

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

567281

15
h-index

713466

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. | 4.1 | 5,128 |
| 2 | phangorn: phylogenetic analysis in R. <i>Bioinformatics</i> , 2011, 27, 592-593. | 4.1 | 2,840 |
| 3 | Intertwining phylogenetic trees and networks. <i>Methods in Ecology and Evolution</i> , 2017, 8, 1212-1220. | 5.2 | 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. | 4.1 | 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, . | 4.1 | 97 |
| 6 | phyloregion: R package for biogeographical regionalization and macroecology. <i>Methods in Ecology and Evolution</i> , 2020, 11, 1483-1491. | 5.2 | 70 |
| 7 | Host-associated microbiomes are predicted by immune system complexity and climate. <i>Genome Biology</i> , 2020, 21, 23. | 8.8 | 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. | 8.9 | 43 |
| 9 | rWind: download, edit and include wind data in ecological and evolutionary analysis. <i>Ecography</i> , 2019, 42, 804-810. | 4.5 | 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. | 4.6 | 37 |
| 11 | Harvesting Evolutionary Signals in a Forest of Prokaryotic Gene Trees. <i>Molecular Biology and Evolution</i> , 2011, 28, 1393-1405. | 8.9 | 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. | 2.3 | 24 |
| 14 | apex : phylogenetics with multiple genes. <i>Molecular Ecology Resources</i> , 2017, 17, 19-26. | 4.8 | 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. | 1.6 | 20 |
| 16 | Linking genomics and population genetics with R. <i>Molecular Ecology Resources</i> , 2017, 17, 54-66. | 4.8 | 17 |
| 17 | Graphs in phylogenetic comparative analysis: Anscombe's quartet revisited. <i>Methods in Ecology and Evolution</i> , 2018, 9, 2145-2154. | 5.2 | 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. | 3.9 | 9 |

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|----|--|-----|-----------|
| 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. | 1.1 | 6 |
| 20 | Linking morphological and molecular sources to disentangle the case of <i>Xylodon australis</i> . <i>Scientific Reports</i> , 2020, 10, 22004. | 3.3 | 4 |
| 21 | Brain Allometry Across Macroevolutionary Scales in Squamates Suggests a Conserved Pattern in Snakes. <i>Zoology</i> , 2021, 146, 125926. | 1.2 | 3 |