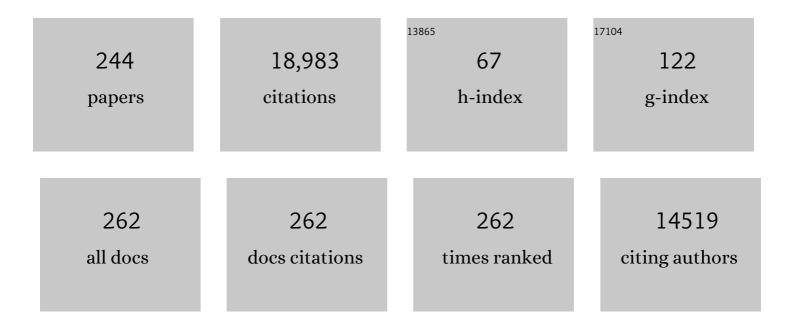
Dieter Ebert

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Conceptual issues in local adaptation. Ecology Letters, 2004, 7, 1225-1241.	6.4	2,964
2	Experimental evolution. Trends in Ecology and Evolution, 2012, 27, 547-560.	8.7	631
3	Host–parasite â€~Red Queen' dynamics archived in pond sediment. Nature, 2007, 450, 870-873.	27.8	537
4	Experimental Evolution of Parasites. , 1998, 282, 1432-1436.		489
5	Virulence and Local Adaptation of a Horizontally Transmitted Parasite. Science, 1994, 265, 1084-1086.	12.6	455
6	GENETIC VARIATION IN A HOST-PARASITE ASSOCIATION: POTENTIAL FOR COEVOLUTION AND FREQUENCY-DEPENDENT SELECTION. Evolution; International Journal of Organic Evolution, 2001, 55, 1136-1145.	2.3	443
7	On the evolutionary ecology of specific immune defence. Trends in Ecology and Evolution, 2003, 18, 27-32.	8.7	397
8	Sex against virulence: the coevolution of parasitic diseases. Trends in Ecology and Evolution, 1996, 11, 79-82.	8.7	305
9	Challenging the trade-off model for the evolution of virulence: is virulence management feasible?. Trends in Microbiology, 2003, 11, 15-20.	7.7	284
10	The evolution of parasitic diseases. Parasitology Today, 1996, 12, 96-101.	3.0	252
11	Intensive fish farming and the evolution of pathogen virulence: the case of columnaris disease in Finland. Proceedings of the Royal Society B: Biological Sciences, 2010, 277, 593-600.	2.6	230
12	The Effect of Parasites on Host Population Density and Extinction: Experimental Epidemiology with Daphnia and Six Microparasites. American Naturalist, 2000, 156, 459-477.	2.1	228
13	Within–and between–population variation for resistance of Daphnia magna to the bacterial endoparasite Pasteuria ramosa. Proceedings of the Royal Society B: Biological Sciences, 1998, 265, 2127-2134.	2.6	224
14	The First Myriapod Genome Sequence Reveals Conservative Arthropod Gene Content and Genome Organisation in the Centipede Strigamia maritima. PLoS Biology, 2014, 12, e1002005.	5.6	221
15	A Selective Advantage to Immigrant Genes in a Daphnia Metapopulation. Science, 2002, 295, 485-488.	12.6	220
16	The population dynamics of vertically and horizontally transmitted parasites. Proceedings of the Royal Society B: Biological Sciences, 1995, 260, 321-327.	2.6	210
17	The Evolution of Virulence When Parasites Cause Host Castration and Gigantism. American Naturalist, 2004, 164, S19-S32.	2.1	205
18	The Epidemiology and Evolution of Symbionts with Mixed-Mode Transmission. Annual Review of Ecology, Evolution, and Systematics, 2013, 44, 623-643.	8.3	194

#	Article	IF	CITATIONS
19	Host–parasite coevolution: Insights from the Daphnia–parasite model system. Current Opinion in Microbiology, 2008, 11, 290-301.	5.1	187
20	Dose effects and density-dependent regulation of two microparasites of Daphnia magna. Oecologia, 2000, 122, 200-209.	2.0	163
21	Empirical Support for Optimal Virulence in a Castrating Parasite. PLoS Biology, 2006, 4, e197.	5.6	154
22	Development, life cycle, ultrastructure and phylogenetic position of Pasteuria ramosa Metchnikoff 1888: rediscovery of an obligate endoparasite of Daphnia magna Straus. Philosophical Transactions of the Royal Society B: Biological Sciences, 1996, 351, 1689-1701.	4.0	148
23	In deep trouble: Habitat selection constrained by multiple enemies in zooplankton. Proceedings of the National Academy of Sciences of the United States of America, 2002, 99, 5481-5485.	7.1	146
24	The Dscam Homologue of the Crustacean Daphnia Is Diversified by Alternative Splicing Like in Insects. Molecular Biology and Evolution, 2008, 25, 1429-1439.	8.9	145
25	Intensive Farming: Evolutionary Implications for Parasites and Pathogens. Evolutionary Biology, 2010, 37, 59-67.	1.1	145
26	Prevalence, Host Specificity and Impact on Host Fecundity of Microparasites and Epibionts in Three Sympatric Daphnia Species. Journal of Animal Ecology, 1997, 66, 212.	2.8	141
27	A Matching-Allele Model Explains Host Resistance to Parasites. Current Biology, 2013, 23, 1085-1088.	3.9	137
28	Adaptive phenotypic plasticity and local adaptation for temperature tolerance in freshwater zooplankton. Proceedings of the Royal Society B: Biological Sciences, 2014, 281, 20132744.	2.6	136
29	Water fleas require microbiota for survival, growth and reproduction. ISME Journal, 2015, 9, 59-67.	9.8	132
30	Genetic diversity of <i>Daphnia magna</i> populations enhances resistance to parasites. Ecology Letters, 2008, 11, 918-928.	6.4	130
31	Does Internet-based guided-self-help for depression cause harm? An individual participant data meta-analysis on deterioration rates and its moderators in randomized controlled trials. Psychological Medicine, 2016, 46, 2679-2693.	4.5	129
32	Optimal killing for obligate killers: the evolution of life histories and virulence of semelparous parasites. Proceedings of the Royal Society B: Biological Sciences, 1997, 264, 985-991.	2.6	128
33	The curse of the pharaoh : the evolution of virulence in pathogens with long living propagules. Proceedings of the Royal Society B: Biological Sciences, 1996, 263, 715-721.	2.6	127
34	Dose–dependent infection rates of parasites produce the Allee effect in epidemiology. Proceedings of the Royal Society B: Biological Sciences, 2002, 269, 271-279.	2.6	122
35	Reduced flight-to-light behaviour of moth populations exposed to long-term urban light pollution. Biology Letters, 2016, 12, 20160111.	2.3	120
36	THE INFLUENCE OF HOST DEMOGRAPHY ON THE EVOLUTION OF VIRULENCE OF A MICROSPORIDIAN GUT PARASITE. Evolution; International Journal of Organic Evolution, 1997, 51, 1828-1837.	2.3	114

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37	Cloning of the unculturable parasite <i>Pasteuria ramosa</i> and its <i>Daphnia</i> host reveals extreme genotype–genotype interactions. Ecology Letters, 2011, 14, 125-131.	6.4	114
38	THE ORIGIN OF SPECIFICITY BY MEANS OF NATURAL SELECTION: EVOLVED AND NONHOST RESISTANCE IN HOST-PATHOGEN INTERACTIONS. Evolution; International Journal of Organic Evolution, 2013, 67, 1-9.	2.3	114
39	THE EFFECTS OF MULTIPLE INFECTIONS ON THE EXPRESSION AND EVOLUTION OF VIRULENCE IN A <i>DAPHNIA</i> à€ENDOPARASITE SYSTEM. Evolution; International Journal of Organic Evolution, 2008, 62, 1700-1711.	2.3	112
40	Evolution of a morphological novelty occurred before genome compaction in a lineage of extreme parasites. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 15480-15485.	7.1	111
41	The Reduced Genome of the Parasitic Microsporidian Enterocytozoon bieneusi Lacks Genes for Core Carbon Metabolism. Genome Biology and Evolution, 2010, 2, 304-309.	2.5	110
42	Host–parasite co-evolution and its genomic signature. Nature Reviews Genetics, 2020, 21, 754-768.	16.3	110
43	Evolution in Health and Disease: Work in Progress. Quarterly Review of Biology, 2001, 76, 417-432.	0.1	101
44	PARASITE-HOST SPECIFICITY: EXPERIMENTAL STUDIES ON THE BASIS OF PARASITE ADAPTATION. Evolution; International Journal of Organic Evolution, 2006, 60, 31-38.	2.3	100
45	Resolving the infection process reveals striking differences in the contribution of environment, genetics and phylogeny to host-parasite interactions. BMC Biology, 2011, 9, 11.	3.8	100
46	The Influence of Host Demography on the Evolution of Virulence of a Microsporidian Gut Parasite. Evolution; International Journal of Organic Evolution, 1997, 51, 1828.	2.3	97
47	Associations between parasitism and host genotype in natural populations ofDaphnia(Crustacea:) Tj ETQq1 1 0.7	'84314 rg 2.8	BT_/Overlock
48	HOST STARVATION DECREASES PARASITE LOAD AND MEAN HOST SIZE IN EXPERIMENTAL POPULATIONS. Ecology, 2004, 85, 823-833.	3.2	93
49	Ecological implications of parasites in natural Daphnia populations. Oecologia, 2005, 144, 382-390.	2.0	93
50	The effect of size at birth, maturation threshold and genetic differences on the life-history of Daphnia magna. Oecologia, 1991, 86, 243-250.	2.0	92
51	Time-shift experiments as a tool to study antagonistic coevolution. Trends in Ecology and Evolution, 2009, 24, 226-232.	8.7	92
52	Genetic Diversity and Genetic Differentiation in Daphnia Metapopulations With Subpopulations of Known Age. Genetics, 2005, 170, 1809-1820.	2.9	89
53	Daphnia magna transcriptome by RNA-Seq across 12 environmental stressors. Scientific Data, 2016, 3, 160030.	5.3	89
54	RESPONSES OF A BACTERIAL PATHOGEN TO PHOSPHORUS LIMITATION OF ITS AQUATIC INVERTEBRATE HOST. Ecology, 2008, 89, 313-318.	3.2	88

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55	TEMPORAL AND SPATIAL DYNAMICS OF PARASITE RICHNESS IN ADAPHNIAMETAPOPULATION. Ecology, 2001, 82, 3417-3434.	3.2	86
56	STRONG INBREEDING DEPRESSION IN A DAPHNIA METAPOPULATION. Evolution; International Journal of Organic Evolution, 2002, 56, 518-526.	2.3	79
57	Host Sexual Dimorphism and Parasite Adaptation. PLoS Biology, 2012, 10, e1001271.	5.6	79
58	The cause of parasitic infection in natural populations of Daphnia (Crustacea: Cladocera): the role of host genetics. Proceedings of the Royal Society B: Biological Sciences, 2000, 267, 2037-2042.	2.6	78
59	A quantitative test of the relationship between parasite dose and infection probability across different host–parasite combinations. Proceedings of the Royal Society B: Biological Sciences, 2008, 275, 853-859.	2.6	78
60	Comparative metagenomics of Daphnia symbionts. BMC Genomics, 2009, 10, 172.	2.8	78
61	The infection rate of Daphnia magna by Pasteuria ramosa conforms with the mass-action principle. Epidemiology and Infection, 2003, 131, 957-966.	2.1	75
62	A Genome for the Environment. Science, 2011, 331, 539-540.	12.6	74
63	Microbial ecosystems are dominated by specialist taxa. Ecology Letters, 2015, 18, 974-982.	6.4	74
64	Pathogen Dose Infectivity Curves as a Method to Analyze the Distribution of Host Susceptibility: A Quantitative Assessment of Maternal Effects after Food Stress and Pathogen Exposure. American Naturalist, 2010, 175, 106-115.	2.1	73
65	A Maturation Size Threshold and Phenotypic Plasticity of Age and Size at Maturity in Daphnia magna. Oikos, 1994, 69, 309.	2.7	71
66	Transgenerational effects of poor elemental food quality on Daphnia magna. Oecologia, 2010, 162, 865-872.	2.0	70
67	A Population Biology Perspective on the Stepwise Infection Process of the Bacterial Pathogen Pasteuria ramosa inADaphnia. Advances in Parasitology, 2016, 91, 265-310.	3.2	70
68	The Ecological Interactions between a Microsporidian Parasite and its Host Daphnia magna. Journal of Animal Ecology, 1995, 64, 361.	2.8	69
69	TEMPORAL PATTERNS OF GENETIC VARIATION FOR RESISTANCE AND INFECTIVITY IN A DAPHNIA-MICROPARASITE SYSTEM. Evolution; International Journal of Organic Evolution, 2001, 55, 1146-1152.	2.3	69
70	EVIDENCE FOR STRONG HOST CLONE-PARASITE SPECIES INTERACTIONS IN THE DAPHNIA MICROPARASITE SYSTEM. Evolution; International Journal of Organic Evolution, 2003, 57, 784-792.	2.3	69
71	Variation in phenoloxidase activity and its relation to parasite resistance within and between populations of Daphnia magna. Proceedings of the Royal Society B: Biological Sciences, 2004, 271, 1175-1183.	2.6	69
72	Ecological interactions of the microparasite Caullerya mesnili and its host Daphnia galeata. Limnology and Oceanography, 2002, 47, 300-305.	3.1	68

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73	Founder events as determinants of within-island and among-island genetic structure of Daphnia metapopulations. Heredity, 2006, 96, 150-158.	2.6	68
74	The coexistence of hybrid and parental Daphnia : the role of parasites. Proceedings of the Royal Society B: Biological Sciences, 2006, 273, 1977-1983.	2.6	68
75	Climate change affects colonization dynamics in a metacommunity of three <i>Daphnia</i> species. Global Change Biology, 2008, 14, 1209-1220.	9.5	67
76	Draft genome sequence of the Daphnia pathogen Octosporea bayeri: insights into the gene content of a large microsporidian genome and a model for host-parasite interactions. Genome Biology, 2009, 10, R106.	9.6	67
77	Genetics of life history in Daphnia magna. I. Heritabilities at two food levels. Heredity, 1993, 70, 335-343.	2.6	64
78	Benefits of host genetic diversity for resistance to infection depend on parasite diversity. Ecology, 2010, 91, 1263-1268.	3.2	63
79	Populations in small, ephemeral habitat patches may drive dynamics in a Daphnia magna metapopulation. Ecology, 2010, 91, 2975-2982.	3.2	63
80	Virulence and transmission modes of two microsporidia in <i>Daphnia magna</i> . Parasitology, 1995, 111, 133-142.	1.5	62
81	Brood pouch-mediated polystyrene nanoparticle uptake during <i>Daphnia magna</i> embryogenesis. Nanotoxicology, 2017, 11, 1059-1069.	3.0	60
82	A food-independent maturation threshold and size at maturity in Daphnia magna. Limnology and Oceanography, 1992, 37, 878-881.	3.1	56
83	Distributions and impacts of microparasites on Daphnia in a rockpool metapopulation. Oecologia, 1998, 115, 213-221.	2.0	56
84	Experimental evolution of field populations of <i>Daphnia magna</i> in response to parasite treatment. Journal of Evolutionary Biology, 2008, 21, 1068-1078.	1.7	55
85	Mixed inoculations of a microsporidian parasite with horizontal and vertical infections. Oecologia, 2005, 143, 157-166.	2.0	54
86	The first-generation Daphnia magna linkage map. BMC Genomics, 2010, 11, 508.	2.8	54
87	Disentangling the influence of parasite genotype, host genotype and maternal environment on different stages of bacterial infection in <i>Daphnia magna</i> . Proceedings of the Royal Society B: Biological Sciences, 2012, 279, 3176-3183.	2.6	54
88	Local adaptation of sex induction in a facultative sexual crustacean: insights from <scp>QTL</scp> mapping and natural populations of <i><scp>D</scp>aphnia magna</i> . Molecular Ecology, 2013, 22, 3567-3579.	3.9	54
89	Small subunit ribosomal DNA phylogeny of microsporidia that infect Daphnia (Crustacea: Cladocera). Parasitology, 2002, 124, 381-389.	1.5	53
90	PARASITE-MEDIATED SELECTION IN EXPERIMENTAL DAPHNIA MAGNA POPULATIONS. Evolution; International Journal of Organic Evolution, 2003, 57, 249-260.	2.3	53

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91	Haunted by the past: Evidence for dormant stage banks of microparasites and epibionts of Daphnia. Limnology and Oceanography, 2004, 49, 1355-1364.	3.1	52
92	Rethinking "mutualism―in diverse hostâ€symbiont communities. BioEssays, 2016, 38, 100-108.	2.5	52
93	The Evolutionary Consequences of Stepwise Infection Processes. Trends in Ecology and Evolution, 2017, 32, 612-623.	8.7	51
94	The genetic basis of resistance and matching-allele interactions of a host-parasite system: The Daphnia magna-Pasteuria ramosa model. PLoS Genetics, 2017, 13, e1006596.	3.5	51
95	Redescriptjon of Pleistophora intestinalis Chatton, 1907, a microsporidian parasite of Daphnia magna and Daphnia puiex, with establishment of the new genus Glugoides (Microspora, glugeidae). European Journal of Protistology, 1996, 32, 251-261.	1.5	50
96	The interactive effects of temperature, food level and maternal phenotype on offspring size in Daphnia magna. Oecologia, 1996, 107, 189-196.	2.0	50
97	Invasion thresholds and the evolution of nonequilibrium virulence. Evolutionary Applications, 2008, 1, 172-182.	3.1	50
98	GENETIC VARIATION IN A HOST-PARASITE ASSOCIATION: POTENTIAL FOR COEVOLUTION AND FREQUENCY-DEPENDENT SELECTION. Evolution; International Journal of Organic Evolution, 2001, 55, 1136.	2.3	49
99	Sex-specific effects of a parasite evolving in a female-biased host population. BMC Biology, 2012, 10, 104.	3.8	49
100	An SNP-based second-generation genetic map of Daphnia magna and its application to QTL analysis of phenotypic traits. BMC Genomics, 2014, 15, 1033.	2.8	49
101	Genetic architecture of resistance in Daphnia hosts against two species of host-specific parasites. Heredity, 2015, 114, 241-248.	2.6	49
102	TEST OF SYNERGISTIC INTERACTION BETWEEN INFECTION AND INBREEDING IN DAPHNIA MAGNA. Evolution; International Journal of Organic Evolution, 2003, 57, 777-783.	2.3	48
103	Experimental evidence for male biased flightâ€ŧoâ€light behavior in two moth species. Entomologia Experimentalis Et Applicata, 2009, 130, 259-265.	1.4	48
104	Apparent seasonality of parasite dynamics: analysis of cyclic prevalence patterns. Proceedings of the Royal Society B: Biological Sciences, 2006, 273, 199-206.	2.6	47
105	Fractional Resource Allocation into Few Eggs: Daphnia as an Example. Ecology, 1994, 75, 568-571.	3.2	46
106	Differential Adaptation in Spacially Heterogeneous Environments and Host-Parasite Coevolution. , 1998, , 325-342.		45
107	Cytological and molecular description of Hamiltosporidium tvaerminnensis gen. et sp. nov., a microsporidian parasite of Daphnia magna, and establishment of Hamiltosporidium magnivora comb. nov Parasitology, 2011, 138, 447-462.	1.5	45
108	The expression of virulence during double infections by different parasites with conflicting host exploitation and transmission strategies. Journal of Evolutionary Biology, 2011, 24, 1307-1316.	1.7	45

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109	Red Queen dynamics in multi-host and multi-parasite interaction system. Scientific Reports, 2015, 5, 10004.	3.3	45
110	Expression of parasite genetic variation changes over the course of infection: implications of within-host dynamics for the evolution of virulence. Proceedings of the Royal Society B: Biological Sciences, 2015, 282, 20142820.	2.6	45
111	A high-density genetic map reveals variation in recombination rate across the genome of Daphnia magna. BMC Genetics, 2016, 17, 137.	2.7	45
112	Genetics of life history in Daphnia magna. II. Phenotypic plasticity. Heredity, 1993, 70, 344-352.	2.6	44
113	Different mechanisms of transmission of the microsporidium Octosporea bayeri: a cocktail of solutions for the problem of parasite permanence. Parasitology, 2005, 130, 501-509.	1.5	44
114	Phenotypic plasticity of host-parasite interactions in response to the route of infection. Journal of Evolutionary Biology, 2005, 18, 911-921.	1.7	44
115	Physiology of Immunity in the Water Flea Daphnia magna: Environmental and Genetic Aspects of Phenoloxidase Activity. Physiological and Biochemical Zoology, 2003, 76, 836-842.	1.5	43
116	Dietary supply with polyunsaturated fatty acids and resulting maternal effects influence host – parasite interactions. BMC Ecology, 2013, 13, 41.	3.0	43
117	A fossil-calibrated phylogenomic analysis of Daphnia and the Daphniidae. Molecular Phylogenetics and Evolution, 2019, 137, 250-262.	2.7	43
118	Genes mirror geography in <i>DaphniaÂmagna</i> . Molecular Ecology, 2015, 24, 4521-4536.	3.9	41
119	A Photoreceptor Contributes to the Natural Variation of Diapause Induction in <i>Daphnia magna</i> . Molecular Biology and Evolution, 2016, 33, 3194-3204.	8.9	41
120	The microbiota of diapause: How host–microbe associations are formed after dormancy in an aquatic crustacean. Journal of Animal Ecology, 2018, 87, 400-413.	2.8	40
121	The influence of pool volume and summer desiccation on the production of the resting and dispersal stage in a Daphnia metapopulation. Oecologia, 2008, 157, 441-452.	2.0	39
122	<i>Daphnia</i> invest in sexual reproduction when its relative costs are reduced. Proceedings of the Royal Society B: Biological Sciences, 2018, 285, 20172176.	2.6	39
123	Desiccation of Rock Pool Habitats and Its Influence on Population Persistence in a Daphnia Metacommunity. PLoS ONE, 2009, 4, e4703.	2.5	39
124	Within-host dynamics of a microsporidium with horizontal and vertical transmission:Octosporea bayeriinDaphnia magna. Parasitology, 2004, 128, 31-38.	1.5	38
125	Bacterial infection changes the elemental composition of <i>Daphnia magna</i> . Journal of Animal Ecology, 2008, 77, 1265-1272.	2.8	38
126	Ultrastructural study and description of Ordospora colligata gen. et sp. nov. (microspora,) Tj ETQq0 0 0 rgBT /Ove	rlock 10 T 1.5	f 50 67 Td (37

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127	Parasites promote host gene flow in a metapopulation. Evolutionary Ecology, 2007, 21, 561-575.	1.2	37
128	Life history and virulence are linked in the ectoparasitic salmon louse <i>Lepeophtheirus salmonis</i> . Journal of Evolutionary Biology, 2012, 25, 856-861.	1.7	37
129	Response to: Eliot and Gandon and Day: Revisiting virulence management. Trends in Microbiology, 2003, 11, 208-209.	7.7	36
130	Parasite–mediated selection in experimental metapopulations of Daphnia magna. Proceedings of the Royal Society B: Biological Sciences, 2004, 271, 2149-2155.	2.6	36
131	Inference of parasite local adaptation using two different fitness components. Journal of Evolutionary Biology, 2007, 20, 921-929.	1.7	36
132	The Ordospora colligata Genome: Evolution of Extreme Reduction in Microsporidia and Host-To-Parasite Horizontal Gene Transfer. MBio, 2015, 6, .	4.1	36
133	Mitogenome phylogeographic analysis of a planktonic crustacean. Molecular Phylogenetics and Evolution, 2018, 129, 138-148.	2.7	36
134	High and Highly Variable Spontaneous Mutation Rates in <i>Daphnia</i> . Molecular Biology and Evolution, 2020, 37, 3258-3266.	8.9	36
135	Parasite-host specificity: experimental studies on the basis of parasite adaptation. Evolution; International Journal of Organic Evolution, 2006, 60, 31-8.	2.3	36
136	Resistance to a bacterial parasite in the crustacean Daphnia magna shows Mendelian segregation with dominance. Heredity, 2012, 108, 547-551.	2.6	34
137	Singleâ€nucleotide polymorphisms of two closely related microsporidian parasites suggest a clonal population expansion after the last glaciation. Molecular Ecology, 2013, 22, 314-326.	3.9	34
138	A novel approach to parasite population genetics: Experimental infection reveals geographic differentiation, recombination and hostâ€mediated population structure in <i><scp>P</scp>asteuria ramosa</i> , a bacterial parasite of <i><scp>D</scp>aphnia</i> . Molecular Ecology, 2013, 22, 972-986.	3.9	34
139	The effect of temperature on maturation threshold body-length in Daphnia magna. Oecologia, 1996, 108, 627-630.	2.0	33
140	Interactions between environmental stressors: the influence of salinity on host–parasite interactions between Daphnia magna and Pasteuria ramosa. Oecologia, 2013, 171, 789-796.	2.0	33
141	Host-parasite Red Queen dynamics with phase-locked rare genotypes. Science Advances, 2016, 2, e1501548.	10.3	33
142	Strong inbreeding depression in a Daphnia metapopulation. Evolution; International Journal of Organic Evolution, 2002, 56, 518-26.	2.3	33
143	STRONG INBREEDING DEPRESSION IN A DAPHNIA METAPOPULATION. Evolution; International Journal of Organic Evolution, 2002, 56, 518.	2.3	32
144	Diet quality determines interspecific parasite interactions in host populations. Ecology and Evolution, 2014, 4, 3093-3102.	1.9	32

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145	The effects of parasitism and inbreeding on the competitive ability in Daphnia magna: evidence for synergistic epistasis. Journal of Evolutionary Biology, 2003, 16, 976-985.	1.7	31
146	The Red Queen lives: Epistasis between linked resistance loci. Evolution; International Journal of Organic Evolution, 2016, 70, 480-487.	2.3	31
147	Variation and Plasticity of Biomass Allocation in Daphnia. Functional Ecology, 1994, 8, 435.	3.6	30
148	Genetic differences in the interactions of a microsporidian parasite and four clones of its cyclically parthenogenetic host. Parasitology, 1994, 108, 11-16.	1.5	28
149	Quantitative PCR to detect, discriminate and quantify intracellular parasites in their host: an example from three microsporidians in Daphnia. Parasitology, 2006, 133, 11.	1.5	28
150	Identification of a polymorphic collagen-like protein in the crustacean bacteria Pasteuria ramosa. Research in Microbiology, 2009, 160, 792-799.	2.1	28
151	The Combined Effect of Temperature and Host Clonal Line on the Microbiota of a Planktonic Crustacean. Microbial Ecology, 2018, 76, 506-517.	2.8	28
152	Dissecting the genetic architecture of a stepwise infection process. Molecular Ecology, 2019, 28, 3942-3957.	3.9	28
153	Expression of parasite virulence at different host population densities under natural conditions. Oecologia, 2009, 160, 247-255.	2.0	27
154	Characterisation of a large family of polymorphic collagen-like proteins in the endospore-forming bacterium Pasteuria ramosa. Research in Microbiology, 2011, 162, 701-714.	2.1	27
155	Temperature- versus precipitation-limitation shape local temperature tolerance in a Holarctic freshwater crustacean. Proceedings of the Royal Society B: Biological Sciences, 2019, 286, 20190929.	2.6	27
156	Microsporidia with Vertical Transmission Were Likely Shaped by Nonadaptive Processes. Genome Biology and Evolution, 2020, 12, 3599-3614.	2.5	27
157	PARASITE-HOST SPECIFICITY: EXPERIMENTAL STUDIES ON THE BASIS OF PARASITE ADAPTATION. Evolution; International Journal of Organic Evolution, 2006, 60, 31.	2.3	26
158	Population Genetics of Duplicated Alternatively Spliced Exons of the Dscam Gene in Daphnia and Drosophila. PLoS ONE, 2011, 6, e27947.	2.5	25
159	The role of moulting in parasite defence. Proceedings of the Royal Society B: Biological Sciences, 2012, 279, 3049-3054.	2.6	25
160	MORE THAN ONE WAY TO PRODUCE PROTEIN DIVERSITY: DUPLICATION AND LIMITED ALTERNATIVE SPLICING OF AN ADHESION MOLECULE GENE IN BASAL ARTHROPODS. Evolution; International Journal of Organic Evolution, 2013, 67, n/a-n/a.	2.3	25
161	Genetic, ecological and geographic covariables explaining host range and specificity of a microsporidian parasite. Journal of Animal Ecology, 2015, 84, 1711-1719.	2.8	25
162	Male-biased sex-ratio distortion caused by Octosporea bayeri, a vertically and horizontally-transmitted parasite of Daphnia magna. International Journal for Parasitology, 2008, 38, 969-979.	3.1	24

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163	Phylogenetic Characterization and Prevalence of " <i>Spirobacillus cienkowskii</i> ,―a Red-Pigmented, Spiral-Shaped Bacterial Pathogen of Freshwater <i>Daphnia</i> Species. Applied and Environmental Microbiology, 2008, 74, 1575-1582.	3.1	24
164	Covariation of Mitochondrial Genome Size with Gene Lengths: Evidence for Gene Length Reduction During Mitochondrial Evolution. Journal of Molecular Evolution, 2004, 59, 90-6.	1.8	23
165	A short term benefit for outcrossing in a Daphnia metapopulation in relation to parasitism. Journal of the Royal Society Interface, 2007, 4, 777-785.	3.4	23
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