

Stephen E Greiman

List of Publications by Year in descending order

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Version: 2024-02-01

40
papers

572
citations

687363
13
h-index

677142
22
g-index

40
all docs

40
docs citations

40
times ranked

649
citing authors

#	ARTICLE	IF	CITATIONS
1	Building an integrated infrastructure for exploring biodiversity: field collections and archives of mammals and parasites. <i>Journal of Mammalogy</i> , 2019, 100, 382-393.	1.3	61
2	Neorickettsial Endosymbionts of the Digenea. <i>Advances in Parasitology</i> , 2012, 79, 253-297.	3.2	59
3	The Beringian Coevolution Project: holistic collections of mammals and associated parasites reveal novel perspectives on evolutionary and environmental change in the North. <i>Arctic Science</i> , 2017, 3, 585-617.	2.3	50
4	Building Natural History Collections for the Twenty-First Century and Beyond. <i>BioScience</i> , 2020, 70, 674-687.	4.9	40
5	Transmission rates of the bacterial endosymbiont, <i>Neorickettsia risticii</i> , during the asexual reproduction phase of its digenean host, <i>Plagiorchis elegans</i> , within naturally infected lymnaeid snails. <i>Parasites and Vectors</i> , 2013, 6, 303.	2.5	31
6	Large Scale Screening of Digeneans for Neorickettsia Endosymbionts Using Real-Time PCR Reveals New Neorickettsia Genotypes, Host Associations and Geographic Records. <i>PLoS ONE</i> , 2014, 9, e98453.	2.5	31
7	Transformational Principles for NEON Sampling of Mammalian Parasites and Pathogens: A Response to Springer and Colleagues. <i>BioScience</i> , 2016, 66, 917-919.	4.9	28
8	Museum metabarcoding: A novel method revealing gut helminth communities of small mammals across space and time. <i>International Journal for Parasitology</i> , 2018, 48, 1061-1070.	3.1	26
9	Description and phylogenetic relationships of <i>Rodentolepis gnoskei</i> n. sp. (Cyclophyllidea: Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 343-350.	1.3	23
10	New genetic lineages, host associations and circulation pathways of Neorickettsia endosymbionts of digeneans. <i>Acta Parasitologica</i> , 2012, 57, 285-92.	1.1	19
11	Convoluted history and confusing morphology: Molecular phylogenetic analysis of dicrocoeliids reveals true systematic position of the Anenterotrematidae Yamaguti, 1958 (Platyhelminthes, Digenea). <i>Parasitology International</i> , 2018, 67, 501-508.	1.3	17
12	Description and Molecular Differentiation of a New <i>Staphylocystoides</i> (Cyclophyllidea: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 307 Parasitology, 2013, 99, 1045-1049.	0.7	15
13	<i>Nanophyetus salmincola</i> , vector of the salmon poisoning disease agent <i>Neorickettsia helminthoeca</i> , harbors a second pathogenic <i>Neorickettsia</i> species. <i>Veterinary Parasitology</i> , 2016, 229, 107-109.	1.8	14
14	Unravelling the diversity of the Crassiphialinae (Digenea: Diplostomidae) with molecular phylogeny and descriptions of five new species. <i>Current Research in Parasitology and Vector-borne Diseases</i> , 2021, 1, 100051.	1.9	13
15	Ultrastructure of the spermatozoon of the digenean <i>Plagiorchis elegans</i> (Rudolphi, 1802) (Plagiorchioidea, Plagiorchiidae). <i>Journal of Morphology</i> , 2013, 274, 965-972.	1.2	12
16	Vitellogenesis of the digenean <i>Plagiorchis elegans</i> (Rudolphi, 1802) (Plagiorchioidea, Plagiorchiidae). <i>Parasitology International</i> , 2014, 63, 537-543.	1.3	11
17	The numbers game: quantitative analysis of <i>Neorickettsia</i> sp. propagation through complex life cycle of its digenean host using real-time qPCR. <i>Parasitology Research</i> , 2016, 115, 2779-2788.	1.6	11
18	Germs within Worms: Localization of <i>Neorickettsia</i> sp. within Life Cycle Stages of the Digenean <i>Plagiorchis elegans</i> . <i>Applied and Environmental Microbiology</i> , 2016, 82, 2356-2362.	3.1	11

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19	Real-time PCR detection and phylogenetic relationships of <i>Neorickettsia</i> spp. in digeneans from Egypt, Philippines, Thailand, Vietnam and the United States. <i>Parasitology International</i> , 2017, 66, 1003-1007.	1.3	11
20	Build international biorepository capacity. <i>Science</i> , 2020, 370, 773-774.	12.6	9
21	Evidence for an Established Population of Tegu Lizards (<i>Salvator merianae</i>) in Southeastern Georgia, USA. <i>Southeastern Naturalist</i> , 2020, 19, .	0.4	9
22	Distinguishing Features of the Urinary Bacterial Microbiome in Patients with Neurogenic Lower Urinary Tract Dysfunction. <i>Journal of Urology</i> , 2022, 207, 627-634.	0.4	8
23	Laboratory maintenance of the bacterial endosymbiont, <i>Neorickettsia</i> sp., through the life cycle of a digenetic, <i>Plagiorchis elegans</i> . <i>Experimental Parasitology</i> , 2015, 157, 78-83.	1.2	7
24	Phylogenetic relationships and systematic position of the enigmatic <i>Uroterrema Braun</i> , 1900 (Platyhelminthes: Digenea). <i>Parasitology International</i> , 2019, 70, 118-122.	1.3	7
25	A New Species of Sucking Louse from the Long-Tailed Ground Squirrel, <i>Urocitellus undulatus</i> , from Mongolia, with a Key to Species, and a Review of Host Associations and Geographical Distributions of Members of the Genus <i>Linognathoides</i> (Psocidae: Anoplura: Polyplacidae). <i>Journal of Parasitology</i> , 2019, 105, 469.	0.7	7
26	First record of the Holarctic least shrew (<i>Sorex minutissimus</i>) and associated helminths from Canada: new light on northern Pleistocene refugia. <i>Canadian Journal of Zoology</i> , 2016, 94, 367-372.	1.0	6
27	Interrelationships of <i>Anenterotrema</i> (Digenea: Dicrocoeliidae) from Neotropical bats (Mammalia: Tj ETQq1 1 0.784314 rgBT /Overlock Research, 2021, 120, 2003-2016.	1.6	6
28	Hyperparasitism and Non-Nidicolous Mating by Male <i>Ixodes angustus</i> Ticks (Acari: Ixodidae). <i>Journal of Medical Entomology</i> , 2018, 55, 766-768.	1.8	5
29	Microbiomes From Biorepositories? 16S rRNA Bacterial Amplicon Sequencing of Archived and Contemporary Intestinal Samples of Wild Mammals (Eulipotyphla: Soricidae). <i>Frontiers in Ecology and Evolution</i> , 2020, 8, .	2.2	5
30	Method for the Rapid Fixation of Gastrointestinal Helminths in Small Mammals. <i>Acta Parasitologica</i> , 2019, 64, 406-410.	1.1	4
31	<i>Euschoengastia pipistrelli</i> (Acari: Trombiculidae) from American <i>Perimyotis</i> , <i>Perimyotis subflavus</i> (Chiroptera: Vespertilionidae): Novel Stereoscopic and Scanning Electron Microscopy. <i>Journal of Parasitology</i> , 2021, 107, 125-128.	0.7	3
32	A New Species of Sucking Louse from the Mandrill from Gabon with a Review of Host Associations and Geographical Distributions, and Identification Keys to Members of the Genus <i>Pedicinus</i> (Phthiraptera: Tj ETQq0 0 0rgBT /Overlock 10 T		
33	Fleas (Siphonaptera) Parasitizing Peridomestic and Indigenous Mammals in PanamÁ and Screening of Selected Fleas for Vector-Borne Bacterial Pathogens. <i>Journal of Medical Entomology</i> , 2021, 58, 1316-1321.	1.8	2
34	Transmission Biology, Host Associations, Distribution and Molecular Diagnostics of <i>Neorickettsia</i> . , 2016, , 295-325.		2
35	Detection of <i>Splendidofilaria</i> sp. (Onchocercidae:Splendidofilarinae) Microfilaria within Alaskan Ground-Dwelling Birds in the Grouse Subfamily Tetraoninae Using Taqman Probe-Based Real-Time PCR. <i>Journal of Parasitology</i> , 2022, 108, 192-198.	0.7	2
36	Description and Molecular Differentiation of a New Skrjabinoptera (Nematode: Physalopteridae) from <i>Eutropis macularia</i> (Sauria: Scincidae) in North-Central Vietnam. <i>Journal of Parasitology</i> , 2021, 107, 172-178.	0.7	1

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37	Two New Species of Sucking Lice (Phthiraptera: Anoplura: Hoplopleuridae and Polyplacidae) from Grant's Rock Mouse, <i>Micaelamys granti</i> , in South Africa. <i>Journal of Parasitology</i> , 2020, 106, 478.	0.7	1
38	Description and Molecular Differentiation of a New Falcaustra (Nematode: Kathlaniidae) from the Indochinese Water Dragon, <i>Physignathus cocincinus</i> (Squamata: Agamidae) in North-Central Vietnam. <i>Journal of Parasitology</i> , 2020, 107, 98-107.	0.7	1
39	A molecular reconstruction of holarctic Heligmosomidae reveals a new species of <i>Heligmosomoides</i> (Nematoda: Heligmosomidae) in <i>Peromyscus maniculatus</i> (Neotominae) from Canada. <i>Systematics and Biodiversity</i> , 2022, 20, 1-19.	1.2	1
40	A New Species of Sucking Louse from the Long-tailed Ground Squirrel, , from Mongolia, with a Key to Species, and a Review of Host Associations and Geographical Distributions of Members of the Genus (Psocodea: Anoplura: Polyplacidae). <i>Journal of Parasitology</i> , 2019, 105, 469-479.	0.7	0