

Johan A A Nylander

List of Publications by Year in descending order

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Version: 2024-02-01

19
papers

5,220
citations

471371

17
h-index

839398

18
g-index

19
all docs

19
docs citations

19
times ranked

6617
citing authors

#	ARTICLE	IF	CITATIONS
1	Metagenomic Analysis of the Indian Ocean Picocyanobacterial Community: Structure, Potential Function and Evolution. PLoS ONE, 2016, 11, e0155757.	1.1	54
2	Phylogeny, Evolution and Classification of Gall Wasps: The Plot Thickens. PLoS ONE, 2015, 10, e0123301.	1.1	136
3	Local hopping mobile DNA implicated in pseudogene formation and reductive evolution in an obligate cyanobacteria-plant symbiosis. BMC Genomics, 2015, 16, 193.	1.2	20
4	Functional Tradeoffs Underpin Salinity-Driven Divergence in Microbial Community Composition. PLoS ONE, 2014, 9, e89549.	1.1	184
5	An evaluation of new parsimony-based versus parametric inference methods in biogeography: a case study using the globally distributed plant family Sapindaceae. Journal of Biogeography, 2011, 38, 531-550.	1.4	171
6	The phylogenetic placement and biogeographical origins of the New Zealand stick insects (Phasmatodea). Systematic Entomology, 2010, 35, 207-225.	1.7	80
7	Biogeographical history of cuckoo-shrikes (Aves: Passeriformes): transoceanic colonization of Africa from Australo-Papua. Journal of Biogeography, 2010, 37, 1767-1781.	1.4	37
8	Genome Erosion in a Nitrogen-Fixing Vertically Transmitted Endosymbiotic Multicellular Cyanobacterium. PLoS ONE, 2010, 5, e11486.	1.1	178
9	Revision and phylogenetics of the genus Paraulax Kieffer (Hymenoptera, Cynipidae) with biological notes and description of a new tribe, a new genus, and five new species. Zootaxa, 2009, 2200, 1-40.	0.2	33
10	Plastid and nuclear DNA markers reveal intricate relationships at subfamilial and tribal levels in the soapberry family (Sapindaceae). Molecular Phylogenetics and Evolution, 2009, 51, 238-258.	1.2	131
11	Tracing the impact of the Andean uplift on Neotropical plant evolution. Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 9749-9754.	3.3	550
12	Accounting for Phylogenetic Uncertainty in Biogeography: A Bayesian Approach to Dispersal-Vicariance Analysis of the Thrushes (Aves: Turdus). Systematic Biology, 2008, 57, 257-268.	2.7	336
13	AWTY (are we there yet?): a system for graphical exploration of MCMC convergence in Bayesian phylogenetics. Bioinformatics, 2008, 24, 581-583.	1.8	1,390
14	The phylogeny and evolution of Figitidae (Hymenoptera: Cynipoidea). Cladistics, 2007, 23, 403-431.	1.5	79
15	Phylogenetic relationships of typical antbirds (Thamnophilidae) and test of incongruence based on Bayes factors. BMC Evolutionary Biology, 2004, 4, 23.	3.2	57
16	Bayesian Phylogenetic Analysis of Combined Data. Systematic Biology, 2004, 53, 47-67.	2.7	1,636
17	A Maximum-Likelihood Analysis of Eight Phylogenetic Markers in Gallwasps (Hymenoptera: Cynipidae): Implications for Insect Phylogenetic Studies. Molecular Phylogenetics and Evolution, 2002, 22, 206-219.	1.2	98
18	A test of monophyly of the gutless Phalloporinae (Oligochaeta, Tubificidae) and the use of a 573-bp region of the mitochondrial cytochrome oxidase I gene in analysis of annelid phylogeny. Zoologica Scripta, 1999, 28, 305-313.	0.7	46

#	ARTICLE	IF	CITATIONS
19	Life history of Parnips and the evolutionary origin of gall wasps. Journal of Hymenoptera Research, 0, 65, 91-110.	0.8	4