

# Tod W Reeder

## List of Publications by Citations

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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	Partitioned Bayesian analyses, partition choice, and the phylogenetic relationships of scincid lizards. <i>Systematic Biology</i> , <b>2005</b> , 54, 373-90	8.4	598
38	Molecular systematics of the Eastern Fence Lizard ( <i>Sceloporus undulatus</i> ): a comparison of Parsimony, Likelihood, and Bayesian approaches. <i>Systematic Biology</i> , <b>2002</b> , 51, 44-68	8.4	501
37	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> , <b>2006</b> , 168, 579-96	3.7	323
36	Hylid frog phylogeny and sampling strategies for speciose clades. <i>Systematic Biology</i> , <b>2005</b> , 54, 778-807	8.4	252
35	Resolving the phylogeny of lizards and snakes (Squamata) with extensive sampling of genes and species. <i>Biology Letters</i> , <b>2012</b> , 8, 1043-6	3.6	231
34	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> , <b>2006</b> , 60, 123-41	3.8	224
33	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> , <b>2006</b> , 60, 123-141	3.8	220
32	Rapid development of multiple nuclear loci for phylogenetic analysis using genomic resources: an example from squamate reptiles. <i>Molecular Phylogenetics and Evolution</i> , <b>2008</b> , 47, 129-42	4.1	196
31	Species delimitation using Bayes factors: simulations and application to the <i>Sceloporus scalaris</i> species group (Squamata: Phrynosomatidae). <i>Systematic Biology</i> , <b>2014</b> , 63, 119-33	8.4	192
30	Integrated analyses resolve conflicts over squamate reptile phylogeny and reveal unexpected placements for fossil taxa. <i>PLoS ONE</i> , <b>2015</b> , 10, e0118199	3.7	182
29	Combining phylogenomics and fossils in higher-level squamate reptile phylogeny: molecular data change the placement of fossil taxa. <i>Systematic Biology</i> , <b>2010</b> , 59, 674-88	8.4	173
28	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> , <b>2011</b> , 61, 363-80	4.1	159
27	Branch lengths, support, and congruence: testing the phylogenomic approach with 20 nuclear loci in snakes. <i>Systematic Biology</i> , <b>2008</b> , 57, 420-31	8.4	153
26	Phylogeny of the Spiny Lizards ( <i>Sceloporus</i> ) Based on Molecular and Morphological Evidence. <i>Herpetological Monographs</i> , <b>1997</b> , 11, 1	1.5	143
25	A phylogeny of the Australian Sphenomorphus 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> , <b>2009</b> , 27, 281-27	4.1	120
24	Combining Data Sets with Different Numbers of Taxa for Phylogenetic Analysis. <i>Systematic Biology</i> , <b>1995</b> , 44, 548-558	8.4	108
23	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> , <b>2007</b> , 61, 1188-207	3.8	103

22	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> , <b>2010</b> , 54, 150-61	4.1	93
21	Loss and re-evolution of complex life cycles in marsupial frogs: does ancestral trait reconstruction mislead?. <i>Evolution; International Journal of Organic Evolution</i> , <b>2007</b> , 61, 1886-99	3.8	83
20	Estimating divergence dates and evaluating dating methods using phylogenomic and mitochondrial data in squamate reptiles. <i>Molecular Phylogenetics and Evolution</i> , <b>2012</b> , 65, 974-91	4.1	82
19	Phylogenetic Insights on Evolutionary Novelties in Lizards and Snakes: Sex, Birth, Bodies, Niches, and Venom. <i>Annual Review of Ecology, Evolution, and Systematics</i> , <b>2011</b> , 42, 227-244	13.5	68
18	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> , <b>2002</b> , 56, 1498-513	3.8	62
17	When do species-tree and concatenated estimates disagree? An empirical analysis with higher-level scincid lizard phylogeny. <i>Molecular Phylogenetics and Evolution</i> , <b>2015</b> , 82 Pt A, 146-55	4.1	56
16	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> , <b>2017</b> , 121, 379-394	1.9	40
15	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> , <b>1999</b> , 53, 1884-1897	3.8	39
14	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> , <b>2013</b> , 108, 127-143	1.9	27
13	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> , <b>2008</b> , 46, 484-502	4.1	27
12	Multilocus phylogeny of alligator lizards ( <i>Elgaria</i> , Anguidae): Testing mtDNA introgression as the source of discordant molecular phylogenetic hypotheses. <i>Molecular Phylogenetics and Evolution</i> , <b>2017</b> , 110, 104-121	4.1	22
11	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> , <b>2017</b> , 106, 103-117	4.1	22
10	Intercontinental dispersal by a microendemic burrowing reptile (Dibamidae). <i>Proceedings of the Royal Society B: Biological Sciences</i> , <b>2011</b> , 278, 2568-74	4.4	22
9	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> , <b>2018</b> , 45, 64-75	4.1	11
8	Complex patterns of hybridization and introgression across evolutionary timescales in Mexican whiptail lizards ( <i>Aspidoscelis</i> ). <i>Molecular Phylogenetics and Evolution</i> , <b>2019</b> , 132, 284-295	4.1	11
7	Biogeographical history and coalescent species delimitation of Pacific island skinks (Squamata: Scincidae: <i>Emoia cyanura</i> species group). <i>Journal of Biogeography</i> , <b>2016</b> , 43, 1917-1929	4.1	10
6	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> , <b>2006</b> , 60, 123	3.8	9
5	Squamate Phylogenetics, Molecular Branch Lengths, and Molecular Apomorphies: A Response to McMahan et al.. <i>Copeia</i> , <b>2016</b> , 104, 702-707	1.1	7

4	Phylogenetic Affinities of the Rare and Enigmatic Limb-Reduced Anelytropsis (Reptilia: Squamata) as Inferred with Mitochondrial 16S rRNA Sequence Data. <i>Journal of Herpetology</i> , <b>2008</b> , 42, 303-311	1.1	4
3	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> , <b>2020</b> , 190, 181-226	2.4	2
2	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> , <b>2011</b> , 2011, 113-120	1.1	1
1	A New Diploid Parthenogenetic Whiptail Lizard from Sonora, Mexico, Is the "Missing Link" in the Evolutionary Transition to Polyploidy. <i>American Naturalist</i> , <b>2021</b> , 198, 295-309	3.7	1