

Dirk KrÃ¼ger

List of Publications by Year in descending order

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Version: 2024-02-01

40
papers

2,590
citations

236912

25
h-index

289230

40
g-index

41
all docs

41
docs citations

41
times ranked

3574
citing authors

#	ARTICLE	IF	CITATIONS
1	Animal Evolution and the Molecular Signature of Radiations Compressed in Time. <i>Science</i> , 2005, 310, 1933-1938.	12.6	357
2	Life in leaf litter: novel insights into community dynamics of bacteria and fungi during litter decomposition. <i>Molecular Ecology</i> , 2016, 25, 4059-4074.	3.9	297
3	Preserving Accuracy in GenBank. <i>Science</i> , 2008, 319, 1616-1616.	12.6	198
4	Wood decay rates of 13 temperate tree species in relation to wood properties, enzyme activities and organismic diversities. <i>Forest Ecology and Management</i> , 2017, 391, 86-95.	3.2	151
5	Linking molecular deadwood-inhabiting fungal diversity and community dynamics to ecosystem functions and processes in Central European forests. <i>Fungal Diversity</i> , 2016, 77, 367-379.	12.3	140
6	Network Analysis Reveals Ecological Links between N-Fixing Bacteria and Wood-Decaying Fungi. <i>PLoS ONE</i> , 2014, 9, e88141.	2.5	129
7	A pyrosequencing insight into sprawling bacterial diversity and community dynamics in decaying deadwood logs of <i>Fagus sylvatica</i> and <i>Picea abies</i> . <i>Scientific Reports</i> , 2015, 5, 9456.	3.3	101
8	Widespread Occurrence of Expressed Fungal Secretory Peroxidases in Forest Soils. <i>PLoS ONE</i> , 2014, 9, e95557.	2.5	91
9	Molecular evidence strongly supports deadwood-inhabiting fungi exhibiting unexpected tree species preferences in temperate forests. <i>ISME Journal</i> , 2018, 12, 289-295.	9.8	90
10	Uncoupling of microbial community structure and function in decomposing litter across beech forest ecosystems in Central Europe. <i>Scientific Reports</i> , 2014, 4, 7014.	3.3	65
11	Influence of Different Forest System Management Practices on Leaf Litter Decomposition Rates, Nutrient Dynamics and the Activity of Ligninolytic Enzymes: A Case Study from Central European Forests. <i>PLoS ONE</i> , 2014, 9, e93700.	2.5	65
12	Spatial Distribution of Fungal Communities in an Arable Soil. <i>PLoS ONE</i> , 2016, 11, e0148130.	2.5	63
13	Changes within a single land-use category alter microbial diversity and community structure: Molecular evidence from wood-inhabiting fungi in forest ecosystems. <i>Journal of Environmental Management</i> , 2014, 139, 109-119.	7.8	61
14	Effects of resource availability and quality on the structure of the micro-food web of an arable soil across depth. <i>Soil Biology and Biochemistry</i> , 2012, 50, 1-11.	8.8	60
15	Dynamics of fungal community composition, decomposition and resulting deadwood properties in logs of <i>Fagus sylvatica</i> , <i>Picea abies</i> and <i>Pinus sylvestris</i> . <i>Forest Ecology and Management</i> , 2016, 382, 129-142.	3.2	58
16	Effects of Forest Management Practices in Temperate Beech Forests on Bacterial and Fungal Communities Involved in Leaf Litter Degradation. <i>Microbial Ecology</i> , 2015, 69, 905-913.	2.8	56
17	Are correlations between deadwood fungal community structure, wood physico-chemical properties and lignin-modifying enzymes stable across different geographical regions?. <i>Fungal Ecology</i> , 2016, 22, 98-105.	1.6	47
18	Correlations between the composition of modular fungal communities and litter decomposition-associated ecosystem functions. <i>Fungal Ecology</i> , 2016, 22, 106-114.	1.6	46

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19	Determinants of Deadwood-Inhabiting Fungal Communities in Temperate Forests: Molecular Evidence From a Large Scale Deadwood Decomposition Experiment. <i>Frontiers in Microbiology</i> , 2018, 9, 2120.	3.5	43
20	The Lycoperdales. A molecular approach to the systematics of some gasteroid mushrooms. <i>Mycologia</i> , 2001, 93, 947-957.	1.9	42
21	Basidiomycetous Yeasts from Boletales Fruiting Bodies and Their Interactions with the Mycoparasite <i>Sepedonium chrysospermum</i> and the Host Fungus <i>Paxillus</i> . <i>Microbial Ecology</i> , 2012, 63, 295-303.	2.8	42
22	Secondary structure of ITS2 rRNA provides taxonomic characters for systematic studies – a case in Lycoperdaceae (Basidiomycota). <i>Mycological Research</i> , 2008, 112, 316-330.	2.5	35
23	Resource Type and Availability Regulate Fungal Communities Along Arable Soil Profiles. <i>Microbial Ecology</i> , 2015, 70, 390-399.	2.8	32
24	Chilenopectins A and B, Peptaibols from the Chilean <i>Sepedonium</i> aff. <i>chalcipori</i> KSH 883. <i>Journal of Natural Products</i> , 2016, 79, 929-938.	3.0	32
25	Comparing fungal richness and community composition in coarse woody debris in Central European beech forests under three types of management. <i>Mycological Progress</i> , 2014, 13, 959-964.	1.4	31
26	Actinobacteria may influence white truffle (<i>Tuber magnatum</i> Pico) nutrition, ascocarp degradation and interactions with other soil fungi. <i>Fungal Ecology</i> , 2013, 6, 527-538.	1.6	27
27	Increasing N deposition impacts neither diversity nor functions of deadwood-inhabiting fungal communities, but adaptation and functional redundancy ensure ecosystem function. <i>Environmental Microbiology</i> , 2018, 20, 1693-1710.	3.8	26
28	Patterns of laccase and peroxidases in coarse woody debris of <i>Fagus sylvatica</i> , <i>Picea abies</i> and <i>Pinus sylvestris</i> and their relation to different wood parameters. <i>European Journal of Forest Research</i> , 2016, 135, 109-124.	2.5	24
29	Home-Field Advantage in Wood Decomposition Is Mainly Mediated by Fungal Community Shifts at “Home” Versus “Away”. <i>Microbial Ecology</i> , 2019, 78, 725-736.	2.8	24
30	First insight into dead wood protistan diversity: a molecular sampling of bright-spored Myxomycetes (Amoebozoa, slime-moulds) in decaying beech logs. <i>FEMS Microbiology Ecology</i> , 2015, 91, .	2.7	23
31	The Lycoperdales. A Molecular Approach to the Systematics of Some Gasteroid Mushrooms. <i>Mycologia</i> , 2001, 93, 947.	1.9	21
32	Molecular phylogenies and mating study data in <i>Polyporus</i> with special emphasis on group “ <i>Melanopus</i> ” (Basidiomycota). <i>Mycological Progress</i> , 2006, 5, 185-206.	1.4	18
33	Diversity and Interactions of Wood-Inhabiting Fungi and Beetles after Deadwood Enrichment. <i>PLoS ONE</i> , 2015, 10, e0143566.	2.5	18
34	Influence of Commonly Used Primer Systems on Automated Ribosomal Intergenic Spacer Analysis of Bacterial Communities in Environmental Samples. <i>PLoS ONE</i> , 2015, 10, e0118967.	2.5	18
35	Diversity Measures in Environmental Sequences Are Highly Dependent on Alignment Quality – Data from ITS and New LSU Primers Targeting Basidiomycetes. <i>PLoS ONE</i> , 2012, 7, e32139.	2.5	15
36	Application of next-generation sequencing technologies to conservation of wood-inhabiting fungi. <i>Conservation Biology</i> , 2019, 33, 716-724.	4.7	13

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37	A better understanding of functional roles of fungi in the decomposition process: using precursor rRNA containing ITS regions as a marker for the active fungal community. <i>Annals of Forest Science</i> , 2012, 69, 659-662.	2.0	12
38	The tropical <i>Polyporus tricholoma</i> (Polyporaceae) – Taxonomy, phylogeny, and the development of methods to detect cryptic species. <i>Mycological Progress</i> , 2004, 3, 65-79.	1.4	9
39	New measures of topological stability in phylogenetic trees – Taking taxon composition into account. <i>Bioinformatics</i> , 2006, 1, 327-330.	0.5	9
40	Assessing the Mycorrhizal Diversity of Soils and Identification of Fungus Fruiting Bodies and Axenic Cultures. <i>Soil Biology</i> , 2009, , 159-188.	0.8	1