

Eva ĀEisovskĀ; BazsalovicsovĀ;

List of Publications by Year in descending order

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#	ARTICLE	IF	CITATIONS
1	Genetic interrelationships of <i>Spirometra erinaceieuropaei</i> (Cestoda: Diphyllbothriidea), the causative agent of sparganosis in Europe. <i>Parasite</i> , 2022, 29, 8.	2.0	5
2	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.	1.5	5
3	The first records of <i>Spirometra erinaceieuropaei</i> (Cestoda: Diphyllbothriidae), a causative agent of human sparganosis, in Latvian wildlife. <i>Parasitology Research</i> , 2021, 120, 365-371.	1.6	11
4	Ups and downs of infections with the broad fish tapeworm <i>Dibothriocephalus latus</i> in Europe from 1900 to 2020: Part I. <i>Advances in Parasitology</i> , 2021, 114, 75-166.	3.2	7
5	Development of 14 Microsatellite Markers for Zoonotic Tapeworm <i>Dibothriocephalus dendriticus</i> (Cestoda: Diphyllbothriidea). <i>Genes</i> , 2020, 11, 782.	2.4	5
6	Comparative analysis of monozoic fish tapeworms <i>Caryophyllaeus laticeps</i> (Pallas, 1781) and recently described <i>Caryophyllaeus chondrostomi</i> BarÅk, Oros, HanzelovÅi, Scholz, 2017, using microsatellite markers. <i>Parasitology Research</i> , 2020, 119, 3995-4004.	1.6	1
7	Host Switching of Zoonotic Broad Fish Tapeworm (<i>Dibothriocephalus latus</i>) to Salmonids, Patagonia. <i>Emerging Infectious Diseases</i> , 2019, 25, 2156-2158.	4.3	15
8	Endohelminths of European Perch (<i>Perca fluviatilis</i>) from Selected Localities in Poland with an Emphasis on Search of the Broad Fish Tapeworm <i>Dibothriocephalus latus</i> . <i>Acta Parasitologica</i> , 2019, 64, 544-550.	1.1	3
9	Results on search for the broad fish tapeworm <i>Dibothriocephalus latus</i> (Linnaeus, 1758), (syn.) Tj ETQq1 1 0.784314 rgBT /Overlock 10 256-260.	0.9	4
10	A study of the endohelminths of the European perch <i>Perca fluviatilis</i> L. from the central region of the Danube river basin in Slovakia. <i>ZooKeys</i> , 2019, 899, 47-58.	1.1	6
11	Tour around the globe: The case of invasive tapeworm <i>Atractolytocestus huronensis</i> (Cestoda:) Tj ETQq1 1 0.784314 rgBT /Overlock 10 1.3 5	1.3	5
12	Development of microsatellite loci in zoonotic tapeworm <i>Dibothriocephalus latus</i> (Linnaeus, 1758), LÅ¼he, 1899 (syn. <i>Diphyllbothrium latum</i>) using microsatellite library screening. <i>Molecular and Biochemical Parasitology</i> , 2018, 225, 1-3.	1.1	10
13	Mitochondrial genotyping of <i>Fascioloides magna</i> from Bavaria, Germany. <i>Acta Parasitologica</i> , 2017, 62, 870-874.	1.1	3
14	Transmission risk assessment of invasive fluke <i>Fascioloides magna</i> using GIS-modelling and multicriteria analysis methods. <i>Helminthologia</i> , 2017, 54, 119-131.	0.9	4
15	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.	2.5	12
16	A long-term survey of <i>Fascioloides magna</i> in red deer (<i>Cervus elaphus</i>) in Slovakia (Danube floodplain) Tj ETQq0 0 0 rgBT /Overlock 10 0.9 4	0.9	4
17	A genetic structure of novel population of <i>Fascioloides magna</i> from Poland, Podkarpackie Province, indicates an expanding second European natural focus of fascioloidosis. <i>Acta Parasitologica</i> , 2016, 61, 790-795.	1.1	5
18	The Giant Liver Fluke, <i>Fascioloides magna</i> : Past, Present and Future Research. <i>Springer Briefs in Animal Sciences</i> , 2016, , .	0.1	18

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19	Distribution of <i>Fascioloides magna</i> . Springer Briefs in Animal Sciences, 2016, , 17-40.	0.1	1
20	Modern Approaches in <i>Fascioloides magna</i> Studies. Springer Briefs in Animal Sciences, 2016, , 81-104.	0.1	0
21	Genetic interrelationships of North American populations of giant liver fluke <i>Fascioloides magna</i> . Parasites and Vectors, 2015, 8, 288.	2.5	13
22	Asian fish tapeworm, <i>Khawia japonensis</i> (Yamaguti, 1934), has expanded its European invasive range. Parasitology Research, 2015, 114, 2035-2039.	1.6	7
23	Cytogenetics of <i>Aspidogaster limacoides</i> (Trematoda, Aspidogastrea): karyotype, spermatocyte division, and genome size. Parasitology Research, 2015, 114, 1473-1483.	1.6	9
24	Molecular characterization of <i>Fascioloides magna</i> (Trematoda: Fasciolidae) from south-western Poland based on mitochondrial markers. Acta Parasitologica, 2015, 60, 544-7.	1.1	12
25	Development of microsatellite markers in <i>Caryophyllaeus laticeps</i> (Cestoda: Caryophyllidea), monozoic fish tapeworm, using next-generation sequencing approach. Parasitology Research, 2015, 114, 721-726.	1.6	10
26	Conflict between morphology and molecular data: a case of the genus <i>Caryophyllaeus</i> (Cestoda: Caryophyllidea). Parasitology Research, 2014, 113, 19-24.	1.3	19
27	Molecular evidence of cryptic diversity in <i>Paracaryophyllaeus</i> (Cestoda: Caryophyllidea), parasites of loaches (Cobitidae) in Eurasia, including description of <i>P. vladkae</i> n. sp.. Parasitology International, 2014, 63, 841-850.	1.3	19
28	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. Molecular and Biochemical Parasitology, 2014, 195, 30-33.	1.1	11
29	Conflict between morphology and molecular data: a case of the genus <i>Caryophyllaeus</i> (Cestoda: Caryophyllidea). Parasitology Research, 2014, 113, 19-24.	1.3	19
30	The tapeworm <i>Atractolytocestus tenuicollis</i> (Cestoda: Caryophyllidea) is a sister species or ancestor of an invasive <i>A. huronensis</i> ?. Parasitology Research, 2013, 112, 3379-3388.	1.6	13
31	Ribosomal ITS2 structure in <i>Caryophyllaeus laticeps</i> and <i>Caryophyllaeus brachycollis</i> (Cestoda: Caryophyllidea). Parasitology Research, 2013, 112, 2661-2666.	0.9	7
32	The origin of the giant liver fluke, <i>Fascioloides magna</i> (Trematoda: Fasciolidae) from Croatia determined by high-resolution melting screening of mitochondrial <i>cox1</i> haplotypes. Parasitology Research, 2013, 112, 2661-2666.	1.6	13
33	Sequence structure and intragenomic variability of ribosomal ITS2 in monozoic tapeworms of the genus <i>Khawia</i> (Cestoda: Caryophyllidea), parasites of cyprinid fish. Parasitology Research, 2012, 111, 1621-1627.	1.6	18
34	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> . Parasitology Research, 2012, 110, 1621-1629.	1.6	14
35	Substitution saturation and nuclear paralogs of commonly employed phylogenetic markers in the Caryophyllidea, an unusual group of non-segmented tapeworms (Platyhelminthes). International Journal for Parasitology, 2012, 42, 259-267.	3.1	53
36	Multiple origins of European populations of the giant liver fluke <i>Fascioloides magna</i> (Trematoda: Fasciolidae). Parasitology Research, 2012, 110, 1621-1629.	3.1	52

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37	Development of high-resolution melting (HRM) analysis for population studies of <i>Fascioloides magna</i> (Trematoda: Fasciolidae), the giant liver fluke of ruminants. <i>Parasitology Research</i> , 2011, 108, 201-209.	1.6	22
38	Population study of <i>Atractolytocestus huronensis</i> (Cestoda: Caryophyllidea), an invasive parasite of common carp introduced to Europe: mitochondrial <i>cox1</i> haplotypes and intragenomic ribosomal ITS2 variants. <i>Parasitology Research</i> , 2011, 109, 125-131.	1.6	25
39	A comparative study of karyotypes and chromosomal location of rDNA genes in important liver flukes <i>Fasciola hepatica</i> and <i>Fascioloides magna</i> (Trematoda: Fasciolidae). <i>Parasitology Research</i> , 2011, 109, 1021-1028.	1.6	12
40	Revision of <i>Khawia</i> spp. (Cestoda: Caryophyllidea), parasites of cyprinid fish, including a key to their identification and molecular phylogeny. <i>Folia Parasitologica</i> , 2011, 58, 197-223.	1.3	43
41	Revision of <i>Khawia</i> spp. (Cestoda: Caryophyllidea), parasites of cyprinid fish, including a key to their identification and molecular phylogeny. <i>Folia Parasitologica</i> , 2011, 58, 197-223.	1.3	16
42	Determination of ribosomal internal transcribed spacer 2 (ITS2) interspecific markers in <i>Fasciola hepatica</i> , <i>Fascioloides magna</i> , <i>Dicrocoelium dendriticum</i> and <i>Paramphistomum cervi</i> (Trematoda), parasites of wild and domestic ruminants. <i>Helminthologia</i> , 2010, 47, 76-82.	0.9	49
43	A description of karyotype of the giant liver fluke <i>Fascioloides magna</i> (Trematoda, Platyhelminthes) and a review of Fasciolidae cytogenetics. <i>Helminthologia</i> , 2010, 47, 69-75.	0.9	6
44	Molecular discrimination of eggs of cervid trematodes using the Teflon (PTFE) technique for eggshell disruption. <i>Helminthologia</i> , 2010, 47, 147-151.	0.9	7
45	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.	3.1	43
46	Karyotype, chromosomal characteristics of multiple rDNA clusters and intragenomic variability of ribosomal ITS2 in <i>Caryophyllaeides fennica</i> (Cestoda). <i>Parasitology International</i> , 2010, 59, 351-357.	1.3	26