development of conventional multiplex PCR method for discrimination between <i>Dispharynx nasuta</i> and <i>Cheilospirura hamulosa</i> (Nematoda: Acuariidae) parasitizing poultry. <i>Journal of Veterinary Medical Science</i> , 2021, 83, 226-229.	0.9	0
14	A rebuttal letter to Letter to the Editor by P. Heneberg on "Taxonomic comments on the validity of <i>Echinostoma miyagawai</i> Ishii, 1932 (Trematoda: Echinostomatidae)". <i>Parasitology International</i> , 2020, 74, 101971.	1.3	1
15	The prevalence and molecular characterization of <i>Acarapis woodi</i> and <i>Varroa destructor</i> mites in honeybees in the Tohoku region of Japan. <i>Parasitology International</i> , 2020, 75, 102052.	1.3	2
16	Multigene typing and phylogenetic analysis of <i>Fasciola</i> from endemic foci in Iran. <i>Infection, Genetics and Evolution</i> , 2020, 80, 104202.	2.3	5
17	Phylogenetic relationships between Lymnaeidae in relation to infection with <i>Fasciola</i> sp. in Hokkaido, Japan. <i>Molluscan Research</i> , 2020, 40, 160-168.	0.7	6
18	First report of <i>Paragonimus skrjabini miyazakii</i> metacercariae in <i>Geothelphusa dehaani</i> (Sawagani) occurring in Iwate Prefecture, Japan. <i>Journal of Veterinary Medical Science</i> , 2019, 81, 1109-1112.	0.9	3

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19	Characterization of <i>Echinostoma revolutum</i> and <i>Echinostoma robustum</i> from ducks in Bangladesh based on morphology, nuclear ribosomal ITS2 and mitochondrial nad1 sequences. <i>Parasitology International</i> , 2019, 69, 1-7.	1.3	21
20	Morphological and genetic characterization of green-banded broodsacs of <i>Leucochloridium</i> (<i>Leucochloridiidae</i> : Trematoda) sporocysts detected in <i>Succinea lauta</i> in Hokkaido, Japan. <i>Parasitology International</i> , 2019, 68, 53-56.	1.3	2
21	Infection by and Molecular Features of <i>Learedius learedi</i> (Digenea: Schistosomatoidea) in Green Sea Turtles (<i>Chelonia mydas</i>) on the Ogasawara Islands, Japan. <i>Journal of Parasitology</i> , 2019, 105, 533.	0.7	4
22	Infection by and Molecular Features of (Digenea: Schistosomatoidea) in Green Sea Turtles () on the Ogasawara Islands, Japan. <i>Journal of Parasitology</i> , 2019, 105, 533-538.	0.7	1
23	Hybrid origin of Asian aspermic &Fasciola& flukes is confirmed by analyzing two single-copy genes, &pepck& and &pold&. <i>Journal of Veterinary Medical Science</i> , 2018, 80, 98-102.	0.9	21
24	Molecular characterization and phylogenetic analysis of <i>Fasciola gigantica</i> from Nigeria. <i>Parasitology International</i> , 2017, 66, 893-897.	1.3	25
25	Gastrointestinal pseudoparasitism by chestnut weevil (<i>Curculio sikkimensis</i>) larvae in a dog. <i>Journal of Small Animal Practice</i> , 2017, 58, 302-302.	1.2	0
26	Nuclear and mitochondrial DNA analysis reveals that hybridization between <i>Fasciola hepatica</i> and <i>Fasciola gigantica</i> occurred in China. <i>Parasitology</i> , 2017, 144, 206-213.	1.5	40
27	Molecular and phylogenetic analyses of the liver amphistome <i>Explanatum explanatum</i> (Creplin, 1847) Fukui, 1929 in ruminants from Bangladesh and Nepal based on nuclear ribosomal ITS2 and mitochondrial nad1 sequences. <i>Journal of Helminthology</i> , 2017, 91, 497-503.	1.0	7
28	Phylogenetic relationships between <i>Dicrocoelium chinensis</i> populations in Japan and China based on mitochondrial nad1 gene sequences. <i>Parasitology Research</i> , 2017, 116, 2605-2609.	1.6	9
29	First report of <i>Fasciola</i> larva infection in <i>Galba truncatula</i> (Müller, 1774) (Gastropoda,) Tj ETQq1 1 0.784314 rgBT /Over Science, 2017, 79, 1381-1383.	0.9	7
30	Molecular characterization and phylogenetic analysis of <i>Explanatum explanatum</i> in India based on nucleotide sequences of ribosomal ITS2 and the mitochondrial gene <i>nad1</i>. <i>Journal of Veterinary Medical Science</i> , 2016, 78, 1745-1748.	0.9	2
31	Molecular characterization of &Fasciola gigantica& in Delhi, India and its phylogenetic relation to the species from South Asian countries. <i>Journal of Veterinary Medical Science</i> , 2016, 78, 1529-1532.	0.9	17
32	Novel methods for the molecular discrimination of <i>Fasciola</i> spp. on the basis of nuclear protein-coding genes. <i>Parasitology International</i> , 2016, 65, 180-183.	1.3	50
33	Molecular characterization and phylogenetic analysis of <i>Fasciola hepatica</i> from Peru. <i>Parasitology International</i> , 2016, 65, 171-174.	1.3	21
34	Reappraisal of <i>Hydatigera taeniaeformis</i> (Batsch, 1786) (Cestoda: Taeniidae) sensu lato with description of <i>Hydatigera kamiyai</i> n. sp.. <i>International Journal for Parasitology</i> , 2016, 46, 361-374.	3.1	40
35	Sequence differences in the internal transcribed spacer 1 and 5.8S ribosomal RNA among three &Moniezia& species isolated from ruminants in Japan. <i>Journal of Veterinary Medical Science</i> , 2015, 77, 105-107.	0.9	13
36	Molecular phylogenetic analysis of <i>Fasciola</i> flukes from eastern India. <i>Parasitology International</i> , 2015, 64, 334-338.	1.3	34

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37	Protection against <i>Fasciola gigantica</i> infection in mice by vaccination with recombinant juvenile-specific cathepsin L. <i>Vaccine</i> , 2015, 33, 1596-1601.	3.8	18
38	Morphological and molecular characterization of <i>Eurytrema cladorchis</i> parasitizing cattle (<i>Bos</i>) in Thailand. <i>Journal of Parasitology</i> , 2010, 100, 107-114.	1.6	10
39	Vaccine potential of recombinant pro- and mature cathepsinL1 against fasciolosis <i>gigantica</i> in mice. <i>Acta Tropica</i> , 2015, 150, 71-78.	2.0	10
40	Molecular characterization of <i>Cryptosporidium parvum</i> from two different Japanese prefectures, Okinawa and Hokkaido. <i>Parasitology International</i> , 2015, 64, 161-166.	1.3	19
41	Molecular detection and characterization of <i>Cryptosporidium</i> species in household dogs, pet shop puppies, and dogs kept in a school of veterinary nursing in Japan. <i>Veterinary Parasitology</i> , 2014, 200, 284-288.	1.8	14
42	Molecular phylogenetic identification of <i>Fasciola</i> flukes in Nepal. <i>Parasitology International</i> , 2014, 63, 758-762.	1.3	33
43	Characteristics and molecular phylogeny of <i>Fasciola</i> flukes from Bangladesh, determined based on spermatogenesis and nuclear and mitochondrial DNA analyses. <i>Parasitology Research</i> , 2014, 113, 2493-2501.	1.6	52
44	Distinct Distribution of <i>Dicrocoelium dendriticum</i> and <i>D. chinensis</i> in Iwate Prefecture, Japan, and a New Final Host Record for <i>D. chinensis</i> . <i>Journal of Veterinary Medical Science</i> , 2014, 76, 1415-1417.	0.9	8
45	Molecular identification of <i>Fasciola</i> spp. (Digenea: Platyhelminthes) in cattle from Vietnam. <i>Parasite</i> , 2012, 19, 85-89.	2.0	23
46	Molecular Analysis of Aspermic <i>Fasciola</i> Flukes from Korea on the Basis of the Nuclear ITS1 Region and Mitochondrial DNA Markers and Comparison with Japanese Aspermic <i>Fasciola</i> Flukes. <i>Journal of Veterinary Medical Science</i> , 2012, 74, 899-904.	0.9	20
47	Identification of <i>Fasciola</i> flukes in Thailand based on their spermatogenesis and nuclear ribosomal DNA, and their intraspecific relationships based on mitochondrial DNA. <i>Parasitology International</i> , 2012, 61, 545-549.	1.3	37
48	Characterization of <i>Fasciola</i> spp. in Myanmar on the basis of spermatogenesis status and nuclear and mitochondrial DNA markers. <i>Parasitology International</i> , 2011, 60, 474-479.	1.3	39
49	Prevalence of intestinal parasites and genotyping of <i>Giardia intestinalis</i> in pet shop puppies in east Japan. <i>Veterinary Parasitology</i> , 2011, 176, 74-78.	1.8	23
50	Hybridization experiments indicate incomplete reproductive isolating mechanism between <i>Fasciola hepatica</i> and <i>Fasciola gigantica</i> . <i>Parasitology</i> , 2011, 138, 1278-1284.	1.5	26
51	Identification of <i>Fasciola</i> species isolated from Egypt based on sequence analysis of genomic (ITS1 and) <i>Tj ETQq1 1_0,784314,rgBT /Ove</i>	1.3	77
52	DNA Types of Aspermic <i>Fasciola</i> Species in Japan. <i>Journal of Veterinary Medical Science</i> , 2010, 72, 1371-1374.	0.9	27
53	Discrimination of the ITS1 types of <i>Fasciola</i> spp. based on a PCR-RFLP method. <i>Parasitology Research</i> , 2010, 106, 757-761.	1.6	77
54	Molecular characterization of <i>Fasciola hepatica</i> , <i>Fasciola gigantica</i> , and aspermic <i>Fasciola</i> sp. in China based on nuclear and mitochondrial DNA. <i>Parasitology Research</i> , 2009, 105, 809-815.	1.6	89

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55	Occurrence of spermic diploid and aspermic triploid forms of <i>Fasciola</i> in Vietnam and their molecular characterization based on nuclear and mitochondrial DNA. <i>Parasitology International</i> , 2009, 58, 81-85.	1.3	83
56	Prevalence of <i>Giardia intestinalis</i> Infection in Household Cats of Tohoku District in Japan. <i>Journal of Veterinary Medical Science</i> , 2006, 68, 161-163.	0.9	14
57	Molecular Characterization of Parthenogenic <i>Fasciola</i> sp. in Korea on the Basis of DNA Sequences of Ribosomal ITS1 and Mitochondrial NDI Gene. <i>Journal of Veterinary Medical Science</i> , 2005, 67, 1115-1118.	0.9	94
58	Genotyping of <i>Giardia intestinalis</i> from domestic and wild animals in Japan using glutamate dehydrogenase gene sequencing. <i>Veterinary Parasitology</i> , 2005, 133, 283-287.	1.8	82
59	Genetic characterization of parthenogenic <i>Fasciola</i> sp. in Japan on the basis of the sequences of ribosomal and mitochondrial DNA. <i>Parasitology</i> , 2005, 131, 679-685.	1.5	153
60	Detection of <i>Giardia</i> Antigen in Puppies Using Enzyme-linked Immunosorbent Assay. <i>Nippon Juishikai Zasshi Journal of the Japan Veterinary Medical Association</i> , 2004, 57, 579-582.	0.1	4
61	Discrimination of Three Amphistome Species by PCR-RFLP Based on rDNA ITS2 Markers. <i>Journal of Veterinary Medical Science</i> , 2003, 65, 931-933.	0.9	57
62	An Investigation of Heavy Metal Exposure and Risks to Wildlife in the Kafue Flats of Zambia.. <i>Journal of Veterinary Medical Science</i> , 2001, 63, 315-318.	0.9	13
63	Comparative Study of the Reproductive Organs of <i>Fasciola</i> Groups by Optical Microscope.. <i>Journal of Veterinary Medical Science</i> , 2001, 63, 735-742.	0.9	25
64	Mitochondrial DNA polymorphism of a triploid form of <i>Fasciola</i> in Japan. <i>Journal of Helminthology</i> , 2001, 75, 193-6.	1.0	3
65	Triploid form of <i>Fasciola</i> in Japan: genetic relationships between <i>Fasciola hepatica</i> and <i>Fasciola gigantica</i> determined by ITS-2 sequence of nuclear rDNA. <i>International Journal for Parasitology</i> , 1998, 28, 777-781.	3.1	107
66	Taxonomic Status of the Japanese Triploid Forms of <i>Fasciola</i> : Comparison of Mitochondrial ND1 and COI Sequences with <i>F. hepatica</i> and <i>F. gigantica</i> . <i>Journal of Parasitology</i> , 1998, 84, 445.	0.7	43
67	Taxonomic status of the Japanese triploid forms of <i>Fasciola</i> : comparison of mitochondrial ND1 and COI sequences with <i>F. hepatica</i> and <i>F. gigantica</i> . <i>Journal of Parasitology</i> , 1998, 84, 445-8.	0.7	9
68	Infectivity of Three Species of <i>Fasciola</i> to Wistar Rats.. <i>Journal of Veterinary Medical Science</i> , 1994, 56, 977-979.	0.9	10
69	Susceptibility of Japanese <i>Lymnaea ollula</i> to Nigerian <i>Fasciola gigantica</i> .. <i>Nihon Juigaku Zasshi</i> , 1989, 51, 853-854.	0.3	4
70	Experimental infection of Japanese <i>Lymnaea</i> snails with Australian <i>Fasciola hepatica</i> .. <i>Nihon Juigaku Zasshi</i> , 1988, 50, 1085-1091.	0.3	9
71	The snail host of <i>Fasciola</i> sp. in the Tempoku district of Hokkaido.. <i>Nihon Juigaku Zasshi</i> , 1986, 48, 323-328.	0.3	5