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List of Publications by Year in descending order

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Version: 2024-02-01

37
papers

644
citations

687220

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610775

24
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39
all docs

39
docs citations

39
times ranked

631
citing authors

#	ARTICLE	IF	CITATIONS
1	Geography and host specificity: Two forces behind the genetic structure of the freshwater fish parasite <i>Ligula intestinalis</i> (Cestoda: Diphyllbothriidae). <i>International Journal for Parasitology</i> , 2008, 38, 1465-1479.	1.3	61
2	Population Bottlenecks during the Infectious Cycle of the Lyme Disease Spirochete <i>Borrelia burgdorferi</i> . <i>PLoS ONE</i> , 2014, 9, e101009.	1.1	60
3	A hitchhikers guide to the Galápagos: co-phylogeography of Galápagos mockingbirds and their parasites. <i>BMC Evolutionary Biology</i> , 2011, 11, 284.	3.2	57
4	Multiple origins of European populations of the giant liver fluke <i>Fascioloides magna</i> (Trematoda: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 6	1.3	52
5	Interplay of host specificity and biogeography in the population structure of a cosmopolitan endoparasite: microsatellite study of <i>Ligula intestinalis</i> (Cestoda). <i>Molecular Ecology</i> , 2009, 18, 1187-1206.	2.0	51
6	Intra-individual internal transcribed spacer 1 (ITS1) and ITS2 ribosomal sequence variation linked with multiple rDNA loci: A case of triploid <i>Atractolytocestus huronensis</i> , the monozoic cestode of common carp. <i>International Journal for Parasitology</i> , 2010, 40, 175-181.	1.3	43
7	The phylogeny of diphyllbothriid tapeworms (Cestoda: Pseudophyllidea) based on ITS-2 rDNA sequences. <i>Parasitology Research</i> , 2004, 94, 10-15.	0.6	32
8	Host generalists and specialists emerging side by side: an analysis of evolutionary patterns in the cosmopolitan chewing louse genus <i>Menacanthus</i> . <i>International Journal for Parasitology</i> , 2015, 45, 63-73.	1.3	27
9	Host specificity driving genetic structure and diversity in ectoparasite populations: Coevolutionary patterns in <i>Apodemus</i> mice and their lice. <i>Ecology and Evolution</i> , 2018, 8, 10008-10022.	0.8	27
10	Population study of <i>Atractolytocestus huronensis</i> (Cestoda: Caryophyllidea), an invasive parasite of common carp introduced to Europe: mitochondrial cox1 haplotypes and intragenomic ribosomal ITS2 variants. <i>Parasitology Research</i> , 2011, 109, 125-131.	0.6	25
11	Host specificity and genealogy of the louse <i>Polyplax serrata</i> on field mice, <i>Apodemus</i> species: A case of parasite duplication or colonisation?. <i>International Journal for Parasitology</i> , 2008, 38, 731-741.	1.3	21
12	Balancing selection and genetic drift create unusual patterns of $MHCII$ variation in Galápagos mockingbirds. <i>Molecular Ecology</i> , 2016, 25, 4757-4772.	2.0	17
13	Molecular characterization of <i>Atractolytocestus sagittatus</i> (Cestoda: Caryophyllidea), monozoic parasite of common carp, and its differentiation from the invasive species <i>Atractolytocestus huronensis</i> . <i>Parasitology Research</i> , 2012, 110, 1621-1629.	0.6	14
14	The tapeworm <i>Atractolytocestus tenuicollis</i> (Cestoda: Caryophyllidea) a sister species or ancestor of an invasive <i>A. huronensis</i> ?. <i>Parasitology Research</i> , 2013, 112, 3379-3388.	0.6	13
15	Genetic interrelationships of North American populations of giant liver fluke <i>Fascioloides magna</i> . <i>Parasites and Vectors</i> , 2015, 8, 288.	1.0	13
16	Population structure and dispersal routes of an invasive parasite, <i>Fascioloides magna</i> , in North America and Europe. <i>Parasites and Vectors</i> , 2016, 9, 547.	1.0	12
17	Development and characterization of multiplex panels of polymorphic microsatellite loci in giant liver fluke <i>Fascioloides magna</i> (Trematoda: Fasciolidae), using next-generation sequencing approach. <i>Molecular and Biochemical Parasitology</i> , 2014, 195, 30-33.	0.5	11
18	Distinct haplotype structure at the innate immune receptor Toll-like receptor 2 across bank vole populations and lineages in Europe. <i>Biological Journal of the Linnean Society</i> , 2015, 116, 124-133.	0.7	10

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19	Development of microsatellite loci in zoonotic tapeworm <i>Dibothriocephalus latus</i> (Linnaeus, 1758), L1/4he, 1899 (syn. <i>Diphyllobothrium latum</i>) using microsatellite library screening. <i>Molecular and Biochemical Parasitology</i> , 2018, 225, 1-3.	0.5	10
20	Early evidence of establishment of the tropical bedbug (<i>Cimex hemipterus</i>) in Central Europe. <i>Medical and Veterinary Entomology</i> , 2021, 35, 462-467.	0.7	10
21	Pathways of cryptic invasion in a fish parasite traced using coalescent analysis and epidemiological survey. <i>Biological Invasions</i> , 2013, 15, 1907-1923.	1.2	9
22	Isolation and characterization of microsatellite loci in the tapeworm <i>Ligula intestinalis</i> (Cestoda:) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 6	1.7	8
23	Population co-divergence in common cuttlefish (<i>Sepia officinalis</i>) and its dicyemid parasite in the Mediterranean Sea. <i>Scientific Reports</i> , 2019, 9, 14300.	1.6	8
24	Phylogeography of the parasitic mite <i>Laelaps agilis</i> in Western Palearctic shows lineages lacking host specificity but possessing different demographic histories. <i>BMC Zoology</i> , 2022, 7, .	0.3	8
25	��Parasite turnover zone� at secondary contact: A new pattern in host� parasite population genetics. <i>Molecular Ecology</i> , 2020, 29, 4653-4664.	2.0	7
26	Different phylogenomic methods support monophyly of enigmatic � Mesozoa� (Dicyemida +) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 6	1.2	7
27	From taxonomic deflation to newly detected cryptic species: Hidden diversity in a widespread African squeaker catfish. <i>Scientific Reports</i> , 2019, 9, 15748.	1.6	6
28	Tour around the globe: The case of invasive tapeworm <i>Atractolytocestus huronensis</i> (Cestoda:) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 38	0.6	5
29	Unique genetic structure of the human tapeworm <i>Dibothriocephalus latus</i> from the Alpine lakes region � a successful adaptation?. <i>Parasitology</i> , 2022, 149, 1106-1118.	0.7	5
30	Development of polymorphic microsatellites for the invasive Asian fish tapeworm <i>Schyzocotyle acheilognathi</i> . <i>Parasitology International</i> , 2018, 67, 341-343.	0.6	4
31	Discordant population histories of host and its parasite: A role for ecological permeability of extreme environment?. <i>PLoS ONE</i> , 2017, 12, e0175286.	1.1	3
32	Characterisation of microsatellite loci in two species of lice, <i>Polyplax serrata</i> (Phthiraptera:) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 227 Td	0.7	3
33	<i>Cimex lectularius</i> and <i>Cimex hemipterus</i> (bed bugs). <i>Trends in Parasitology</i> , 2022, 38, 919-920.	1.5	2
34	Comparative analysis of monozoic fish tapeworms <i>Caryophyllaeus laticeps</i> (Pallas, 1781) and recently described <i>Caryophyllaeus chondrostomi</i> Bar�k, Oros, Hanzelov�, Scholz, 2017, using microsatellite markers. <i>Parasitology Research</i> , 2020, 119, 3995-4004.	0.6	1
35	Association between louse abundance and MHC II supertypes in Gal�pagos mockingbirds. <i>Parasitology Research</i> , 2020, 119, 1597-1605.	0.6	1
36	Comparative phylogeography of two bat species and their mites in Iran shows impact of host sociality and vagility on population structure. <i>Journal of Zoological Systematics and Evolutionary Research</i> , 2021, 59, 1557-1582.	0.6	1

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37	Genetic analysis of dicyemid intrapopulations suggests sexual reproduction and host colonization by multiple individuals is common. <i>Organisms Diversity and Evolution</i> , 2021, 21, 437.	0.7	0