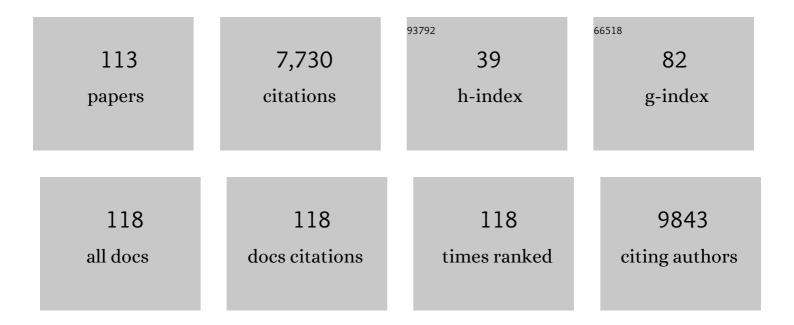
## Sandrine Pavoine

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8506251/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Traitâ€habitat associations explain novel bird assemblages mixing native and alien species across New Zealand landscapes. Diversity and Distributions, 2022, 28, 38-52.	1.9	6
2	A new parametric measure of functional dissimilarity: Bridging the gap between the Bray-Curtis dissimilarity and the Euclidean distance. Ecological Modelling, 2022, 466, 109880.	1.2	8
3	Effects of life-history traits and network topological characteristics on the robustness of marine food webs. Global Ecology and Conservation, 2022, 34, e02048.	1.0	2
4	Trade-offs in the conservation of phylogenetically distinctive species. Biological Conservation, 2022, 270, 109565.	1.9	3
5	Functional imbalance not functional evenness is the third component of community structure. Ecological Indicators, 2022, 140, 109035.	2.6	6
6	A new method for indicator species analysis in the framework of multivariate analysis of variance. Journal of Vegetation Science, 2021, 32, e13013.	1.1	3
7	Beta redundancy for functional ecology. Methods in Ecology and Evolution, 2021, 12, 1062-1069.	2.2	5
8	Revisiting species and areas of interest for conserving global mammalian phylogenetic diversity. Nature Communications, 2021, 12, 3694.	5.8	25
9	Biomass of slow life history species increases as local bottom trawl effort decreases in the Celtic sea. Journal of Environmental Management, 2021, 290, 112634.	3.8	0
10	A framework for understanding how biodiversity patterns unfold across multiple spatial scales in urban ecosystems. Ecosphere, 2021, 12, e03650.	1.0	24
11	Disturbed habitats locally reduce the signal of deep evolutionary history in functional traits of plants. New Phytologist, 2021, 232, 1849-1862.	3.5	7
12	Towards a unifying framework for diversity and dissimilarity coefficients. Ecological Indicators, 2021, 129, 107971.	2.6	10
13	On the relationships between rarity, uniqueness, distinctiveness, originality and functional/phylogenetic diversity. Biological Conservation, 2021, 263, 109356.	1.9	8
14	On two dissimilarity-based measures of functional beta diversity. Ecological Informatics, 2021, 66, 101458.	2.3	1
15	A global database for metacommunity ecology, integrating species, traits, environment and space. Scientific Data, 2020, 7, 6.	2.4	28
16	Environment outweighs the effects of fishing in regulating demersal community structure in an exploited marine ecosystem. Global Change Biology, 2020, 26, 2106-2119.	4.2	27
17	Assembly rules of helminth parasite communities in grey mullets: combining components of diversity. International Journal for Parasitology, 2020, 50, 1089-1098.	1.3	13
18	adiv: An <scp>r</scp> package to analyse biodiversity in ecology. Methods in Ecology and Evolution, 2020, 11, 1106-1112.	2.2	63

1

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19	The worldwide impact of urbanisation on avian functional diversity. Ecology Letters, 2020, 23, 962-972.	3.0	95
20	Comparing taxon- and trait-environment relationships in stream communities. Ecological Indicators, 2020, 117, 106625.	2.6	7
21	From alpha to beta functional and phylogenetic redundancy. Methods in Ecology and Evolution, 2020, 11, 487-493.	2.2	19
22	Urbanization Effects on Biodiversity Revealed by a Two-Scale Analysis of Species Functional Uniqueness vs. Redundancy. Frontiers in Ecology and Evolution, 2020, 8, .	1.1	44
23	Rarefaction of beta diversity. Ecological Indicators, 2019, 107, 105606.	2.6	6
24	Explosive breeding in tropical anurans: environmental triggers, community composition and acoustic structure. BMC Ecology, 2019, 19, 28.	3.0	23
25	A simple translation from indices of species diversity to indices of phylogenetic diversity. Ecological Indicators, 2019, 101, 552-561.	2.6	22
26	Reconciling the concepts and measures of diversity, rarity and originality in ecology and evolution. Biological Reviews, 2019, 94, 1317-1337.	4.7	67
27	Species splitting increases estimates of evolutionary history at risk. Biological Conservation, 2019, 235, 27-35.	1.9	19
28	Mammal extinctions and the increasing isolation of humans on the tree of life. Ecology and Evolution, 2019, 9, 914-924.	0.8	12
29	An ordination approach to explore similarities among communities. Journal of Theoretical Biology, 2019, 462, 85-96.	0.8	3
30	Measuring functional dissimilarity among plots: Adapting old methods to new questions. Ecological Indicators, 2019, 97, 67-72.	2.6	15
31	Ecological versatility and the assembly of multiple competitors: cautionary notes for assembly inferences. Ecology, 2018, 99, 1173-1183.	1.5	7
32	Trait and phylogenetic diversity provide insights into community assembly of reefâ€associated shrimps (Palaemonidae) at different spatial scales across the Chagos Archipelago. Ecology and Evolution, 2018, 8, 4098-4107.	0.8	7
33	Predicting the impacts of co-extinctions on phylogenetic diversity in mutualistic networks. Biological Conservation, 2018, 219, 161-171.	1.9	8
34	A new method for quantifying the phylogenetic redundancy of biological communities. Oecologia, 2018, 186, 339-346.	0.9	10
35	A Generalized Framework for Analyzing Taxonomic, Phylogenetic, and Functional Community Structure Based on Presence–Absence Data. Mathematics, 2018, 6, 250.	1.1	9

Relating Species Traits to Environment. , 2018, , 223-237.

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37	Analysing Spatial Structures. , 2018, , 239-260.		1
38	Analysing Patterns of Biodiversity. , 2018, , 281-294.		0
39	Analysing Changes in Structures. , 2018, , 167-204.		0
40	Analysing Phylogenetic Structures. , 2018, , 261-280.		0
41	Multivariate Analysis of Ecological Data with ade4. , 2018, , .		206
42	Indicators for the Expected Loss of Phylogenetic Diversity. , 2018, , 73-91.		14
43	Priority Areas for Phylogenetic Diversity: Maximising Gains in the Mediterranean Basin. , 2018, , 145-166.		4
44	Predicting loss of evolutionary history: Where are we?. Biological Reviews, 2017, 92, 271-291.	4.7	67
45	From inselberg to inselberg: Floristic patterns across scales in French Guiana (South America). Flora: Morphology, Distribution, Functional Ecology of Plants, 2017, 229, 147-158.	0.6	18
46	Urbanisation and the loss of phylogenetic diversity in birds. Ecology Letters, 2017, 20, 721-729.	3.0	145
47	From phylogenetic to functional originality: Guide through indices and new developments. Ecological Indicators, 2017, 82, 196-205.	2.6	47
48	A guide to phylogenetic metrics for conservation, community ecology and macroecology. Biological Reviews, 2017, 92, 698-715.	4.7	570
49	Ancestrality and evolution of trait syndromes in finches (Fringillidae). Ecology and Evolution, 2017, 7, 9935-9953.	0.8	3
50	Phylogenies and traits provide distinct insights about the historical and contemporary assembly of aquatic insect communities. Ecology and Evolution, 2016, 6, 2925-2937.	0.8	30
51	Measuring the functional redundancy of biological communities: a quantitative guide. Methods in Ecology and Evolution, 2016, 7, 1386-1395.	2.2	197
52	The Evolutionary Legacy of Diversification Predicts Ecosystem Function. American Naturalist, 2016, 188, 398-410.	1.0	14
53	A family of functional dissimilarity measures for presence and absence data. Ecology and Evolution, 2016, 6, 5383-5389.	0.8	16
54	â€~Equivalent numbers' for species, phylogenetic or functional diversity in a nested hierarchy of multiple scales. Methods in Ecology and Evolution, 2016, 7, 1152-1163.	2.2	30

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55	Integrating dataâ€deficient species in analyses of evolutionary history loss. Ecology and Evolution, 2016, 6, 8502-8514.	0.8	20
56	Loss and conservation of evolutionary history in the Mediterranean Basin. BMC Ecology, 2016, 16, 43.	3.0	12
57	A guide through a family of phylogenetic dissimilarity measures among sites. Oikos, 2016, 125, 1719-1732.	1.2	31
58	Measuring similarity among plots including similarity among species: an extension of traditional approaches. Journal of Vegetation Science, 2015, 26, 1061-1067.	1.1	13
59	Acoustic indices for biodiversity assessments: Analyses of bias based on simulated bird assemblages and recommendations for field surveys. Biological Conservation, 2015, 191, 306-312.	1.9	87
60	A multiple-site dissimilarity measure for species presence/absence data and its relationship with nestedness and turnover. Ecological Indicators, 2015, 54, 203-206.	2.6	20
61	Considering external information to improve the phylogenetic comparison of microbial communities: a new approach based on constrained Double Principal Coordinates Analysis ( <scp>cDPCoA</scp> ). Molecular Ecology Resources, 2015, 15, 242-249.	2.2	19
62	A cautionary note on some phylogenetic dissimilarity measures. Journal of Plant Ecology, 2015, 8, 12-16.	1.2	9
63	First description of underwater acoustic diversity in three temperate ponds. PeerJ, 2015, 3, e1393.	0.9	47
64	Statistical ecology comes of age. Biology Letters, 2014, 10, 20140698.	1.0	40
65	Acoustic Indices for Biodiversity Assessment and Landscape Investigation. Acta Acustica United With Acustica, 2014, 100, 772-781.	0.8	336
66	Combining the fourthâ€corner and the RLQ methods for assessing trait responses to environmental variation. Ecology, 2014, 95, 14-21.	1.5	398
67	Species living in harsh environments have low clade rank and are localized on former Laurasian continents: a case study of <i>Willemia</i> (Collembola). Journal of Biogeography, 2014, 41, 353-365.	1.4	3
68	Functional and phylogenetic similarity among communities. Methods in Ecology and Evolution, 2014, 5, 666-675.	2.2	53
69	New biodiversity measure that includes consistent interspecific and intraspecific components. Methods in Ecology and Evolution, 2014, 5, 165-172.	2.2	13
70	Temporal and spatial variability of animal sound within a neotropical forest. Ecological Informatics, 2014, 21, 133-143.	2.3	86
71	Monitoring temporal change of bird communities with dissimilarity acoustic indices. Methods in Ecology and Evolution, 2014, 5, 495-505.	2.2	69
72	Life history traits, but not phylogeny, drive compositional patterns in a butterfly metacommunity. Ecology, 2014, 95, 3304-3313.	1.5	31

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73	Correlations between phylogenetic and functional diversity: mathematical artefacts or true ecological and evolutionary processes?. Journal of Vegetation Science, 2013, 24, 781-793.	1.1	103
74	Does trait conservatism guarantee that indicators of phylogenetic community structure will reveal nicheâ€based assembly processes along stress gradients?. Journal of Vegetation Science, 2013, 24, 820-833.	1.1	31
75	Specialists leave fewer descendants within a region than generalists. Global Ecology and Biogeography, 2013, 22, 213-222.	2.7	23
76	TESTING FOR PHYLOGENETIC SIGNAL IN BIOLOGICAL TRAITS: THE UBIQUITY OF CROSS-PRODUCT STATISTICS. Evolution; International Journal of Organic Evolution, 2013, 67, 828-840.	1.1	38
77	A guide for using functional diversity indices to reveal changes in assembly processes along ecological gradients. Journal of Vegetation Science, 2013, 24, 794-806.	1.1	316
78	A New Technique for Analysing Interacting Factors Affecting Biodiversity Patterns: Crossed-DPCoA. PLoS ONE, 2013, 8, e54530.	1.1	14
79	Assessing biodiversity with sound: Do acoustic diversity indices reflect phylogenetic and functional diversities of bird communities?. Ecological Indicators, 2013, 25, 279-287.	2.6	143
80	Biodiversity Sampling Using a Global Acoustic Approach: Contrasting Sites with Microendemics in New Caledonia. PLoS ONE, 2013, 8, e65311.	1.1	88
81	A New Freshwater Biodiversity Indicator Based on Fish Community Assemblages. PLoS ONE, 2013, 8, e80968.	1.1	10
82	Ecophylogenetics: advances and perspectives. Biological Reviews, 2012, 87, 769-785.	4.7	341
83	Local gardening practices shape urban lawn floristic communities. Landscape and Urban Planning, 2012, 105, 53-61.	3.4	91
84	Monitoring animal diversity using acoustic indices: Implementation in a temperate woodland. Ecological Indicators, 2012, 13, 46-54.	2.6	294
85	Links between the species abundance distribution and the shape of the corresponding rank abundance curve. Ecological Indicators, 2012, 14, 1-6.	2.6	14
86	ASSESSING PHYLOGENETIC SIGNAL WITH MEASUREMENT ERROR: A COMPARISON OF MANTEL TESTS, BLOMBERG ET AL'S K, AND PHYLOGENETIC DISTOGRAMS. Evolution; International Journal of Organic Evolution, 2012, 66, 2614-2621.	1.1	59
87	Functional rarefaction for species abundance data. Methods in Ecology and Evolution, 2012, 3, 519-525.	2.2	40
88	Clarifying and developing analyses of biodiversity: towards a generalisation of current approaches. Methods in Ecology and Evolution, 2012, 3, 509-518.	2.2	35
89	Threat Diversity Will Erode Mammalian Phylogenetic Diversity in the Near Future. PLoS ONE, 2012, 7, e46235.	1.1	24
90	New concentration measures as kinds of the quadratic entropy. Ecological Indicators, 2011, 11, 540-544.	2.6	2

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91	Linking patterns in phylogeny, traits, abiotic variables and space: a novel approach to linking environmental filtering and plant community assembly. Journal of Ecology, 2011, 99, 165-175.	1.9	141
92	Diet and fuelling of the globally threatened aquatic warbler at autumn migration stopover as compared with two congeners. Animal Conservation, 2011, 14, 261-270.	1.5	17
93	Measuring biodiversity to explain community assembly: a unified approach. Biological Reviews, 2011, 86, 792-812.	4.7	489
94	Integrating functional diversity into tropical forest plantation designs to study ecosystem processes. Annals of Forest Science, 2010, 67, 303-303.	0.8	14
95	Putting phylogeny into the analysis of biological traits: A methodological approach. Journal of Theoretical Biology, 2010, 264, 693-701.	0.8	60
96	Using biological traits to assess how urbanization filters plant species of small woodlands. Applied Vegetation Science, 2010, 13, 412-424.	0.9	62
97	Decomposition of trait diversity among the nodes of a phylogenetic tree. Ecological Monographs, 2010, 80, 485-507.	2.4	72
98	Variation within and between Closely Related Species Uncovers High Intra-Specific Variability in Dispersal. PLoS ONE, 2010, 5, e11123.	1.1	80
99	Host range as an axis of niche partitioning in the plant-feeding nematode community of banana agroecosystems. Soil Biology and Biochemistry, 2009, 41, 1139-1145.	4.2	13
100	Biological diversity: Distinct distributions can lead to the maximization of Rao's quadratic entropy. Theoretical Population Biology, 2009, 75, 153-163.	0.5	19
101	On the challenge of treating various types of variables: application for improving the measurement of functional diversity. Oikos, 2009, 118, 391-402.	1.2	473
102	Hierarchical partitioning of evolutionary and ecological patterns in the organization of phylogeneticallyâ€structured species assemblages: application to rockfish (genus: <i>Sebastes</i> ) in the Southern California Bight. Ecology Letters, 2009, 12, 898-908.	3.0	71
103	Corrigendum. Ecology Letters, 2009, 12, 999-999.	3.0	21
104	Relationships between channelization structures, environmental characteristics, and plant communities in four French streams in the Seine–Normandy catchment. Journal of the North American Benthological Society, 2009, 28, 596-610.	3.0	14
105	Testing for phylogenetic signal in phenotypic traits: New matrices of phylogenetic proximities. Theoretical Population Biology, 2008, 73, 79-91.	0.5	111
106	Can hostâ€range allow niche differentiation of invasive polyphagous fruit flies (Diptera: Tephritidae) in La Réunion?. Ecological Entomology, 2008, 33, 439-452.	1.1	38
107	Rapid Acoustic Survey for Biodiversity Appraisal. PLoS ONE, 2008, 3, e4065.	1.1	448
108	A NEW TECHNIQUE FOR ORDERING ASYMMETRICAL THREE-DIMENSIONAL DATA SETS IN ECOLOGY. Ecology, 2007, 88, 512-523.	1.5	10

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109	New analysis for consistency among markers in the study of genetic diversity: development and application to the description of bacterial diversity. BMC Evolutionary Biology, 2007, 7, 156.	3.2	10
110	Is the originality of a species measurable?. Ecology Letters, 2005, 8, 579-586.	3.0	168
111	The apportionment of quadratic entropy: a useful alternative for partitioning diversity in ecological data. Environmental and Ecological Statistics, 2005, 12, 125-138.	1.9	103
112	Measuring diversity from dissimilarities with Rao's quadratic entropy: Are any dissimilarities suitable?. Theoretical Population Biology, 2005, 67, 231-239.	0.5	116
113	From dissimilarities among species to dissimilarities among communities: a double principal coordinate analysis. Journal of Theoretical Biology, 2004, 228, 523-537.	0.8	184