

Tod W Reeder

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

39
papers

4,577
citations

27
h-index

40
g-index

40
ext. papers

5,003
ext. citations

4.5
avg, IF

5.6
L-index

#	Paper	IF	Citations
39	A New Diploid Parthenogenetic Whiptail Lizard from Sonora, Mexico, Is the "Missing Link" in the Evolutionary Transition to Polyploidy. <i>American Naturalist</i> , 2021 , 198, 295-309	3.7	1
38	Introgression obscures lineage boundaries and phylogeographic history in the western banded gecko, <i>Coleonyx variegatus</i> (Squamata: Eublepharidae). <i>Zoological Journal of the Linnean Society</i> , 2020 , 190, 181-226	2.4	2
37	Complex patterns of hybridization and introgression across evolutionary timescales in Mexican whiptail lizards (<i>Aspidoscelis</i>). <i>Molecular Phylogenetics and Evolution</i> , 2019 , 132, 284-295	4.1	11
36	Pleistocene climatic fluctuations drive isolation and secondary contact in the red diamond rattlesnake (<i>Crotalus ruber</i>) in Baja California. <i>Journal of Biogeography</i> , 2018 , 45, 64-75	4.1	11
35	Multilocus phylogeny of alligator lizards (<i>Elgaria</i> , Anguillidae): Testing mtDNA introgression as the source of discordant molecular phylogenetic hypotheses. <i>Molecular Phylogenetics and Evolution</i> , 2017 , 110, 104-121	4.1	22
34	Phylogenetic inference and divergence dating of snakes using molecules, morphology and fossils: new insights into convergent evolution of feeding morphology and limb reduction. <i>Biological Journal of the Linnean Society</i> , 2017 , 121, 379-394	1.9	40
33	Lineage diversification of fringe-toed lizards (Phrynosomatidae: <i>Uma notata</i> complex) in the Colorado Desert: Delimiting species in the presence of gene flow. <i>Molecular Phylogenetics and Evolution</i> , 2017 , 106, 103-117	4.1	22
32	Squamate Phylogenetics, Molecular Branch Lengths, and Molecular Apomorphies: A Response to McMahan et al.. <i>Copeia</i> , 2016 , 104, 702-707	1.1	7
31	Biogeographical history and coalescent species delimitation of Pacific island skinks (Squamata: Scincidae: <i>Emoia cyanura</i> species group). <i>Journal of Biogeography</i> , 2016 , 43, 1917-1929	4.1	10
30	When do species-tree and concatenated estimates disagree? An empirical analysis with higher-level scincid lizard phylogeny. <i>Molecular Phylogenetics and Evolution</i> , 2015 , 82 Pt A, 146-55	4.1	56
29	Integrated analyses resolve conflicts over squamate reptile phylogeny and reveal unexpected placements for fossil taxa. <i>PLoS ONE</i> , 2015 , 10, e0118199	3.7	182
28	Species delimitation using Bayes factors: simulations and application to the <i>Sceloporus scalaris</i> species group (Squamata: Phrynosomatidae). <i>Systematic Biology</i> , 2014 , 63, 119-33	8.4	192
27	Contrasting global-scale evolutionary radiations: phylogeny, diversification, and morphological evolution in the major clades of iguanian lizards. <i>Biological Journal of the Linnean Society</i> , 2013 , 108, 127-143	1.9	27
26	Resolving the phylogeny of lizards and snakes (Squamata) with extensive sampling of genes and species. <i>Biology Letters</i> , 2012 , 8, 1043-6	3.6	231
25	Estimating divergence dates and evaluating dating methods using phylogenomic and mitochondrial data in squamate reptiles. <i>Molecular Phylogenetics and Evolution</i> , 2012 , 65, 974-91	4.1	82
24	Phylogenetic Insights on Evolutionary Novelty in Lizards and Snakes: Sex, Birth, Bodies, Niches, and Venom. <i>Annual Review of Ecology, Evolution, and Systematics</i> , 2011 , 42, 227-244	13.5	68
23	Phylogeny of iguanian lizards inferred from 29 nuclear loci, and a comparison of concatenated and species-tree approaches for an ancient, rapid radiation. <i>Molecular Phylogenetics and Evolution</i> , 2011 , 61, 363-80	4.1	159

22	Phylogenetic Relationships within the Australian Limb-Reduced Lizard Genus <i>Hemiergis</i> (Scincidae: Squamata) as Inferred from the Bayesian Analysis of Mitochondrial rRNA Gene Sequences. <i>Copeia</i> , 2011 , 2011, 113-120	1.1	1
21	Intercontinental dispersal by a microendemic burrowing reptile (Dibamidae). <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2011 , 278, 2568-74	4.4	22
20	Combining phylogenomics and fossils in higher-level squamate reptile phylogeny: molecular data change the placement of fossil taxa. <i>Systematic Biology</i> , 2010 , 59, 674-88	8.4	173
19	Phylogenetic relationships of phrynosomatid lizards based on nuclear and mitochondrial data, and a revised phylogeny for <i>Sceloporus</i> . <i>Molecular Phylogenetics and Evolution</i> , 2010 , 54, 150-61	4.1	93
18	Branch lengths, support, and congruence: testing the phylogenomic approach with 20 nuclear loci in snakes. <i>Systematic Biology</i> , 2008 , 57, 420-31	8.4	153
17	Novel patterns of historical isolation, dispersal, and secondary contact across Baja California in the Rosy Boa (<i>Lichanura trivirgata</i>). <i>Molecular Phylogenetics and Evolution</i> , 2008 , 46, 484-502	4.1	27
16	Rapid development of multiple nuclear loci for phylogenetic analysis using genomic resources: an example from squamate reptiles. <i>Molecular Phylogenetics and Evolution</i> , 2008 , 47, 129-42	4.1	196
15	Phylogenetic Affinities of the Rare and Enigmatic Limb-Reduced <i>Anelytropsis</i> (Reptilia: Squamata) as Inferred with Mitochondrial 16S rRNA Sequence Data. <i>Journal of Herpetology</i> , 2008 , 42, 303-311	1.1	4
14	A phylogenetic perspective on elevational species richness patterns in Middle American treefrogs: why so few species in lowland tropical rainforests?. <i>Evolution; International Journal of Organic Evolution</i> , 2007 , 61, 1188-207	3.8	103
13	Loss and re-evolution of complex life cycles in marsupial frogs: does ancestral trait reconstruction mislead?. <i>Evolution; International Journal of Organic Evolution</i> , 2007 , 61, 1886-99	3.8	83
12	WHY DOES A TRAIT EVOLVE MULTIPLE TIMES WITHIN A CLADE? REPEATED EVOLUTION OF SNAKELIKE BODY FORM IN SQUAMATE REPTILES. <i>Evolution; International Journal of Organic Evolution</i> , 2006 , 60, 123	3.8	9
11	Evolutionary and ecological causes of the latitudinal diversity gradient in hylid frogs: treefrog trees unearth the roots of high tropical diversity. <i>American Naturalist</i> , 2006 , 168, 579-96	3.7	323
10	WHY DOES A TRAIT EVOLVE MULTIPLE TIMES WITHIN A CLADE? REPEATED EVOLUTION OF SNAKELINE BODY FORM IN SQUAMATE REPTILES. <i>Evolution; International Journal of Organic Evolution</i> , 2006 , 60, 123-141	3.8	220
9	Why does a trait evolve multiple times within a clade? Repeated evolution of snakelike body form in squamate reptiles. <i>Evolution; International Journal of Organic Evolution</i> , 2006 , 60, 123-41	3.8	224
8	Hylid frog phylogeny and sampling strategies for speciose clades. <i>Systematic Biology</i> , 2005 , 54, 778-807	8.4	252
7	Partitioned Bayesian analyses, partition choice, and the phylogenetic relationships of scincid lizards. <i>Systematic Biology</i> , 2005 , 54, 373-90	8.4	598
6	A phylogeny of the Australian <i>Sphenomorphus</i> group (Scincidae: Squamata) and the phylogenetic placement of the crocodile skinks (<i>Tribolonotus</i>): Bayesian approaches to assessing congruence and obtaining confidence in maximum likelihood inferred relationships. <i>Molecular Phylogenetics and Evolution</i> , 2003 , 27, 384-97	4.1	120
5	Evidence for parallel ecological speciation in scincid lizards of the <i>Eumeces skiltonianus</i> species group (Squamata: Scincidae). <i>Evolution; International Journal of Organic Evolution</i> , 2002 , 56, 1498-513	3.8	62

4	Molecular systematics of the Eastern Fence Lizard (<i>Sceloporus undulatus</i>): a comparison of Parsimony, Likelihood, and Bayesian approaches. <i>Systematic Biology</i> , 2002 , 51, 44-68	8.4	501
3	MOLECULAR PHYLOGENETICS AND EVOLUTION OF SEXUAL DICHROMATISM AMONG POPULATIONS OF THE YARROWS SPINY LIZARD (<i>SCELOPORUS JARROVII</i>). <i>Evolution; International Journal of Organic Evolution</i> , 1999 , 53, 1884-1897	3.8	39
2	Phylogeny of the Spiny Lizards (<i>Sceloporus</i>) Based on Molecular and Morphological Evidence. <i>Herpetological Monographs</i> , 1997 , 11, 1	1.5	143
1	Combining Data Sets with Different Numbers of Taxa for Phylogenetic Analysis. <i>Systematic Biology</i> , 1995 , 44, 548-558	8.4	108