

# Klaus Peter Schliep

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8589402/publications.pdf>

Version: 2024-02-01

21  
papers

8,812  
citations

566801

15  
h-index

713013

21  
g-index

24  
all docs

24  
docs citations

24  
times ranked

16853  
citing authors

#	ARTICLE	IF	CITATIONS
1	ape 5.0: an environment for modern phylogenetics and evolutionary analyses in R. <i>Bioinformatics</i> , 2019, 35, 526-528.	1.8	5,128
2	phangorn: phylogenetic analysis in R. <i>Bioinformatics</i> , 2011, 27, 592-593.	1.8	2,840
3	Intertwining phylogenetic trees and networks. <i>Methods in Ecology and Evolution</i> , 2017, 8, 1212-1220.	2.2	211
4	Putative Origins of Cell-Free DNA in Humans: A Review of Active and Passive Nucleic Acid Release Mechanisms. <i>International Journal of Molecular Sciences</i> , 2020, 21, 8062.	1.8	103
5	Local Mobile Gene Pools Rapidly Cross Species Boundaries To Create Endemicity within Global <i>Vibrio cholerae</i> Populations. <i>MBio</i> , 2011, 2, .	1.8	97
6	phyloregion: R package for biogeographical regionalization and macroecology. <i>Methods in Ecology and Evolution</i> , 2020, 11, 1483-1491.	2.2	70
7	Host-associated microbiomes are predicted by immune system complexity and climate. <i>Genome Biology</i> , 2020, 21, 23.	3.8	54
8	Let Them Fall Where They May: Congruence Analysis in Massive Phylogenetically Messy Data Sets. <i>Molecular Biology and Evolution</i> , 2011, 28, 2773-2785.	3.5	43
9	rWind: download, edit and include wind data in ecological and evolutionary analysis. <i>Ecography</i> , 2019, 42, 804-810.	2.1	39
10	Of woods and webs: possible alternatives to the tree of life for studying genomic fluidity in <i>E. coli</i> . <i>Biology Direct</i> , 2011, 6, 39; discussion 39.	1.9	37
11	Harvesting Evolutionary Signals in a Forest of Prokaryotic Gene Trees. <i>Molecular Biology and Evolution</i> , 2011, 28, 1393-1405.	3.5	33
12	Life history linked to immune investment in developing amphibians. , 2016, 4, cow025.		28
13	Phylogenetic signal and evolutionary correlates of urban tolerance in a widespread neotropical lizard clade*. <i>Evolution; International Journal of Organic Evolution</i> , 2020, 74, 1274-1288.	1.1	24
14	apex : phylogenetics with multiple genes. <i>Molecular Ecology Resources</i> , 2017, 17, 19-26.	2.2	23
15	The evolution of social behaviour in Blaberid cockroaches with diverse habitats and social systems: phylogenetic analysis of behavioural sequences. <i>Biological Journal of the Linnean Society</i> , 2014, 111, 58-77.	0.7	20
16	Linking genomics and population genetics with R. <i>Molecular Ecology Resources</i> , 2017, 17, 54-66.	2.2	17
17	Graphs in phylogenetic comparative analysis: Anscombe's quartet revisited. <i>Methods in Ecology and Evolution</i> , 2018, 9, 2145-2154.	2.2	9
18	Using Genomics to Shape the Definition of the Agglutinin-Like Sequence (ALS) Family in the Saccharomycetales. <i>Frontiers in Cellular and Infection Microbiology</i> , 2021, 11, 794529.	1.8	9

#	ARTICLE	IF	CITATIONS
19	Comparison of the Serum Tumor Markers S100 and Melanoma-inhibitory Activity (MIA) in the Monitoring of Patients with Metastatic Melanoma Receiving Vaccination Immunotherapy with Dendritic Cells. <i>Anticancer Research</i> , 2017, 37, 5033-5037.	0.5	6
20	Linking morphological and molecular sources to disentangle the case of <i>Xylodon australis</i> . <i>Scientific Reports</i> , 2020, 10, 22004.	1.6	4
21	Brain Allometry Across Macroevolutionary Scales in Squamates Suggests a Conserved Pattern in Snakes. <i>Zoology</i> , 2021, 146, 125926.	0.6	3