

Olaf Rp: Bininda-Emonds

List of Publications by Year in descending order

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114
papers

14,690
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41344

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117
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117
times ranked

15901
citing authors

#	ARTICLE	IF	CITATIONS
1	18S rRNA variability maps reveal three highly divergent, conserved motifs within Rotifera. <i>Bmc Ecology and Evolution</i> , 2021, 21, 118.	1.6	6
2	Lethal effects and ultrastructure of cellular uptake of ingested gold nanoparticles in the freshwater rotifer <i>Brachionus calyciflorus</i> (Monogononta: Brachionidae). <i>Environmental Pollution</i> , 2021, 289, 117897.	7.5	5
3	The evolution of Synchaetidae (Rotifera: Monogononta) with a focus on <i>Synchaeta</i> : An integrative approach combining molecular and morphological data. <i>Journal of Zoological Systematics and Evolutionary Research</i> , 2020, 58, 823-857.	1.4	7
4	Is the valid species <i>Synchaeta monopus</i> Plate, 1889 (Rotifera: Monogononta) a product of preparation artefacts?. <i>Journal of Natural History</i> , 2019, 53, 413-423.	0.5	3
5	Cloning of a novel trypsin inhibitor from the Traditional Chinese medicine decoction pieces, <i>Radix Trichosanthis</i> . <i>Analytical Biochemistry</i> , 2019, 578, 23-28.	2.4	1
6	A weighted taxonomic matrix key for species of the rotifer genus <i>Synchaeta</i> (Rotifera, Monogononta). <i>Tj ETQq0 0 Q rrgBT /Overlock 10 T</i>	1.1	6
7	A comprehensive and integrative re-description of <i>Synchaeta oblonga</i> and its relationship to <i>Synchaeta tremula</i> , <i>Synchaeta rufina</i> and <i>Synchaeta littoralis</i> (Rotifera: Monogononta). <i>Organisms Diversity and Evolution</i> , 2018, 18, 407-423.	1.6	5
8	Immediate and heritable costs of desiccation on the life history of the bdelloid rotifer <i>Philodina roseola</i> . <i>Organisms Diversity and Evolution</i> , 2018, 18, 399-406.	1.6	2
9	On the importance of robust species descriptions for Rotifera: re-descriptions of <i>Synchaeta stylata</i> and <i>Synchaeta longipes</i> and a comparison to <i>Synchaeta jollyae</i> . <i>Zoologischer Anzeiger</i> , 2018, 277, 42-54.	0.9	4
10	A comprehensive and integrative re-description of <i>Synchaeta tremula</i> (Müller, 1786) and the newly rediscovered <i>Synchaeta tremuloida</i> Pourriot, 1965 (Rotifera: Synchaetidae). <i>Zootaxa</i> , 2017, 4276, 503.	0.5	4
11	Phenotypic influences on the reproductive strategy of the facultative sexual rotifer <i>Brachionus rubens</i> (Monogononta). <i>Organisms Diversity and Evolution</i> , 2017, 17, 779-788.	1.6	2
12	Does retinal configuration make the head and eyes of foveate birds move?. <i>Scientific Reports</i> , 2017, 7, 38406.	3.3	25
13	Evidence Supporting the Uptake and Genomic Incorporation of Environmental DNA in the Ancient Asexual Bdelloid Rotifer <i>Philodina roseola</i> . <i>Life</i> , 2016, 6, 38.	2.4	8
14	Population structuring in the monogonont rotifer <i>Synchaeta pectinata</i> : high genetic divergence on a small geographical scale. <i>Freshwater Biology</i> , 2015, 60, 1364-1378.	2.4	15
15	The diversity and evolution of anuran skin peptides. <i>Peptides</i> , 2015, 63, 96-117.	2.4	126
16	Phylogeographical analysis of <i>Ligia oceanica</i> (Crustacea: Isopoda) reveals two deeply divergent mitochondrial lineages. <i>Biological Journal of the Linnean Society</i> , 2014, 112, 16-30.	1.6	17
17	Phylogenetically related and ecologically similar carnivores harbour similar parasite assemblages. <i>Journal of Animal Ecology</i> , 2014, 83, 671-680.	2.8	74
18	Evolution of the Cation Chloride Cotransporter Family: Ancient Origins, Gene Losses, and Subfunctionalization through Duplication. <i>Molecular Biology and Evolution</i> , 2014, 31, 434-447.	8.9	54

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19	An Introduction to Supertree Construction (and Partitioned Phylogenetic Analyses) with a View Toward the Distinction Between Gene Trees and Species Trees. , 2014, , 49-76.		10
20	Unraveling the origin of Cladocera by identifying heterochrony in the developmental sequences of Branchiopoda. <i>Frontiers in Zoology</i> , 2013, 10, 35.	2.0	22
21	Molecular cloning of the trypsin inhibitor from the skin secretion of the Madagascan Tomato Frog, <i>Dyscophus guineti</i> (Microhylidae), and insights into its potential defensive role. <i>Organisms Diversity and Evolution</i> , 2013, 13, 453-461.	1.6	3
22	How does the "ancient" asexual <i>Philodina roseola</i> (Rotifera: Bdelloidea) handle potential UVB-induced mutations?. <i>Journal of Experimental Biology</i> , 2013, 216, 3090-5.	1.7	12
23	Amalgamating Source Trees with Different Taxonomic Levels. <i>Systematic Biology</i> , 2013, 62, 231-249.	5.6	17
24	Time-dependent Gene Expression Analysis of the Developing Superior Olivary Complex. <i>Journal of Biological Chemistry</i> , 2013, 288, 25865-25879.	3.4	32
25	Response to Comment on "Impacts of the Cretaceous Terrestrial Revolution and KPg Extinction on Mammal Diversification". <i>Science</i> , 2012, 337, 34-34.	12.6	2
26	Are Voluntary Wheel Running and Open-Field Behavior Correlated in Mice? Different Answers from Comparative and Artificial Selection Approaches. <i>Behavior Genetics</i> , 2012, 42, 830-844.	2.1	41
27	Species status and population structure of mussels (Mollusca: Bivalvia: <i>Mytilus</i> spp.) in the Wadden Sea of Lower Saxony (Germany). <i>Organisms Diversity and Evolution</i> , 2012, 12, 387-402.	1.6	10
28	Comment on "Impacts of the Cretaceous Terrestrial Revolution and KPg Extinction on Mammal Diversification". <i>Science</i> , 2012, 337, 34-34.	12.6	4
29	Antimicrobial peptides and alytesin are co-secreted from the venom of the Midwife toad, <i>Alytes maurus</i> (Alytidae, Anura): Implications for the evolution of frog skin defensive secretions. <i>Toxicon</i> , 2012, 60, 967-981.	1.6	21
30	Molecular cloning of skin peptide precursor-encoding cDNAs from tibial gland secretion of the Giant Monkey Frog, <i>Phyllomedusa bicolor</i> (Hylidae, Anura). <i>Peptides</i> , 2012, 38, 371-376.	2.4	11
31	Molecular time scale of diversification of feeding strategy and morphology in New World Leaf-Nosed Bats (Phyllostomidae): a phylogenetic perspective. , 2012, , 385-409.		64
32	A novel method for comparative analysis of retinal specialization traits from topographic maps. <i>Journal of Vision</i> , 2012, 12, 13-13.	0.3	30
33	Updating the evolutionary history of Carnivora (Mammalia): a new species-level supertree complete with divergence time estimates. <i>BMC Biology</i> , 2012, 10, 12.	3.8	354
34	Complete, accurate, mammalian phylogenies aid conservation planning, but not much. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2011, 366, 2652-2660.	4.0	59
35	Evidence for convergent evolution in the antimicrobial peptide system in anuran amphibians. <i>Peptides</i> , 2011, 32, 20-25.	2.4	34
36	The genome of <i>Oryctes rhinoceros</i> nudivirus provides novel insight into the evolution of nuclear arthropod-specific large circular double-stranded DNA viruses. <i>Virus Genes</i> , 2011, 42, 444-456.	1.6	53

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37	Supporting species in ODE: explaining and citing. <i>Organisms Diversity and Evolution</i> , 2011, 11, 1-2.	1.6	10
38	Inferring the Tree of Life: chopping a phylogenomic problem down to size?. <i>BMC Biology</i> , 2011, 9, 59.	3.8	3
39	Quantifying the Phylodynamic Forces Driving Papillomavirus Evolution. <i>Molecular Biology and Evolution</i> , 2011, 28, 2101-2113.	8.9	114
40	Automated Removal of Noisy Data in Phylogenomic Analyses. <i>Journal of Molecular Evolution</i> , 2010, 71, 319-331.	1.8	61
41	A decade of ODE: looking back and looking forward. <i>Organisms Diversity and Evolution</i> , 2010, 10, 1-4.	1.6	1
42	How Many Bootstrap Replicates Are Necessary?. <i>Journal of Computational Biology</i> , 2010, 17, 337-354.	1.6	800
43	A comprehensive phylogeny of extant horses, rhinos and tapirs (Perissodactyla) through data combination. <i>Zoosystematics and Evolution</i> , 2009, 85, 277-292.	1.1	21
44	Exploration strategies map along fast-slow metabolic and life-history continua in muroid rodents. <i>Functional Ecology</i> , 2009, 23, 150-156.	3.6	127
45	Geographical variation in predictors of mammalian extinction risk: big is bad, but only in the tropics. <i>Ecology Letters</i> , 2009, 12, 538-549.	6.4	496
46	A phylogenetic supertree of the fowls (Galloanserae, Aves). <i>Zoologica Scripta</i> , 2009, 38, 465-481.	1.7	49
47	Phylogenetics, evolution, and medical importance of polyomaviruses. <i>Infection, Genetics and Evolution</i> , 2009, 9, 784-799.	2.3	59
48	How Many Bootstrap Replicates Are Necessary?. <i>Lecture Notes in Computer Science</i> , 2009, , 184-200.	1.3	263
49	PanTHERIA: a species-level database of life history, ecology, and geography of extant and recently extinct mammals. <i>Ecology</i> , 2009, 90, 2648-2648.	3.2	1,322
50	Evolution of four BK virus subtypes. <i>Infection, Genetics and Evolution</i> , 2008, 8, 632-643.	2.3	43
51	Correlates of substitution rate variation in mammalian protein-coding sequences. <i>BMC Evolutionary Biology</i> , 2008, 8, 53.	3.2	146
52	A genus-level supertree of Adephaga (Coleoptera). <i>Organisms Diversity and Evolution</i> , 2008, 7, 255-269.	1.6	33
53	Can heterochrony help explain the high morphological diversity within the genus <i>Niphargus</i> (Crustacea: Amphipoda)?. <i>Organisms Diversity and Evolution</i> , 2008, 8, 146-162.	1.6	21
54	Phylogenetic trees and the future of mammalian biodiversity. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 11556-11563.	7.1	131

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55	Fast Genes and Slow Clades: Comparative Rates of Molecular Evolution in Mammals. <i>Evolutionary Bioinformatics</i> , 2007, 3, 117693430700300.	1.2	26
56	The Fast-Slow Continuum in Mammalian Life History: An Empirical Reevaluation. <i>American Naturalist</i> , 2007, 169, 748-757.	2.1	343
57	When genes meet nomenclature: Tortoise phylogeny and the shifting generic concepts of Testudo and Geochelone. <i>Zoology</i> , 2007, 110, 298-307.	1.2	73
58	The delayed rise of present-day mammals. <i>Nature</i> , 2007, 446, 507-512.	27.8	1,832
59	An integrative approach identifies developmental sequence heterochronies in freshwater basommatophoran snails. <i>Evolution & Development</i> , 2007, 9, 122-130.	2.0	33
60	Forelimb-hindlimb developmental timing changes across tetrapod phylogeny. <i>BMC Evolutionary Biology</i> , 2007, 7, 182.	3.2	93
61	Phylogeny and divergence of the pinnipeds (Carnivora: Mammalia) assessed using a multigene dataset. <i>BMC Evolutionary Biology</i> , 2007, 7, 216.	3.2	166
62	Fast genes and slow clades: comparative rates of molecular evolution in mammals. <i>Evolutionary Bioinformatics</i> , 2007, 3, 59-85.	1.2	26
63	A higher-level MRP supertree of placental mammals. <i>BMC Evolutionary Biology</i> , 2006, 6, 93.	3.2	97
64	BATS, CLOCKS, AND ROCKS: DIVERSIFICATION PATTERNS IN CHIROPTERA. <i>Evolution; International Journal of Organic Evolution</i> , 2005, 59, 2243-2255.	2.3	135
65	transAlign: using amino acids to facilitate the multiple alignment of protein-coding DNA sequences. <i>BMC Bioinformatics</i> , 2005, 6, 156.	2.6	185
66	A complete phylogeny of the whales, dolphins and even-toed hoofed mammals (Cetartiodactyla). <i>Biological Reviews</i> , 2005, 80, 445-473.	10.4	234
67	Getting to the Roots of Matrix Representation. <i>Systematic Biology</i> , 2005, 54, 668-672.	5.6	33
68	Supertree Construction in the Genomic Age. <i>Methods in Enzymology</i> , 2005, 395, 745-757.	1.0	27
69	A New Technique for Identifying Sequence Heterochrony. <i>Systematic Biology</i> , 2005, 54, 230-240.	5.6	106
70	Multiple Causes of High Extinction Risk in Large Mammal Species. <i>Science</i> , 2005, 309, 1239-1241.	12.6	1,035
71	BATS, CLOCKS, AND ROCKS: DIVERSIFICATION PATTERNS IN CHIROPTERA. <i>Evolution; International Journal of Organic Evolution</i> , 2005, 59, 2243.	2.3	5
72	Trees Versus Characters and the Supertree/Supermatrix "Paradox". <i>Systematic Biology</i> , 2004, 53, 356-359.	5.6	39

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73	A species-level phylogenetic supertree of marsupials. <i>Journal of Zoology</i> , 2004, 264, 11-31.	1.7	181
74	The Impact of Species Concept on Biodiversity Studies. <i>Quarterly Review of Biology</i> , 2004, 79, 161-179.	0.1	483
75	Evidence for Multiple Alleles at the DGAT1 Locus Better Explains a Quantitative Trait Locus With Major Effect on Milk Fat Content in Cattle. <i>Genetics</i> , 2004, 167, 1873-1881.	2.9	111
76	The evolution of supertrees. <i>Trends in Ecology and Evolution</i> , 2004, 19, 315-322.	8.7	288
77	New uses for old phylogenies. <i>Computational Biology</i> , 2004, , 3-14.	0.2	7
78	Garbage in, Garbage out. <i>Computational Biology</i> , 2004, , 267-280.	0.2	63
79	Is Sequence Heterochrony an Important Evolutionary Mechanism in Mammals?. <i>Journal of Mammalian Evolution</i> , 2003, 10, 335-361.	1.8	56
80	The adaptive significance of coloration in lagomorphs. <i>Biological Journal of the Linnean Society</i> , 2003, 79, 309-328.	1.6	122
81	MORPHOLOGICAL VARIABILITY AND EVOLUTION OF THE BACULUM (OS PENIS) IN MUSTELIDAE (CARNIVORA). <i>Journal of Mammalogy</i> , 2003, 84, 673-690.	1.3	44
82	Genomic organization of the DGAT2/MOGAT gene family in cattle (<i>Bos taurus</i>) and other mammals. <i>Cytogenetic and Genome Research</i> , 2003, 102, 42-47.	1.1	26
83	Supertrees Are a Necessary Not-So-Evil: A Comment on Gatesy et al.. <i>Systematic Biology</i> , 2003, 52, 724-729.	5.6	34
84	Inverting the hourglass: quantitative evidence against the phylotypic stage in vertebrate development. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2003, 270, 341-346.	2.6	85
85	Novel Versus Unsupported Clades: Assessing the Qualitative Support for Clades in MRP Supertrees. <i>Systematic Biology</i> , 2003, 52, 839-848.	5.6	46
86	Novel Versus Unsupported Clades: Assessing the Qualitative Support for Clades in MRP Supertrees. <i>Systematic Biology</i> , 2003, 52, 839-848.	5.6	6
87	Novel versus unsupported clades: assessing the qualitative support for clades in MRP supertrees. <i>Systematic Biology</i> , 2003, 52, 839-48.	5.6	45
88	The (Super)Tree of Life: Procedures, Problems, and Prospects. <i>Annual Review of Ecology, Evolution, and Systematics</i> , 2002, 33, 265-289.	6.7	222
89	Analyzing Developmental Sequences Within a Phylogenetic Framework. <i>Systematic Biology</i> , 2002, 51, 478-491.	5.6	91
90	Flight of the Dodo. <i>Science</i> , 2002, 295, 1683-1683.	12.6	143

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91	Sealing the fate of a historical taxonomy. <i>Trends in Ecology and Evolution</i> , 2002, 17, 109.	8.7	0
92	From Haeckel to event-pairing: the evolution of developmental sequences. <i>Theory in Biosciences</i> , 2002, 121, 297-320.	1.4	51
93	A phylogenetic supertree of the bats (Mammalia: Chiroptera). <i>Biological Reviews</i> , 2002, 77, 223-259.	10.4	322
94	Analyzing evolutionary patterns in amniote embryonic development*. <i>Evolution & Development</i> , 2002, 4, 292-302.	2.0	79
95	From Haeckel to event-pairing: the evolution of developmental sequences. <i>Theory in Biosciences</i> , 2002, 121, 297-320.	1.4	12
96	Flippers versus feet: comparative trends in aquatic and non-aquatic carnivores. <i>Journal of Animal Ecology</i> , 2001, 70, 386-400.	2.8	20
97	Comparative methods in developmental biology. <i>Zoology</i> , 2001, 104, 278-283.	1.2	35
98	The utility of chemical signals as phylogenetic characters: an example from the Felidae. <i>Biological Journal of the Linnean Society</i> , 2001, 72, 1-15.	1.6	33
99	Assessment of the Accuracy of Matrix Representation with Parsimony Analysis Supertree Construction. <i>Systematic Biology</i> , 2001, 50, 565-579.	5.6	83
100	Assessment of the Accuracy of Matrix Representation with Parsimony Analysis Supertree Construction. <i>Systematic Biology</i> , 2001, 50, 565-579.	5.6	122
101	The utility of chemical signals as phylogenetic characters: an example from the Felidae. <i>Biological Journal of the Linnean Society</i> , 2001, 72, 1-15.	1.6	1
102	Factors Influencing Phylogenetic Inference: A Case Study Using the Mammalian Carnivores. <i>Molecular Phylogenetics and Evolution</i> , 2000, 16, 113-126.	2.7	23
103	ARE PINNIPEDS FUNCTIONALLY DIFFERENT FROM FISSIPED CARNIVORES? THE IMPORTANCE OF PHYLOGENETIC COMPARATIVE ANALYSES. <i>Evolution; International Journal of Organic Evolution</i> , 2000, 54, 1011-1023.	2.3	63
104	PINNIPED BRAIN SIZES. <i>Marine Mammal Science</i> , 2000, 16, 469-481.	1.8	19
105	ARE PINNIPEDS FUNCTIONALLY DIFFERENT FROM FISSIPED CARNIVORES? THE IMPORTANCE OF PHYLOGENETIC COMPARATIVE ANALYSES. <i>Evolution; International Journal of Organic Evolution</i> , 2000, 54, 1011.	2.3	5
106	Considering evolutionary processes in conservation biology. <i>Trends in Ecology and Evolution</i> , 2000, 15, 290-295.	8.7	1,567
107	The calculus of biodiversity: integrating phylogeny and conservation. <i>Trends in Ecology and Evolution</i> , 2000, 15, 92-94.	8.7	24
108	Building large trees by combining phylogenetic information: a complete phylogeny of the extant Carnivora (Mammalia). <i>Biological Reviews</i> , 1999, 74, 143-175.	10.4	552

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109	Building large trees by combining phylogenetic information: a complete phylogeny of the extant Carnivora (Mammalia). <i>Biological Reviews</i> , 1999, 74, 143-175.	10.4	56
110	Supraspecific taxa as terminals in cladistic analysis: implicit assumptions of monophyly and a comparison of methods. <i>Biological Journal of the Linnean Society</i> , 1998, 64, 101-133.	1.6	84
111	Supraspecific taxa as terminals in cladistic analysis: implicit assumptions of monophyly and a comparison of methods. <i>Biological Journal of the Linnean Society</i> , 1998, 64, 101-133.	1.6	15
112	Flight style in bats as predicted from wing morphometry: the effects of specimen preservation. <i>Journal of Zoology</i> , 1994, 234, 275-287.	1.7	16
113	Effects of preservation on wing morphometry of the little brown bat (<i>Myotis lucifugus</i>). <i>Journal of Zoology</i> , 1993, 230, 141-158.	1.7	2
114	Rocking clocks and clocking rocks: a critical look at divergence time estimation in mammals. , 0, , 38-82.		5