Matthew K Fujita

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4875389/publications.pdf

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58 papers

4,448 citations

249298 26 h-index 56 g-index

61 all docs

61 docs citations

times ranked

61

6631 citing authors

#	Article	IF	CITATIONS
1	Giant Tree Frog diversification in West and Central Africa: Isolation by physical barriers, climate, and reproductive traits. Molecular Ecology, 2022, 31, 3979-3998.	2.0	7
2	Phylogenomics, introgression, and demographic history of South American true toads (<i>Rhinella</i>). Molecular Ecology, 2022, 31, 978-992.	2.0	14
3	Reduced mitochondrial respiration in hybrid asexual lizards. American Naturalist, 2022, 199, 719-728.	1.0	O
4	Ecology drives patterns of spectral transmission in the ocular lenses of frogs and salamanders. Functional Ecology, 2022, 36, 850-864.	1.7	8
5	Parthenogenesis doubles the rate of amino acid substitution in whiptail mitochondria. Evolution; International Journal of Organic Evolution, 2022, , .	1.1	2
6	Eyeâ€body allometry across biphasic ontogeny in anuran amphibians. Evolutionary Ecology, 2021, 35, 337-359.	0.5	14
7	The first mitochondrial genome of a South America parthenogenetic lizard (Squamata:) Tj ETQq1 1 0.784314 rgBT	Overlock	2 10 Tf 50 50
8	Who's your daddy? On the identity and distribution of the paternal hybrid ancestor of the parthenogenetic gecko Lepidodactylus lugubris (Reptilia: Squamata: Gekkonidae). Zootaxa, 2021, 4999, 87-100.	0.2	8
9	Delimitation despite discordance: Evaluating the species limits of a confounding species complex in the face of mitonuclear discordance. Ecology and Evolution, 2021, 11, 12739-12753.	0.8	11
10	Evolutionary Dynamics and Consequences of Parthenogenesis in Vertebrates. Annual Review of Ecology, Evolution, and Systematics, 2020, 51, 191-214.	3.8	27
11	Eye size and investment in frogs and toads correlate with adult habitat, activity pattern and breeding ecology. Proceedings of the Royal Society B: Biological Sciences, 2020, 287, 20201393.	1.2	32
12	Finding complexity in complexes: Assessing the causes of mitonuclear discordance in a problematic species complex of Mesoamerican toads. Molecular Ecology, 2020, 29, 3543-3559.	2.0	29
13	Transcriptome sequencing reveals signatures of positive selection in the Spot-Tailed Earless Lizard. PLoS ONE, 2020, 15, e0234504.	1.1	5
14	Impacts of the Toba eruption and montane forest expansion on diversification in Sumatran parachuting frogs (<i>Rhacophorus</i>). Molecular Ecology, 2020, 29, 2994-3009.	2.0	4
15	Comparative phylogeography of West African amphibians and reptiles. Evolution; International Journal of Organic Evolution, 2020, 74, 716-724.	1.1	31
16	Phylogeography of montane dragons could shed light on the history of forests and diversification processes on Sumatra. Molecular Phylogenetics and Evolution, 2020, 149, 106840.	1.2	8
17	Exploring rain forest diversification using demographic model testing in the African foamâ€nest treefrog <i>Chiromantis rufescens</i> . Journal of Biogeography, 2019, 46, 2706-2721.	1.4	28
18	Geographic variation in West African <i>Agama picticauda</i> : insights from genetics, morphology and ecology. African Journal of Herpetology, 2019, 68, 33-49.	0.3	3

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19	A Tale of Two Skates: Comparative Phylogeography of North American Skate Species with Implications for Conservation. Copeia, 2019, 107, 297.	1.4	5
20	Mitochondrial genetic variation within and between Holbrookia lacerata lacerata and Holbrookia lacerata subcaudalis, the spot-tailed earless lizards of Texas. Journal of Natural History, 2018, 52, 1017-1027.	0.2	7
21	Synchronous diversification of parachuting frogs (Genus Rhacophorus) on Sumatra and Java. Molecular Phylogenetics and Evolution, 2018, 123, 101-112.	1.2	8
22	Withinâ€island diversification underlies parachuting frog (⟨i⟩Rhacophorus⟨/i⟩) species accumulation on the Sunda Shelf. Journal of Biogeography, 2018, 45, 929-940.	1.4	23
23	Coalescent species delimitation of a Sumatran parachuting frog. Zoologica Scripta, 2018, 47, 33-43.	0.7	4
24	Sky, sea, and forest islands: Diversification in the African leafâ€folding frog <i>Afrixalus paradorsalis</i> (Anura: Hyperoliidae) of the Lower Guineoâ€Congolian rain forest. Journal of Biogeography, 2018, 45, 1781-1794.	1.4	33
25	Insight into the roles of selection in speciation from genomic patterns of divergence and introgression in secondary contact in venomous rattlesnakes. Ecology and Evolution, 2017, 7, 3951-3966.	0.8	34
26	Evaluating mechanisms of diversification in a Guineoâ€Congolian tropical forest frog using demographic model selection. Molecular Ecology, 2017, 26, 5245-5263.	2.0	157
27	Geographical features are the predominant driver of molecular diversification in widely distributed North American whipsnakes. Molecular Ecology, 2017, 26, 5729-5751.	2.0	19
28	Bayesian inference of species diffusion in the West African <i>Agama agama</i> species group (Reptilia,) Tj ETQc	0 0 0 rgB1	「/Overlock 10 22
29	A new species of Gonatodes (Squamata: Sphaerodactylidae) from the western versant of the Cordillera de Mérida, Venezuela. Zootaxa, 2017, 4291, 549.	0.2	5
30	High-coverage sequencing and annotated assembly of the genome of the Australian dragon lizard Pogona vitticeps. GigaScience, 2015, 4, 45.	3.3	97
31	Limitations of Climatic Data for Inferring Species Boundaries: Insights from Speckled Rattlesnakes. PLoS ONE, 2015, 10, e0131435.	1.1	29
32	Genetic divergence and diversity in the Mona and Virgin Islands Boas, Chilabothrus monensis (Epicrates monensis) (Serpentes: Boidae), West Indian snakes of special conservation concern. Molecular Phylogenetics and Evolution, 2015, 88, 144-153.	1,2	9
33	Two Low Coverage Bird Genomes and a Comparison of Reference-Guided versus De Novo Genome Assemblies. PLoS ONE, 2014, 9, e106649.	1.1	30
34	Three crocodilian genomes reveal ancestral patterns of evolution among archosaurs. Science, 2014, 346, 1254449.	6.0	300
35	Two Antarctic penguin genomes reveal insights into their evolutionary history and molecular changes related to the Antarctic environment. GigaScience, 2014, 3, 27.	3.3	72
36	Introgression and Phenotypic Assimilation in Zimmerius Flycatchers (Tyrannidae): Population Genetic and Phylogenetic Inferences from Genome-Wide SNPs. Systematic Biology, 2014, 63, 134-152.	2.7	84

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37	Diversification and asymmetrical gene flow across time and space: lineage sorting and hybridization in polytypic barking frogs. Molecular Ecology, 2014, 23, 3273-3291.	2.0	78
38	Species Delimitation using Genome-Wide SNP Data. Systematic Biology, 2014, 63, 534-542.	2.7	390
39	Geckos: The Animal Answer Guide. By Aaron M. Bauer. Baltimore (Maryland): Johns Hopkins University Press. \$50.00 (hardcover); \$26.95 (paper). xv + 159 p. + 16 pl.; ill.; index. ISBN: 978-1-4214-0852-1 (hc); 978-1-4214-0853-8 (pb); 978-1-4214-0925-2 (eb). 2013 Quarterly Review of Biology, 2014, 89, 191-192.	0.0	O
40	The Burmese python genome reveals the molecular basis for extreme adaptation in snakes. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 20645-20650.	3.3	260
41	Speciation on the Rocks: Integrated Systematics of the Heteronotia spelea Species Complex (Gekkota;) Tj ETQq1	1.0,78431 1.1	.4.rgBT /Ov
42	Report from the First Snake Genomics and Integrative Biology Meeting. Standards in Genomic Sciences, 2012, 7, 150-152.	1.5	4
43	Coalescent-based species delimitation in an integrative taxonomy. Trends in Ecology and Evolution, 2012, 27, 480-488.	4.2	716
44	A Phylogenomic Approach to Vertebrate Phylogeny Supports a Turtle-Archosaur Affinity and a Possible Paraphyletic Lissamphibia. PLoS ONE, 2012, 7, e48990.	1.1	61
45	The genus Astylosternus in the Upper Guinea rainforests, West Africa, with the description of a new species (Amphibia: Anura: Arthroleptidae). Zootaxa, 2012, 3245, 1.	0.2	11
46	The genome of the green anole lizard and a comparative analysis with birds and mammals. Nature, 2011, 477, 587-591.	13.7	575
47	Evaluating phylogenetic informativeness and data-type usage for new protein-coding genes across Vertebrata. Molecular Phylogenetics and Evolution, 2011, 61, 300-307.	1.2	32
48	The genetic legacy of aridification: Climate cycling fostered lizard diversification in Australian montane refugia and left low-lying deserts genetically depauperate. Molecular Phylogenetics and Evolution, 2011, 61, 750-759.	1.2	56
49	Palaeoclimate change drove diversification among isolated mountain refugia in the Australian arid zone. Molecular Ecology, 2011, 20, 1529-1545.	2.0	7 5
50	NONADAPTIVE EVOLUTION OF MITOCHONDRIAL GENOME SIZE. Evolution; International Journal of Organic Evolution, 2011, 65, 2706-2711.	1.1	20
51	Molecular systematics of Stenodactylus (Gekkonidae), an Afro-Arabian gecko species complex. Molecular Phylogenetics and Evolution, 2011, 58, 71-75.	1.2	27
52	A coalescent perspective on delimiting and naming species: a reply to Bauer et al Proceedings of the Royal Society B: Biological Sciences, 2011, 278, 493-495.	1.2	65
53	The Anolis Lizard Genome: An Amniote Genome without Isochores. Genome Biology and Evolution, 2011, 3, 974-984.	1.1	44
54	DIVERSIFICATION AND PERSISTENCE AT THE ARID-MONSOONAL INTERFACE: AUSTRALIA-WIDE BIOGEOGRAPHY OF THE BYNOE'S GECKO (HETERONOTIA BINOEI; GEKKONIDAE). Evolution; International Journal of Organic Evolution, 2010, 64, no-no.	1.1	96

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55	Bayesian species delimitation in West African forest geckos (<i>Hemidactylus fasciatus</i>). Proceedings of the Royal Society B: Biological Sciences, 2010, 277, 3071-3077.	1.2	485
56	Genome Evolution in Reptilia, the Sister Group of Mammals. Annual Review of Genomics and Human Genetics, 2010, 11, 239-264.	2.5	78
57	Multiple Origins and Rapid Evolution of Duplicated Mitochondrial Genes in Parthenogenetic Geckos (Heteronotia binoei; Squamata, Gekkonidae). Molecular Biology and Evolution, 2007, 24, 2775-2786.	3.5	59
58	Turtle phylogeny: insights from a novel nuclear intron. Molecular Phylogenetics and Evolution, 2004, 31, 1031-1040.	1.2	114