

# Anna Papadopoulou

## List of Publications by Year in descending order

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Version: 2024-02-01

28  
papers

2,708  
citations

471477  
17  
h-index

477281  
29  
g-index

30  
all docs

30  
docs citations

30  
times ranked

4138  
citing authors

#	ARTICLE	IF	CITATIONS
1	A Comprehensive Phylogeny of Beetles Reveals the Evolutionary Origins of a Superradiation. <i>Science</i> , 2007, 318, 1913-1916.	12.6	729
2	Revisiting the Insect Mitochondrial Molecular Clock: The Mid-Aegean Trench Calibration. <i>Molecular Biology and Evolution</i> , 2010, 27, 1659-1672.	8.9	729
3	Toward a paradigm shift in comparative phylogeography driven by trait-based hypotheses. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 8018-8024.	7.1	170
4	A roadmap for island biology: 50 fundamental questions after 50Âyears of <i>The Theory of Island Biogeography</i>. <i>Journal of Biogeography</i> , 2017, 44, 963-983.	3.0	167
5	Comparative phylogeography of tenebrionid beetles in the Aegean archipelago: the effect of dispersal ability and habitat preference. <i>Molecular Ecology</i> , 2009, 18, 2503-2517.	3.9	119
6	Speciation and DNA barcodes: testing the effects of dispersal on the formation of discrete sequence clusters. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2008, 363, 2987-2996.	4.0	104
7	Genomic tests of the species-pump hypothesis: Recent island connectivity cycles drive population divergence but not speciation in Caribbean crickets across the Virgin Islands. <i>Evolution; International Journal of Organic Evolution</i> , 2015, 69, 1501-1517.	2.3	88
8	Dense Taxonomic EST Sampling and Its Applications for Molecular Systematics of the Coleoptera (Beetles). <i>Molecular Biology and Evolution</i> , 2006, 23, 268-278.	8.9	86
9	Testing the Speciesâ€“Genetic Diversity Correlation in the Aegean Archipelago: Toward a Haplotype-Based Macroecology?. <i>American Naturalist</i> , 2011, 178, 241-255.	2.1	86
10	Speciesâ€“specific responses to island connectivity cycles: refined models for testing phylogeographic concordance across a <scp>M</scp>editerranean <scp>P</scp>leistocene <scp>A</scp>ggreate <scp>I</scp>sland <scp>C</scp>omplex. <i>Molecular Ecology</i> , 2015, 24, 4252-4268.	3.9	67
11	Sampling Error Does Not Invalidate the Yule-Coalescent Model for Species Delimitation. A Response to Lohse (2009). <i>Systematic Biology</i> , 2009, 58, 442-444.	5.6	59
12	Metagenome skimming for phylogenetic community ecology: a new era in biodiversity research. <i>Molecular Ecology</i> , 2015, 24, 3515-3517.	3.9	34
13	Glacial refugia, recolonization patterns and diversification forces in Alpineâ€“endemic <i>Megabunus</i> harvestmen. <i>Molecular Ecology</i> , 2016, 25, 2904-2919.	3.9	34
14	DNA taxonomy and phylogeography of beetles of the Falkland Islands (Islas Malvinas). <i>Molecular Phylogenetics and Evolution</i> , 2009, 53, 935-947.	2.7	27
15	Diversity and diversification of Eumolpinae (Coleoptera: Chrysomelidae) in New Caledonia. <i>Zoological Journal of the Linnean Society</i> , 2013, 168, 473-495.	2.3	27
16	Linking microâ€“and macroevolutionary perspectives to evaluate the role of Quaternary seaâ€“level oscillations in island diversification. <i>Evolution; International Journal of Organic Evolution</i> , 2017, 71, 2901-2917.	2.3	25
17	Genomic data reveal deep genetic structure but no support for current taxonomic designation in a grasshopper species complex. <i>Molecular Ecology</i> , 2019, 28, 3869-3886.	3.9	23
18	Ecological constraints from incumbent clades drive trait evolution across the treeâ€“ofâ€“life of freshwater macroinvertebrates. <i>Ecography</i> , 2018, 41, 1049-1063.	4.5	21

#	ARTICLE	IF	CITATIONS
19	Automated <scp>DNA</scp>-based plant identification for large-scale biodiversity assessment. Molecular Ecology Resources, 2015, 15, 136-152.	4.8	17
20	Genomic footprints of an old affair: Single nucleotide polymorphism data reveal historical hybridization and the subsequent evolution of reproductive barriers in two recently diverged grasshoppers with partly overlapping distributions. Molecular Ecology, 2020, 29, 2254-2268.	3.9	17
21	High-throughput biodiversity analysis: Rapid assessment of species richness and ecological interactions of Chrysomelidae (Coleoptera) in the tropics. ZooKeys, 2016, 597, 3-26.	1.1	15
22	Community metabarcoding reveals the relative role of environmental filtering and spatial processes in metacommunity dynamics of soil microarthropods across a mosaic of montane forests. Molecular Ecology, 2023, 32, 6110-6128.	3.9	15
23	Evaluation of bias on the assessment of diet breadth of herbivorous insects using molecular methods. Insect Science, 2017, 24, 194-209.	3.0	12
24	&lt;p&gt;&lt;strong&gt;A new species of striped &lt;em&gt;Ichthyophis&lt;/em&gt; Fitzinger, 1826 &lt;/strong&gt;&lt;br /&gt;&lt;strong&gt;(Amphibia: Gymnophiona: Ichthyophiidae) from Myanmar&lt;/strong&gt;&lt;/p&gt;. Zootaxa, 2014, 3785, 45.	0.5	11
25	Dispersal versus vicariance in the Aegean: combining molecular and morphological phylogenies of eastern Mediterranean <i>Dendarus</i> (Coleoptera: Tenebrionidae) sheds new light on the phylogeography of the Aegean area. Zoological Journal of the Linnean Society, 2020, 190, 824-843.	2.3	9
26	The systematics of Boulengerula fischeri (Amphibia: Gymnophiona: Caeciliidae) based on morphological and molecular data. Zootaxa, 2011, 2767, 14.	0.5	6
27	A second horizon scan of biogeography: Golden Ages, Midas touches, and the Red Queen. Frontiers of Biogeography, 2016, 8, .	1.8	3
28	A key to Russian and Eastern European species of Blaps Fabricius, 1775 (Coleoptera: Tenebrionidae:) Tj ETQq0 0 0 rgBT /Overlock 10 Tf morphological and molecular data. Zootaxa, 2022, 5116, 267-291.	0.5	1