

Rosemary G Gillespie

List of Publications by Year in descending order

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159
papers

10,908
citations

53794

45
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37204

96
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172
all docs

172
docs citations

172
times ranked

12859
citing authors

#	ARTICLE	IF	CITATIONS
1	Approaching a state shift in Earth's biosphere. <i>Nature</i> , 2012, 486, 52-58.	27.8	1,518
2	EARLY BURSTS OF BODY SIZE AND SHAPE EVOLUTION ARE RARE IN COMPARATIVE DATA. <i>Evolution; International Journal of Organic Evolution</i> , 2010, 64, no-no.	2.3	672
3	Phylogenetic analysis of community assembly and structure over space and time. <i>Trends in Ecology and Evolution</i> , 2008, 23, 619-630.	8.7	559
4	Community Assembly Through Adaptive Radiation in Hawaiian Spiders. <i>Science</i> , 2004, 303, 356-359.	12.6	521
5	Long-distance dispersal: a framework for hypothesis testing. <i>Trends in Ecology and Evolution</i> , 2012, 27, 47-56.	8.7	450
6	Arthropods on Islands: Colonization, Speciation, and Conservation. <i>Annual Review of Entomology</i> , 2002, 47, 595-632.	11.8	424
7	Islands as model systems in ecology and evolution: prospects fifty years after MacArthur & Wilson. <i>Ecology Letters</i> , 2015, 18, 200-217.	6.4	356
8	Topography-driven isolation, speciation and a global increase of endemism with elevation. <i>Global Ecology and Biogeography</i> , 2016, 25, 1097-1107.	5.8	243
9	EVOLUTION AND ECOLOGY OF SPIDER COLORATION. <i>Annual Review of Entomology</i> , 1998, 43, 619-643.	11.8	238
10	Chelex without boiling, a rapid and easy technique to obtain stable amplifiable DNA from small amounts of ethanol-preserved spiders. <i>Molecular Ecology Resources</i> , 2012, 12, 136-141.	4.8	230
11	Estimating and mitigating amplification bias in qualitative and quantitative arthropod metabarcoding. <i>Scientific Reports</i> , 2017, 7, 17668.	3.3	188
12	The black widow spider genus <i>Latrodectus</i> (Araneae: Theridiidae): phylogeny, biogeography, and invasion history. <i>Molecular Phylogenetics and Evolution</i> , 2004, 31, 1127-1142.	2.7	176
13	Treating Fossils as Terminal Taxa in Divergence Time Estimation Reveals Ancient Vicariance Patterns in the Palpimanoid Spiders. <i>Systematic Biology</i> , 2013, 62, 264-284.	5.6	175
14	Speciation and phylogeography of Hawaiian terrestrial arthropods. <i>Molecular Ecology</i> , 1998, 7, 519-531.	3.9	160
15	The Latitudinal Diversity Gradient: Novel Understanding through Mechanistic Eco-evolutionary Models. <i>Trends in Ecology and Evolution</i> , 2019, 34, 211-223.	8.7	151
16	Impacts of global climate change on the floras of oceanic islands – Projections, implications and current knowledge. <i>Perspectives in Plant Ecology, Evolution and Systematics</i> , 2015, 17, 160-183.	2.7	147
17	Comparing Adaptive Radiations Across Space, Time, and Taxa. <i>Journal of Heredity</i> , 2020, 111, 1-20.	2.4	146
18	Risk-Sensitive Foraging Strategies of Two Spider Populations. <i>Ecology</i> , 1987, 68, 887-899.	3.2	140

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19	From a comb to a tree: phylogenetic relationships of the comb-footed spiders (Araneae, Theridiidae) inferred from nuclear and mitochondrial genes. <i>Molecular Phylogenetics and Evolution</i> , 2004, 31, 225-245.	2.7	138
20	Nanopore sequencing of long ribosomal DNA amplicons enables portable and simple biodiversity assessments with high phylogenetic resolution across broad taxonomic scale. <i>GigaScience</i> , 2019, 8, .	6.4	126
21	Comparative phylogeography of oceanic archipelagos: Hotspots for inferences of evolutionary process. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 7986-7993.	7.1	124
22	Convergent evolution of behavior in an adaptive radiation of Hawaiian web-building spiders. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2004, 101, 16228-16233.	7.1	116
23	Speciation on a Conveyor Belt: Sequential Colonization of the Hawaiian Islands by Orsonwelles Spiders (Araneae, Linyphiidae). <i>Systematic Biology</i> , 2003, 52, 70-88.	5.6	113
24	Biodiversity dynamics in isolated island communities: interaction between natural and human-mediated processes. <i>Molecular Ecology</i> , 2008, 17, 45-57.	3.9	108
25	Are three-dimensional spider webs defensive adaptations?. <i>Ecology Letters</i> , 2002, 6, 13-18.	6.4	105
26	Multiple origins of a spider radiation in Hawaii.. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1994, 91, 2290-2294.	7.1	97
27	Molecular insights into the phylogenetic structure of the spider genus <i>Theridion</i> (Araneae, Tj ETQq1 1 0.784314 rgBT /Overlock 10 T	1.7	95
28	The notes from nature tool for unlocking biodiversity records from museum records through citizen science. <i>ZooKeys</i> , 2012, 209, 219-233.	1.1	85
29	Geology and climate drive diversification. <i>Nature</i> , 2014, 509, 297-298.	27.8	85
30	Influence of volcanic activity on the population genetic structure of Hawaiian Tetragnatha spiders: fragmentation, rapid population growth and the potential for accelerated evolution. <i>Molecular Ecology</i> , 2004, 13, 1729-1743.	3.9	82
31	The Global Museum: natural history collections and the future of evolutionary science and public education. <i>PeerJ</i> , 2020, 8, e8225.	2.0	81
32	Biogeography of spiders on remote oceanic islands of the Pacific: archipelagoes as stepping stones?. <i>Journal of Biogeography</i> , 2002, 29, 655-662.	3.0	78
33	A cost-efficient and simple protocol to enrich prey <scp>DNA</scp> from extractions of predatory arthropods for large-scale gut content analysis by Illumina sequencing. <i>Methods in Ecology and Evolution</i> , 2017, 8, 126-134.	5.2	75
34	Biogeography of the fauna of French Polynesia: diversification within and between a series of hot spot archipelagos. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2008, 363, 3335-3346.	4.0	74
35	Global Island Monitoring Scheme (GIMS): a proposal for the long-term coordinated survey and monitoring of native island forest biota. <i>Biodiversity and Conservation</i> , 2018, 27, 2567-2586.	2.6	72
36	Encyclopedia of Islands. , 2019, , .		64

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37	The role of prey availability in aggregative behaviour of the orb weaving spider <i>Tetragnatha elongata</i> . <i>Animal Behaviour</i> , 1987, 35, 675-681.	1.9	62
38	Community assembly on isolated islands: macroecology meets evolution. <i>Global Ecology and Biogeography</i> , 2016, 25, 769-780.	5.8	62
39	Family ties: molecular phylogeny of crab spiders (Araneae: Thomisidae). <i>Cladistics</i> , 2008, 24, 708-722.	3.3	59
40	A Network Perspective for Community Assembly. <i>Frontiers in Ecology and Evolution</i> , 2019, 7, .	2.2	59
41	COMPOSITIONAL AND FUNCTIONAL STABILITY OF ARTHROPOD COMMUNITIES IN THE FACE OF ANT INVASIONS. , 2008, 18, 1547-1562.		57
42	Island time and the interplay between ecology and evolution in species diversification. <i>Evolutionary Applications</i> , 2016, 9, 53-73.	3.1	57
43	Molecular systematics of <i>Selenops</i> spiders (Araneae: Selenopidae) from North and Central America: implications for Caribbean biogeography. <i>Biological Journal of the Linnean Society</i> , 0, 101, 288-322.	1.6	54
44	Phylogenetic placement of pelican spiders (Archaeidae, Araneae), with insight into evolution of the "neck" and predatory behaviours of the superfamily Palpimanoidea. <i>Cladistics</i> , 2012, 28, 598-626.	3.3	53
45	More data, fewer shifts: Molecular insights into the evolution of the spinning apparatus in non-orb-weaving spiders. <i>Molecular Phylogenetics and Evolution</i> , 2008, 46, 347-368.	2.7	51
46	De novo characterization of the gene-rich transcriptomes of two color-polymorphic spiders, <i>Theridion grallator</i> and <i>T. californicum</i> (Araneae: Theridiidae), with special reference to pigment genes. <i>BMC Genomics</i> , 2013, 14, 862.	2.8	51
47	The founding charter of the Genomic Observatories Network. <i>GigaScience</i> , 2014, 3, 2.	6.4	51
48	Island hopping across the central Pacific: mitochondrial DNA detects sequential colonization of the Austral Islands by crab spiders (Araneae: Thomisidae). <i>Journal of Biogeography</i> , 2006, 33, 201-220.	3.0	49
49	Tarsal Organ Morphology and the Phylogeny of Goblin Spiders (Araneae, Oonopidae), with Notes on Basal Genera. <i>American Museum Novitates</i> , 2012, 3736, 1-52.	0.6	49
50	Non-congruent colonizations and diversification in a coevolving pollination mutualism on oceanic islands. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2013, 280, 20130361.	2.6	49
51	Repeated Diversification of Ecomorphs in Hawaiian Stick Spiders. <i>Current Biology</i> , 2018, 28, 941-947.e3.	3.9	49
52	Island ecology and evolution: challenges in the Anthropocene. <i>Environmental Conservation</i> , 2017, 44, 323-335.	1.3	47
53	SELECTION ON THE COLOR POLYMORPHISM IN HAWAIIAN HAPPY-FACE SPIDERS: EVIDENCE FROM GENETIC STRUCTURE AND TEMPORAL FLUCTUATIONS. <i>Evolution; International Journal of Organic Evolution</i> , 1998, 52, 775-783.	2.3	46
54	Resource Consumption Variance Within and Among Individuals: On Coloniality in Spiders. <i>Ecology</i> , 1995, 76, 196-205.	3.2	45

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55	Unifying macroecology and macroevolution to answer fundamental questions about biodiversity. <i>Global Ecology and Biogeography</i> , 2019, 28, 1925-1936.	5.8	44
56	Diversity despite dispersal: colonization history and phylogeography of Hawaiian crab spiders inferred from multilocus genetic data. <i>Molecular Ecology</i> , 2009, 18, 1746-1764.	3.9	43
57	Are you what you eat? A highly transient and prey-influenced gut microbiome in the grey house spider <i>Badumna longinqua</i> . <i>Molecular Ecology</i> , 2020, 29, 1001-1015.	3.9	39
58	High-throughput sequencing for community analysis: the promise of DNA barcoding to uncover diversity, relatedness, abundances and interactions in spider communities. <i>Development Genes and Evolution</i> , 2020, 230, 185-201.	0.9	39
59	Repeated Evolution of Power-Amplified Predatory Strikes in Trap-Jaw Spiders. <i>Current Biology</i> , 2016, 26, 1057-1061.	3.9	37
60	Portraits of Evolution: Studies of Coloration in Hawaiian Spiders. <i>BioScience</i> , 2001, 51, 521.	4.9	35
61	Chromosome-level reference genome of the European wasp spider <i>Argiope bruennichi</i> : a resource for studies on range expansion and evolutionary adaptation. <i>GigaScience</i> , 2021, 10, .	6.4	35
62	ESTIMATION OF CAPTURE AREAS OF SPIDER ORB WEBS IN RELATION TO ASYMMETRY. <i>Journal of Arachnology</i> , 2002, 30, 70.	0.5	33
63	Species diversification patterns in the Polynesian jumping spider genus <i>Havaika</i> (Araneae, Salticidae). <i>Molecular Phylogenetics and Evolution</i> , 2006, 41, 472-495.	2.7	33
64	Island Biogeography of Remote Archipelagoes. , 2009, , 358-387.		33
65	Genetics of a colour polymorphism in <i>Theridion grallator</i> (Araneae: Theridiidae), the Hawaiian happy-face spider, from Greater Maui. <i>Heredity</i> , 1996, 76, 238-248.	2.6	30
66	Rapid divergence of mussel populations despite incomplete barriers to dispersal. <i>Molecular Ecology</i> , 2018, 27, 1556-1571.	3.9	29
67	The effect of DNA degradation bias in passive sampling devices on metabarcoding studies of arthropod communities and their associated microbiota. <i>PLoS ONE</i> , 2018, 13, e0189188.	2.5	29
68	Biogeography and the evolution of flightlessness in a radiation of Hawaiian moths (Xyloryctidae:). <i>Tj ETQq0 0 0 rgBT/Overlock 10 Tf 50 2</i>	3.0	28
69	Spider webs, stable isotopes and molecular gut content analysis: Multiple lines of evidence support trophic niche differentiation in a community of Hawaiian spiders. <i>Functional Ecology</i> , 2019, 33, 1722-1733.	3.6	28
70	Selection on the Color Polymorphism in Hawaiian Happy-Face Spiders: Evidence from Genetic Structure and Temporal Fluctuations. <i>Evolution; International Journal of Organic Evolution</i> , 1998, 52, 775.	2.3	27
71	Repeated colonization of remote islands by specialized mutualists. <i>Biology Letters</i> , 2012, 8, 258-261.	2.3	26
72	Scaling up <i>DNA</i> barcoding – Primer sets for simple and cost efficient arthropod systematics by multiplex <i>PCR</i> and Illumina amplicon sequencing. <i>Methods in Ecology and Evolution</i> , 2018, 9, 2181-2193.	5.2	26

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73	Effects of Natural Forest Fragmentation on a Hawaiian Spider Community. <i>Environmental Entomology</i> , 2004, 33, 1296-1305.	1.4	25
74	GEOGRAPHICAL CONTEXT OF SPECIATION IN A RADIATION OF HAWAIIAN TETRAGNATHA SPIDERS (ARANEAE, Tj ETQq0 0 0 rgBT /Overl	0.5	25
75	Species Differentiation on a Dynamic Landscape: Shifts in Metapopulation Genetic Structure Using the Chronology of the Hawaiian Archipelago. <i>Evolutionary Biology</i> , 2012, 39, 192-206.	1.1	25
76	HAWAIIAN SPIDERS OF THE GENUS TETRAGNATHA: IV NEW, SMALL SPECIES IN THE SPINY LEG CLADE. <i>Journal of Arachnology</i> , 2002, 30, 159.	0.5	24
77	Why is Madagascar special? The extraordinarily slow evolution of pelican spiders (Araneae, Tj ETQq1 1 0.784314 rgBT /Overl	2.3	24
78	A unified model of species abundance, genetic diversity, and functional diversity reveals the mechanisms structuring ecological communities. <i>Molecular Ecology Resources</i> , 2021, 21, 2782-2800.	4.8	24
79	Evolution of Satellite DNAs in a Radiation of Endemic Hawaiian Spiders: Does Concerted Evolution of Highly Repetitive Sequences Reflect Evolutionary History?. <i>Journal of Molecular Evolution</i> , 2004, 59, 632-641.	1.8	23
80	Oceanic Islands: Models of Diversity. , 2007, , 1-13.		23
81	Sexual dimorphism in venom chemistry in Tetragnatha spiders is not easily explained by adult niche differences. <i>Toxicon</i> , 2016, 114, 45-52.	1.6	23
82	What makes a happy face? Determinants of colour pattern in the Hawaiian happy face spider <i>Theridion grallator</i> (Araneae, Theridiidae). <i>Heredity</i> , 1989, 62, 355-363.	2.6	22
83	Costs and Benefits of Brood Care in the Hawaiian Happy Face Spider <i>Theridion grallator</i> (Araneae, Tj ETQq1 1 0.784314 rgBT /Overl	0.4	22
84	Common origin of the satellite DNAs of the Hawaiian spiders of the genus <i>Tetragnatha</i> : evolutionary constraints on the length and nucleotide composition of the repeats. <i>Gene</i> , 2003, 313, 169-177.	2.2	22
85	COLONIZATION HISTORY AND POPULATION GENETICS OF THE COLOR-POLYMORPHIC HAWAIIAN HAPPY-FACE SPIDER <i>THERIDION GRALLATOR</i> (ARANEAE, THERIDIIDAE). <i>Evolution; International Journal of Organic Evolution</i> , 2012, 66, 2815-2833.	2.3	22
86	Correlates of vulnerability among arthropod species threatened by invasive ants. <i>Biodiversity and Conservation</i> , 2010, 19, 1971-1988.	2.6	21
87	Sexually dimorphic venom proteins in long-jawed orb-weaving spiders (<i>Tetragnatha</i>) comprise novel gene families. <i>PeerJ</i> , 2018, 6, e4691.	2.0	21
88	Predation Through Impalement of Prey: The Foraging Behavior of <i>Doryonychus Raptor</i> (Araneae, Tj ETQq0 0 0 rgBT /Overl	0.9	20
89	Regional patterns in the invasion success of <i>Cheiracanthium</i> spiders (Miturgidae) in vineyard ecosystems. <i>Biological Invasions</i> , 2010, 12, 2499-2508.	2.4	20
90	Stabilizing selection maintains exuberant colour polymorphism in the spider <i>Theridion californicum</i> (Araneae, Theridiidae). <i>Molecular Ecology</i> , 2011, 20, 206-218.	3.9	20

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91	Does biological intimacy shape ecological network structure? A test using a brood pollination mutualism on continental and oceanic islands. <i>Journal of Animal Ecology</i> , 2018, 87, 1160-1171.	2.8	20
92	Co-occurrence of ecologically similar species of Hawaiian spiders reveals critical early phase of adaptive radiation. <i>BMC Evolutionary Biology</i> , 2018, 18, 100.	3.2	20
93	Quantum shifts in the genetic control of a colour polymorphism in <i>Theridion grallator</i> (Araneae: Tj ETQq1 1 0.784314 rgBT /Overlock 19	2.6	19
94	Sharing and reporting benefits from biodiversity research. <i>Molecular Ecology</i> , 2021, 30, 1103-1107.	3.9	19
95	HAWAIIAN SPIDERS OF THE GENUS TETRAGNATHA (ARANEAE, TETRAGNATHIDAE): V. ELONGATE WEB-BUILDERS FROM OAHU. <i>Journal of Arachnology</i> , 2003, 31, 8-19.	0.5	18
96	Evolution of cave living in Hawaiian <i>Schrankia</i> (Lepidoptera: Noctuidae) with description of a remarkable new cave species. <i>Zoological Journal of the Linnean Society</i> , 2009, 156, 114-139.	2.3	18
97	Patterns of habitat affinity and Austral/Holarctic parallelism in dictynoid spiders (Araneae:Entelegynae). <i>Invertebrate Systematics</i> , 2010, 24, 238.	1.3	18
98	Maintaining a happy face: stable colour polymorphism in the spider <i>Theiridion grallator</i> (Araneae,) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50	2.6	17
99	SPIDERS OF THE GENUS TETRAGNATHA (ARANEAE, TETRAGNATHIDAE) IN THE SOCIETY ISLANDS. <i>Journal of Arachnology</i> , 2003, 31, 157-172.	0.5	17
100	Shifting habitats, morphology, and selective pressures: Developmental polyphenism in an adaptive radiation of Hawaiian spiders. <i>Evolution; International Journal of Organic Evolution</i> , 2015, 69, 162-178.	2.3	17
101	Steppingâ€stones across space and time: repeated radiation of Pacific flightless broadâ€nosed weevils (Coleoptera: Curculionidae: Entiminae: <i>Rhyncogonus</i>). <i>Journal of Biogeography</i> , 2017, 44, 784-796.	3.0	17
102	Rapid and cost-effective generation of single specimen multilocus barcoding data from whole arthropod communities by multiple levels of multiplexing. <i>Scientific Reports</i> , 2020, 10, 78.	3.3	17
103	The effects of genetic background on the island-specific control of a colour polymorphism in <i>Theridion grallator</i> (Araneae: Theridiidae), the Hawaiian happy-face spider. <i>Heredity</i> , 1996, 76, 257-266.	2.6	16
104	A comparison of populations of island and adjacent mainland species of Caribbean <i>Selenops</i> (Araneae:) Tj ETQq0 0 0 rgBT /Overlock 10	2.7	16
105	Phytophagous insect community assembly through niche conservatism on oceanic islands. <i>Journal of Biogeography</i> , 2013, 40, 225-235.	3.0	16
106	New sequencing technologies, the development of genomics tools, and their applications in evolutionary arachnology. <i>Journal of Arachnology</i> , 2014, 42, 1-15.	0.5	16
107	Community assembly on remote islands: a comparison of Hawaiian and Mascarene spiders. <i>Journal of Biogeography</i> , 2015, 42, 39-50.	3.0	16
108	Multiplex <i>PCR</i> targeting lineage-specific <i>SNP</i> s: A highly efficient and simple approach to block out predator sequences in molecular gut content analysis. <i>Methods in Ecology and Evolution</i> , 2019, 10, 982-993.	5.2	16

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109	The <i>Tetragnatha kauaiensis</i> Genome Sheds Light on the Origins of Genomic Novelty in Spiders. <i>Genome Biology and Evolution</i> , 2021, 13, .	2.5	16
110	Life history of the spider <i>Selenops occultus</i> (Araneae, Selenopidae) from Brazil with notes on the natural history of the genus. <i>Journal of Natural History</i> , 2008, 42, 2747-2761.	0.5	15
111	Adaptive Radiation: Convergence and Non-equilibrium. <i>Current Biology</i> , 2013, 23, R71-R74.	3.9	15
112	Niche conservatism predominates in adaptive radiation: comparing the diversification of Hawaiian arthropods using ecological niche modelling. <i>Biological Journal of the Linnean Society</i> , 2019, 127, 479-492.	1.6	15
113	Giant Goblins above the waves at the southern end of the world: The biogeography of the spider family Orsolobidae (Araneae, Dysderoidea). <i>Journal of Biogeography</i> , 2019, 46, 332-342.	3.0	15
114	Impaled prey. <i>Nature</i> , 1992, 355, 212-213.	27.8	14
115	FREE-LIVING SPIDERS OF THE GENUS ARIAMNES (ARANEAE, THERIDIIDAE) IN HAWAII. <i>Journal of Arachnology</i> , 2007, 35, 11-37.	0.5	14
116	Sampling across space and time to validate natural experiments: an example with ant invasions in Hawaii. <i>Biological Invasions</i> , 2010, 12, 643-655.	2.4	14
117	Ancient DNA Resolves the History of <i>Tetragnatha</i> (Araneae, Tetragnathidae) Spiders on Rapa Nui. <i>Genes</i> , 2017, 8, 403.	2.4	14
118	Categorization of species as native or nonnative using DNA sequence signatures without a complete reference library. <i>Ecological Applications</i> , 2019, 29, e01914.	3.8	14
119	The Ecology and Evolution of Hawaiian Spider Communities. <i>American Scientist</i> , 2005, 93, 122.	0.1	14
120	MARQUESAN SPIDERS OF THE GENUS TETRAGNATHA (ARANEAE, TETRAGNATHIDAE). <i>Journal of Arachnology</i> , 2003, 31, 62-77.	0.5	13
121	Population structure and dispersal in a patchy landscape: nuclear and mitochondrial markers reveal area effects in the spider <i>Theridion californicum</i> (Araneae: Theridiidae). <i>Biological Journal of the Linnean Society</i> , 2011, 104, 600-620.	1.6	13
122	Adaptive Radiation. , 2001, , 25-44.		13
123	Correlated evolution between coloration and ambush site in predators with visual prey lures. <i>Evolution; International Journal of Organic Evolution</i> , 2017, 71, 2010-2021.	2.3	12
124	Limited Evidence for Microbial Transmission in the Phyllosymbiosis between Hawaiian Spiders and Their Microbiota. <i>MSystems</i> , 2022, 7, e0110421.	3.8	12
125	Pseudorabies in Captive Coyotes. <i>Journal of Wildlife Diseases</i> , 1997, 33, 916-918.	0.8	11
126	Stable isotopes of Hawaiian spiders reflect substrate properties along a chronosequence. <i>PeerJ</i> , 2018, 6, e4527.	2.0	11

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127	The energetics of mimicry: the cost of pedestrian transport in a formicine ant and its mimic, a clubionid spider. <i>Physiological Entomology</i> , 1989, 14, 173-177.	1.5	10
128	Ancient biogeography of generalist predators on remote oceanic islands. <i>Journal of Biogeography</i> , 2017, 44, 1098-1109.	3.0	10
129	Semi-quantitative metabarcoding reveals how climate shapes arthropod community assembly along elevation gradients on Hawaii Island. <i>Molecular Ecology</i> , 2022, 31, 1416-1429.	3.9	10
130	Oceanic Islands: Models of Diversity. , 2001, , 590-599.		9
131	Desert salt flats as oases for the spider <i>Saltonia incerta</i> Banks (Araneae: Dictynidae). <i>Ecology and Evolution</i> , 2014, 4, 3861-3874.	1.9	9
132	Comparative Transcriptomics of Maturity-Associated Color Change in Hawaiian Spiders. <i>Journal of Heredity</i> , 2014, 105, 771-781.	2.4	8
133	Shifting roles of the East China Sea in the phylogeography of red nanmu in East Asia. <i>Journal of Biogeography</i> , 2021, 48, 2486-2501.	3.0	8
134	Foraging Behavior of the Hawaiian Happy Face Spider (Araneae: Theridiidae). <i>Annals of the Entomological Society of America</i> , 1994, 87, 815-822.	2.5	7
135	Chemical Species Recognition in a Tetragnatha Spider (Araneae: Tetragnathidae). <i>Journal of Chemical Ecology</i> , 2021, 47, 63-72.	1.8	7
136	Towards eradicating the nuisance of numts and noise in molecular biodiversity assessment. <i>Molecular Ecology Resources</i> , 2021, 21, 1755-1758.	4.8	7
137	Richness and resilience in the Pacific: <code><scp>DNA</scp></code> metabarcoding enables parallelized evaluation of biogeographic patterns. <i>Molecular Ecology</i> , 2023, 32, 6710-6723.	3.9	7
138	Adaptation under a microscope. <i>Nature</i> , 2007, 446, 386-387.	27.8	6
139	Host and geography together drive early adaptive radiation of Hawaiian planthoppers. <i>Molecular Ecology</i> , 2019, 28, 4513-4528.	3.9	6
140	Shifts in morphology, gene expression, and selection underlie web loss in Hawaiian Tetragnatha spiders. <i>Bmc Ecology and Evolution</i> , 2021, 21, 48.	1.6	6
141	Non-native spiders change assemblages of Hawaiian forest fragment kipuka over space and time. <i>NeoBiota</i> , 0, 55, 1-9.	1.0	6
142	UNUSUALLY LONG HYPTIOTES (ARANEAE, ULOBORIDAE) SEQUENCE FOR SMALL SUBUNIT (18S) RIBOSOMAL RNA SUPPORTS SECONDARY STRUCTURE MODEL UTILITY IN SPIDERS. <i>Journal of Arachnology</i> , 2006, 34, 557-565.	0.5	5
143	A holobiont view of island biogeography: Unravelling patterns driving the nascent diversification of a Hawaiian spider and its microbial associates. <i>Molecular Ecology</i> , 2022, 31, 1299-1316.	3.9	5
144	Rediscovery and Uncertain Future of High-Elevation Haleakala Carabid Beetles (Coleoptera). <i>Pacific Science</i> , 2005, 59, 399-410.	0.6	4

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145	Bringing spiders to the multilocus era: novel anonymous nuclear markers for Harpactocrates ground-dwelling spiders (Araneae: Dysderidae) with application to related genera. Journal of Arachnology, 2011, 39, 506-510.	0.5	4
146	First come, first served: Possible role for priority effects in marine populations under different degrees of dispersal potential. Journal of Biogeography, 2020, 47, 1649-1662.	3.0	4
147	Range contraction and extinction vulnerability: what is natural?. Memoirs of the Museum of Victoria, 1997, 56, 401-409.	0.4	4
148	Cost effective microsatellite isolation and genotyping by high throughput sequencing. Journal of Arachnology, 2019, 47, 190.	0.5	4
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