Wolfgang WÃ¹/₄ster

List of Publications by Year in descending order

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97 papers 7,449 citations

45 h-index 56724 83 g-index

104 all docs

104 docs citations

104 times ranked 5442 citing authors

#	Article	IF	CITATIONS
1	Mohave Rattlesnake (<i>Crotalus scutulatus</i>) Identification Revisited. Wilderness and Environmental Medicine, 2022, 33, 210-218.	0.9	3
2	How do King Cobras move across a major highway? Unintentional wildlife crossing structures may facilitate movement. Ecology and Evolution, 2022, 12, e8691.	1.9	3
3	Interpopulational variation and ontogenetic shift in the venom composition of Lataste's viper (Vipera) Tj ETQq1	1 0.784314 2.4	4 rgBT /Overle
4	Convergent evolution of toxin resistance in animals. Biological Reviews, 2022, 97, 1823-1843.	10.4	20
5	Bacterial Adaptation to Venom in Snakes and Arachnida. Microbiology Spectrum, 2022, 10, .	3.0	13
6	Crowdsourcing snake identification with online communities of professional herpetologists and avocational snake enthusiasts. Royal Society Open Science, 2021, 8, 201273.	2.4	11
7	Convergent evolution of pain-inducing defensive venom components in spitting cobras. Science, 2021, 371, 386-390.	12.6	96
8	Genome-wide data implicate terminal fusion automixis in king cobra facultative parthenogenesis. Scientific Reports, 2021, 11, 7271.	3.3	10
9	Unexpected lack of specialisation in the flow properties of spitting cobra venom. Journal of Experimental Biology, 2021, 224, .	1.7	2
10	Confronting taxonomic vandalism in biology: conscientious community self-organization can preserve nomenclatural stability. Biological Journal of the Linnean Society, 2021, 133, 645-670.	1.6	16
11	An evaluation of the nomina for death adders (Acanthophis Daudin, 1803) proposed by Wells & Samp; Wellington (1985), and confirmation of A. cryptamydros Maddock et al., 2015 as the valid name for the Kimberley death adder. Zootaxa, 2021, 4995, 161-172.	0.5	0
12	Citizen science and online data: Opportunities and challenges for snake ecology and action against snakebite. Toxicon: X, 2021, 9-10, 100071.	2.9	10
13	Promoting co-existence between humans and venomous snakes through increasing the herpetological knowledge base. Toxicon: X, 2021, 12, 100081.	2.9	21
14	King or royal family? Testing for species boundaries in the King Cobra, Ophiophagus hannah (Cantor,) Tj ETQq0 0	0 0 rgBT /O 2.7	verlock 10 Tf 9
15	Evaluating taxonomic inflation: towards evidence-based species delimitation in Eurasian vipers (Serpentes: Viperinae). Amphibia - Reptilia, 2020, 41, 285-311.	0.5	45
16	Fangs for the Memories? A Survey of Pain in Snakebite Patients Does Not Support a Strong Role for Defense in the Evolution of Snake Venom Composition. Toxins, 2020, 12, 201.	3.4	20
17	Deconstructing compassionate conservation. Conservation Biology, 2019, 33, 760-768.	4.7	53
18	Ancient habitat shifts and organismal diversification are decoupled in the African viper genus <i>Bitis</i> (Serpentes: Viperidae). Journal of Biogeography, 2019, 46, 1234-1248.	3.0	26

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19	When one phenotype is not enough: divergent evolutionary trajectories govern venom variation in a widespread rattlesnake species. Proceedings of the Royal Society B: Biological Sciences, 2019, 286, 20182735.	2.6	64
20	The medical threat of mamba envenoming in sub-Saharan Africa revealed by genus-wide analysis of venom composition, toxicity and antivenomics profiling of available antivenoms. Journal of Proteomics, 2018, 172, 173-189.	2.4	80
21	Integration of nuclear and mitochondrial gene sequences and morphology reveals unexpected diversity in the forest cobra (Naja melanoleuca) species complex in Central and West Africa (Serpentes: Elapidae). Zootaxa, 2018, 4455, 68-98.	0.5	33
22	Venom Complexity in a Pitviper Produced by Facultative Parthenogenesis. Scientific Reports, 2018, 8, 11539.	3.3	14
23	Widespread vulnerability of Malagasy predators to the toxins of an introduced toad. Current Biology, 2018, 28, R654-R655.	3.9	22
24	Taxonomy based on science is necessary for global conservation. PLoS Biology, 2018, 16, e2005075.	5.6	149
25	Differential procoagulant effects of saw-scaled viper (Serpentes: Viperidae: Echis) snake venoms on human plasma and the narrow taxonomic ranges of antivenom efficacies. Toxicology Letters, 2017, 280, 159-170.	0.8	69
26	Multi-locus phylogeny and species delimitation of Australo-Papuan blacksnakes (Pseudechis Wagler,) Tj ETQq0 C	0 rgBT /0	Oveglock 10 Tf
27	Venom On-a-Chip: A Fast and Efficient Method for Comparative Venomics. Toxins, 2017, 9, 179.	3.4	17
28	Phylogeography of the Central American lancehead Bothrops asper (SERPENTES: VIPERIDAE). PLoS ONE, 2017, 12, e0187969.	2.5	36
29	Is Hybridization a Source of Adaptive Venom Variation in Rattlesnakes? A Test, Using a Crotalus scutulatus × viridis Hybrid Zone in Southwestern New Mexico. Toxins, 2016, 8, 188.	3.4	29
30	Phylogeny and diversification of mountain vipers (Montivipera, Nilson et al., 2001) triggered by multiple Plioâ€"Pleistocene refugia and high-mountain topography in the Near and Middle East. Molecular Phylogenetics and Evolution, 2016, 101, 336-351.	2.7	43
31	A new species of death adder (Acanthophis: Serpentes: Elapidae)Âfrom north-western Australia. Zootaxa, 2015, 4007, 301-26.	0.5	9
32	Widespread convergence in toxin resistance by predictable molecular evolution. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 11911-11916.	7.1	130
33	Medically important differences in snake venom composition are dictated by distinct postgenomic mechanisms. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 9205-9210.	7.1	253
34	The king cobra genome reveals dynamic gene evolution and adaptation in the snake venom system. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 20651-20656.	7.1	412
35	Complex cocktails: the evolutionary novelty of venoms. Trends in Ecology and Evolution, 2013, 28, 219-229.	8.7	785
36	Phylogeography of the widespread <scp>A</scp> frican puff adder (<i><scp>B</scp>itis arietans</i>) reveals multiple <scp>P</scp> leistocene refugia in southern <scp>A</scp> frica. Molecular Ecology, 2013, 22, 1134-1157.	3.9	71

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37	Anonymous nuclear markers for the African adders (Serpentes: Viperidae: Bitis). Conservation Genetics Resources, 2012, 4, 967-969.	0.8	5
38	Dynamic evolution of venom proteins in squamate reptiles. Nature Communications, 2012, 3, 1066.	12.8	86
39	Venom lethality and diet: Differential responses of natural prey and model organisms to the venom of the saw-scaled vipers (Echis). Toxicon, 2012, 59, 110-116.	1.6	52
40	The structural and functional diversification of the Toxicofera reptile venom system. Toxicon, 2012, 60, 434-448.	1.6	142
41	Domain Loss Facilitates Accelerated Evolution and Neofunctionalization of Duplicate Snake Venom Metalloproteinase Toxin Genes. Molecular Biology and Evolution, 2011, 28, 2637-2649.	8.9	147
42	On the generic classification of the rattlesnakes, with special reference to the Neotropical Crotalus durissus complex (Squamata: Viperidae). Zoologia, 2011, 28, 417-419.	0.5	6
43	Ending the drought: New strategies for improving the flow of affordable, effective antivenoms in Asia and Africa. Journal of Proteomics, 2011, 74, 1735-1767.	2.4	206
44	Gene Tree Parsimony of Multilocus Snake Venom Protein Families Reveals Species Tree Conflict as a Result of Multiple Parallel Gene Loss. Molecular Biology and Evolution, 2011, 28, 1157-1172.	8.9	24
45	What's your poison?. Heredity, 2010, 104, 519-519.	2.6	3
46	Pre-Clinical Assays Predict Pan-African Echis Viper Efficacy for a Species-Specific Antivenom. PLoS Neglected Tropical Diseases, 2010, 4, e851.	3.0	89
47	Destruction of the collection of reptiles and arthropods at Butantan Institute: a view from the United Kingdom. Journal of Venomous Animals and Toxins Including Tropical Diseases, 2010, 16, .	1.4	9
48	Phylogeography and systematic revision of the Egyptian cobra (Serpentes: Elapidae: Naja haje) species complex, with the description of a new species from West Africa. Zootaxa, 2009, 2236, 1-25.	0.5	22
49	In praise of subgenera: taxonomic status of cobras of the genus Naja Laurenti (Serpentes: Elapidae). Zootaxa, 2009, 2236, 26-36.	0.5	50
50	Coevolution of diet and prey-specific venom activity supports the role of selection in snake venom evolution. Proceedings of the Royal Society B: Biological Sciences, 2009, 276, 2443-2449.	2.6	273
51	Comparative venom gland transcriptome surveys of the saw-scaled vipers (Viperidae: Echis) reveal substantial intra-family gene diversity and novel venom transcripts. BMC Genomics, 2009, 10, 564.	2.8	135
52	When continents collide: Phylogeny, historical biogeography and systematics of the medically important viper genus Echis (Squamata: Serpentes: Viperidae). Molecular Phylogenetics and Evolution, 2009, 53, 792-807.	2.7	125
53	Recent Advances in Venomous Snake Systematics. , 2009, , 25-64.		1
54	A nesting of vipers: Phylogeny and historical biogeography of the Viperidae (Squamata: Serpentes). Molecular Phylogenetics and Evolution, 2008, 49, 445-459.	2.7	218

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55	Origin of the eastern brownsnake, Pseudonaja textilis (Dumeril, Bibron and Dumeril) (Serpentes:) Tj ETQq1 1 0.786 on the status of Pseudonaja textilis pughi Hoser 2003. Zootaxa, 2008, 1703, 47.	4314 rgBT 0.5	/Overlock 15
56	Get an eyeful of this: a new species of giant spitting cobra from eastern and north-eastern Africa (Squamata: Serpentes: Elapidae: Naja). Zootaxa, 2007, 1532, 51-68.	0.5	25
57	The phylogeny of cobras inferred from mitochondrial DNA sequences: Evolution of venom spitting and the phylogeography of the African spitting cobras (Serpentes: Elapidae: Naja nigricollis complex). Molecular Phylogenetics and Evolution, 2007, 45, 437-453.	2.7	95
58	Phylogeographic patterns of trans-Amazonian vicariants and Amazonian biogeography: the Neotropical rattlesnake (Crotalus durissus complex) as an example. Journal of Biogeography, 2007, 34, 1296-1312.	3.0	61
59	The good, the bad and the ugly: Australian snake taxonomists and a history of the taxonomy of Australia's venomous snakes. Toxicon, 2006, 48, 919-930.	1.6	25
60	Roles of CITES in Protecting New Species. Science, 2006, 313, 915c-916c.	12.6	3
61	Snakes across the Strait: trans-Torresian phylogeographic relationships in three genera of Australasian snakes (Serpentes: Elapidae: Acanthophis, Oxyuranus, and Pseudechis). Molecular Phylogenetics and Evolution, 2005, 34, 1-14.	2.7	78
62	Tracing an invasion: landbridges, refugia, and the phylogeography of the Neotropical rattlesnake (Serpentes: Viperidae: Crotalus durissus). Molecular Ecology, 2005, 14, 1095-1108.	3.9	174
63	No rattlesnakes in the rainforests: reply to Gosling and Bush. Molecular Ecology, 2005, 14, 3619-3621.	3.9	5
64	Morphological correlates of incipient arboreality and ornithophagy in island pitvipers, and the phylogenetic position of Bothrops insularis. Journal of Zoology, 2005, 266, 1-10.	1.7	46
65	Do aposematism and Batesian mimicry require bright colours? A test, using European viper markings. Proceedings of the Royal Society B: Biological Sciences, 2004, 271, 2495-2499.	2.6	130
66	Assembling an Arsenal: Origin and Evolution of the Snake Venom Proteome Inferred from Phylogenetic Analysis of Toxin Sequences. Molecular Biology and Evolution, 2004, 21, 870-883.	8.9	206
67	A review of the southern African â€~nonâ€spitting' cobras (Serpentes: Elapidae: <i>Naja</i>). African Journal of Herpetology, 2004, 53, 101-122.	0.9	24
68	Molecular Evolution and Phylogeny of Elapid Snake Venom Three-Finger Toxins. Journal of Molecular Evolution, 2003, 57, 110-129.	1.8	319
69	Isolation of a Neurotoxin (?-colubritoxin) from a Nonvenomous Colubrid: Evidence for Early Origin of Venom in Snakes. Journal of Molecular Evolution, 2003, 57, 446-452.	1.8	138
70	A new species of spitting cobra (Naja) from north-eastern Africa (Serpentes: Elapidae). Journal of Zoology, 2003, 259, 345-359.	1.7	27
71	Analysis of Colubroidea snake venoms by liquid chromatography with mass spectrometry: evolutionary and toxinological implications. Rapid Communications in Mass Spectrometry, 2003, 17, 2047-2062.	1.5	141
72	Effectiveness of Snake Antivenom: Species and Regional Venom Variation and Its Clinical Impact. Toxin Reviews, 2003, 22, 23-34.	1.5	64

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73	The conserved structure of snake venom toxins confers extensive immunological cross-reactivity to toxin-specific antibody. Toxicon, 2003, 41, 441-449.	1.6	52
74	A NEW SPECIES OF WOLF SNAKE (SERPENTES: COLUBRIDAE: LYCODON) FROM THE CARDAMOM MOUNTAINS, SOUTHWESTERN CAMBODIA. Herpetologica, 2002, 58, 498-504.	0.4	34
75	Electrospray liquid chromatography/mass spectrometry fingerprinting ofAcanthophis(death adder) venoms: taxonomic and toxinological implications. Rapid Communications in Mass Spectrometry, 2002, 16, 600-608.	1.5	70
76	Combining mitochondrial DNA sequences and morphological data to infer species boundaries: phylogeography of lanceheaded pitvipers in the Brazilian Atlantic forest, and the status of Bothrops pradoi (Squamata: Serpentes: Viperidae). Journal of Evolutionary Biology, 2001, 14, 527-538.	1.7	86
77	Historical Biogeography of the Western Rattlesnake (Serpentes: Viperidae: Crotalus viridis), Inferred from Mitochondrial DNA Sequence Information. Molecular Phylogenetics and Evolution, 2000, 15, 269-282.	2.7	106
78	Synopsis of recent developments in venomous snake systematics, No. 3. Toxicon, 1999, 37, 1123-1129.	1.6	20
79	Evidence that humidity influences snake activity patterns: a field study of the Malayan pit viper Calloselasma rhodostoma. Ecography, 1998, 21, 25-34.	4.5	63
80	Synopsis of recent developments in venomous snake systematics, No. 2. Toxicon, 1998, 36, 299-307.	1.6	30
81	Congruence between morphological variation and altitudinal gradient across a hybrid zone between carrion and hooded crows. Italian Journal of Zoology, 1998, 65, 407-412.	0.6	4
82	Intraspecific Variation in the Feeding Ecology of the Crotaline Snake Calloselasma rhodostoma in Southeast Asia. Journal of Herpetology, 1998, 32, 198.	0.5	42
83	Synopsis of recent developments in venomous snake systematics. Toxicon, 1997, 35, 319-340.	1.6	56
84	Redescription of <i>Naja siamensis</i> (Serpentes: Elapidae), a widely overlooked spitting cobra from S.E. Asia: geographic variation, medical importance and designation of a neotype. Journal of Zoology, 1997, 243, 771-788.	1.7	17
85	Electrophoretic profiles and biological activities: Intraspecific variation in the venom of the malayan pit viper (Calloselasma rhodostoma). Toxicon, 1996, 34, 67-79.	1.6	80
86	Taxonomic changes and toxinology: Systematic revisions of the asiatic cobras (Naja naja species) Tj ETQq0 0 0 n	gBT_lOverl	ock 10 Tf 50
87	Reviews of venomous snake systematics in Toxicon. Toxicon, 1996, 34, 397-398.	1.6	11
88	Venomous snake systematics: implications for snakebite treatment and toxinology. Toxicon, 1996, 34, 143.	1.6	9
89	Diet and snake venom evolution. Nature, 1996, 379, 537-540.	27.8	593
90	Population systematics of the snake genus Naja (Reptilia: Serpentes: Elapidae) in Indochina: Multivariate morphometrics and comparative mitochondrial DNA sequencing (cytochrome oxidase I). Journal of Evolutionary Biology, 1995, 8, 493-510.	1.7	33

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91	Treatment of snake bites by Bothrops species and Lachesis muta in Ecuador: laboratory screening of candidate antivenoms. Transactions of the Royal Society of Tropical Medicine and Hygiene, 1995, 89, 550-554.	1.8	48
92	Naja siamensis, a cryptic species of venomous snake revealed by mtDNA sequencing. Experientia, 1994, 50, 75-79.	1.2	19
93	Defensive Hemipenis Display in the Kukri Snake Oligodon cyclurus. Journal of Herpetology, 1992, 26, 238.	0.5	3
94	Population systematics of Russell's viper: a multivariate study. Biological Journal of the Linnean Society, 1992, 47, 97-113.	1.6	52
95	Geographic variation and population systematics: Distinguishing between ecogenetics and phylogenetics. Bollettino Di Zoologia, 1991, 58, 329-335.	0.3	42
96	Asiatic cobras: Systematics and snakebite. Experientia, 1991, 47, 205-209.	1.2	47
97	Population affinities of the asiatic cobra (Naja naja) species complex in south-east Asia: reliability and random resampling. Biological Journal of the Linnean Society, 1989, 36, 391-409.	1.6	37