

Pierre Legendre

List of Publications by Year in descending order

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Version: 2024-02-01

359
papers

67,947
citations

3919

88
h-index

813

246
g-index

393
all docs

393
docs citations

393
times ranked

51076
citing authors

#	ARTICLE	IF	CITATIONS
1	Evaluating ecological uniqueness over broad spatial extents using species distribution modelling. <i>Oikos</i> , 2022, 2022, .	1.2	12
2	New measures for quantifying directional changes in presence-absence community data. <i>Ecological Indicators</i> , 2022, 136, 108618.	2.6	3
3	Large-scale multi-trophic co-response models and environmental control of pelagic food webs in QuAbec lakes. <i>Oikos</i> , 2021, 130, 377-395.	1.2	4
4	Horizontal gene transfer and recombination analysis of SARS-CoV-2 genes helps discover its close relatives and shed light on its origin. <i>Bmc Ecology and Evolution</i> , 2021, 21, 5.	0.7	44
5	The phyllosphere microbiome of host trees contributes more than leaf phytochemicals to variation in the <i>Agrilus planipennis</i> Fairmaire gut microbiome structure. <i>Scientific Reports</i> , 2021, 11, 15911.	1.6	10
6	Temperature drives local contributions to beta diversity in mountain streams: Stochastic and deterministic processes. <i>Global Ecology and Biogeography</i> , 2020, 29, 420-432.	2.7	30
7	A novel tool to assess the effect of intraspecific spatial niche variation on species distribution shifts under climate change. <i>Global Ecology and Biogeography</i> , 2020, 29, 590-602.	2.7	12
8	Partitioning plant spectral diversity into alpha and beta components. <i>Ecology Letters</i> , 2020, 23, 370-380.	3.0	62
9	Moderate disturbances accelerate forest transition dynamics under climate change in the temperate-boreal ecotone of eastern North America. <i>Global Change Biology</i> , 2020, 26, 4418-4435.	4.2	44
10	What do beta diversity components reveal from presence-absence community data? Let us connect every indicator to an indicandum!. <i>Ecological Indicators</i> , 2020, 117, 106540.	2.6	25
11	Does diversity beget diversity in microbiomes?. <i>ELife</i> , 2020, 9, .	2.8	33
12	Disturbances amplify tree community responses to climate change in the temperate-boreal ecotone. <i>Global Ecology and Biogeography</i> , 2019, 28, 1668-1681.	2.7	67
13	The interaction of phylogeny and community structure: Linking the community composition and trait evolution of clades. <i>Global Ecology and Biogeography</i> , 2019, 28, 1499-1511.	2.7	14
14	Trait-based approach to monitoring marine benthic data along 500 km of coastline. <i>Diversity and Distributions</i> , 2019, 25, 1879-1896.	1.9	35
15	A temporal beta-diversity index to identify sites that have changed in exceptional ways in space-time surveys. <i>Ecology and Evolution</i> , 2019, 9, 3500-3514.	0.8	137
16	Damming interacts with the flood pulse to alter zooplankton communities in an Amazonian river. <i>Freshwater Biology</i> , 2019, 64, 1040-1053.	1.2	19
17	Spatial and temporal analysis of beta diversity in the Barro Colorado Island forest dynamics plot, Panama. <i>Forest Ecosystems</i> , 2019, 6, .	1.3	33
18	Variation in compositional and structural components of community assemblage and its determinants. <i>Journal of Vegetation Science</i> , 2019, 30, 257-268.	1.1	9

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19	Trajectory analysis in community ecology. <i>Ecological Monographs</i> , 2019, 89, e01350.	2.4	74
20	<i>Numerical Ecology</i> . , 2019, , 487-493.		30
21	Flow alterations by dams shaped fish assemblage dynamics in the complex Mekong-3S river system. <i>Ecological Indicators</i> , 2018, 88, 103-114.	2.6	73
22	Box-Cox chord transformations for community composition data prior to beta diversity analysis. <i>Ecography</i> , 2018, 41, 1820-1824.	2.1	67
23	Uniqueness of sampling site contributions to the total variance of macroinvertebrate communities in the Lower Mekong Basin. <i>Ecological Indicators</i> , 2018, 84, 425-432.	2.6	22
24	Spatial organisation of fish communities in the St. Lawrence River: a test for longitudinal gradients and spatial heterogeneities in a large river system. <i>Hydrobiologia</i> , 2018, 809, 155-173.	1.0	17
25	Concomitant impacts of climate change, fragmentation and non-native species have led to reorganization of fish communities since the 1980s. <i>Global Ecology and Biogeography</i> , 2018, 27, 213-222.	2.7	56
26	Bringing multivariate support to multiscale codependence analysis: Assessing the drivers of community structure across spatial scales. <i>Methods in Ecology and Evolution</i> , 2018, 9, 292-304.	2.2	7
27	Biodiversity and trophic ecology of hydrothermal vent fauna associated with tubeworm assemblages on the Juan de Fuca Ridge. <i>Biogeosciences</i> , 2018, 15, 2629-2647.	1.3	18
28	Negative relationships between species richness and temporal variability are common but weak in natural systems. <i>Ecology</i> , 2018, 99, 2592-2604.	1.5	26
29	Summer assessment of zooplankton biodiversity and environmental control in urban waterbodies on the Island of Montréal. <i>Ecosphere</i> , 2018, 9, e02277.	1.0	22
30	Application of Moran Eigenvector Maps (MEM) to irregular sampling designs. <i>Spatial Statistics</i> , 2018, 26, 56-68.	0.9	13
31	<i>Numerical Ecology with R</i> . Use R!, 2018, , .	0.3	439
32	<i>Association Measures and Matrices</i> . Use R!, 2018, , 35-57.	0.3	4
33	<i>Unconstrained Ordination</i> . Use R!, 2018, , 151-201.	0.3	10
34	<i>Canonical Ordination</i> . Use R!, 2018, , 203-297.	0.3	21
35	<i>Spatial Analysis of Ecological Data</i> . Use R!, 2018, , 299-367.	0.3	20
36	<i>Community Diversity</i> . Use R!, 2018, , 369-412.	0.3	4

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37	The Effects of Regional Hydrologic Alteration on Fish Community Structure in Regulated Rivers. <i>River Research and Applications</i> , 2017, 33, 249-257.	0.7	22
38	Predicting microcystin concentrations in lakes and reservoirs at a continental scale: A new framework for modelling an important health risk factor. <i>Global Ecology and Biogeography</i> , 2017, 26, 625-637.	2.7	59
39	Human and natural controls of the variation in aboveground tree biomass in African dry tropical forests. , 2017, 27, 1578-1593.		9
40	Hosts, parasites and their interactions respond to different climatic variables. <i>Global Ecology and Biogeography</i> , 2017, 26, 942-951.	2.7	62
41	Constancy despite variability: Local and regional macrofaunal diversity in intertidal seagrass beds. <i>Journal of Sea Research</i> , 2017, 130, 107-122.	0.6	21
42	Astronomical and atmospheric impacts on deep-sea hydrothermal vent invertebrates. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2017, 284, 20162123.	1.2	29
43	Diatom diversity patterns over the past <i>c</i>. 150 years across the conterminous United States of America: Identifying mechanisms behind beta diversity. <i>Global Ecology and Biogeography</i> , 2017, 26, 1303-1315.	2.7	40
44	Modelling habitat distributions for multiple species using phylogenetics. <i>Ecography</i> , 2017, 40, 1088-1097.	2.1	2
45	Environmental factors structuring benthic primary producers at different spatial scales in the St. Lawrence River (Canada). <i>Aquatic Sciences</i> , 2017, 79, 345-356.	0.6	10
46	Biological and environmental rhythms in (dark) deep-sea hydrothermal ecosystems. <i>Biogeosciences</i> , 2017, 14, 2955-2977.	1.3	26
47	The Willow Microbiome Is Influenced by Soil Petroleum-Hydrocarbon Concentration with Plant Compartment-Specific Effects. <i>Frontiers in Microbiology</i> , 2016, 7, 1363.	1.5	75
48	A spatiallyâ€œexplicit assessment of the fish population response to flow management in a heterogeneous landscape. <i>Ecosphere</i> , 2016, 7, e01252.	1.0	7
49	A new costâ€œeffective approach to survey ecological communities. <i>Oikos</i> , 2016, 125, 975-987.	1.2	12
50	Using fish guilds to assess community responses to temperature and flow regimes in unregulated and regulated Canadian rivers. <i>Freshwater Biology</i> , 2016, 61, 1759-1772.	1.2	12
51	Multi-scale spatial and partitioning analyses of the reef-fish community composition of the Yucatan fringing reef system. <i>Ecological Complexity</i> , 2016, 28, 69-76.	1.4	1
52	Should the Mantel test be used in spatial analysis?. <i>Methods in Ecology and Evolution</i> , 2015, 6, 1239-1247.	2.2	276
53	A Comparison of Electrofishing and Visual Surveying Methods for Estimating Fish Community Structure in Temperate Rivers. <i>River Research and Applications</i> , 2015, 31, 1040-1051.	0.7	19
54	Understanding the Spatio-Temporal Response of Coral Reef Fish Communities to Natural Disturbances: Insights from Beta-Diversity Decomposition. <i>PLoS ONE</i> , 2015, 10, e0138696.	1.1	54

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55	Phylogenetics to help predict active metabolism. <i>Ecosphere</i> , 2015, 6, 1-11.	1.0	4
56	Thirty-year recovery of mollusc communities after nuclear experimentations on Fangataufa atoll (Tuamotu, French Polynesia). <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2015, 282, 20150750.	1.2	22
57	Biodiversity patterns, environmental drivers and indicator species on a high-temperature hydrothermal edifice, Mid-Atlantic Ridge. <i>Deep-Sea Research Part II: Topical Studies in Oceanography</i> , 2015, 121, 177-192.	0.6	76
58	Are the landscape-level drivers of water column and surface sediment diatoms different?. <i>Freshwater Biology</i> , 2015, 60, 267-281.	1.2	17
59	Using intra-individual variation in shrub architecture to explain population cover. <i>Oikos</i> , 2015, 124, 707-716.	1.2	10
60	Oxidative stress modulates the expression of genes involved in cell survival in F508 cystic fibrosis airway epithelial cells. <i>Physiological Genomics</i> , 2014, 46, 634-646.	1.0	20
61	Reconstructing phosphorus levels using models based on the modern diatom assemblages of 55 lakes in southern Quebec. <i>Canadian Journal of Fisheries and Aquatic Sciences</i> , 2014, 71, 887-914.	0.7	14
62	Ward's Hierarchical Agglomerative Clustering Method: Which Algorithms Implement Ward's Criterion?. <i>Journal of Classification</i> , 2014, 31, 274-295.	1.2	2,398
63	Statistical methods for temporal and space-time analysis of community composition data <sup />. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2014, 281, 20132728.	1.2	197
64	Interpreting the replacement and richness difference components of beta diversity. <i>Global Ecology and Biogeography</i> , 2014, 23, 1324-1334.	2.7	705
65	Using phylogenetic information and chemical properties to predict species tolerances to pesticides. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2014, 281, 20133239.	1.2	28
66	Consensus RDA across dissimilarity coefficients for canonical ordination of community composition data. <i>Ecological Monographs</i> , 2014, 84, 491-511.	2.4	31
67	High-resolution dynamics of a deep-sea hydrothermal mussel assemblage monitored by the EMSO-AÅšores MoMAR observatory. <i>Deep-Sea Research Part I: Oceanographic Research Papers</i> , 2014, 90, 62-75.	0.6	29
68	Rhythms and Community Dynamics of a Hydrothermal Tubeworm Assemblage at Main Endeavour Field – A Multidisciplinary Deep-Sea Observatory Approach. <i>PLoS ONE</i> , 2014, 9, e96924.	1.1	55
69	Indicator Species: Computation. , 2013, , 264-268.		22
70	Phylogenetic eigenvector maps: a framework to model and predict species traits. <i>Methods in Ecology and Evolution</i> , 2013, 4, 1120-1131.	2.2	91
71	Dissimilarity measurements and the size structure of ecological communities. <i>Methods in Ecology and Evolution</i> , 2013, 4, 1167-1177.	2.2	50
72	Large-scale geographic patterns of diversity and community structure of pelagic crustacean zooplankton in Canadian lakes. <i>Global Ecology and Biogeography</i> , 2013, 22, 784-795.	2.7	63

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73	Optimization of temporal versus spatial replication in the development of habitat use models to explain among-reach variations of fish density estimates in rivers. <i>Canadian Journal of Fisheries and Aquatic Sciences</i> , 2013, 70, 600-609.	0.7	4
74	Putting the landscape into the genomics of trees: approaches for understanding local adaptation and population responses to changing climate. <i>Tree Genetics and Genomes</i> , 2013, 9, 901-911.	0.6	261
75	Beta diversity as the variance of community data: dissimilarity coefficients and partitioning. <i>Ecology Letters</i> , 2013, 16, 951-963.	3.0	937
76	Potential changes in forest composition could reduce impacts of climate change on boreal wildfires. <i>Ecological Applications</i> , 2013, 23, 21-35.	1.8	117
77	Genetic structure of the white-footed mouse in the context of the emergence of Lyme disease in southern Québec. <i>Ecology and Evolution</i> , 2013, 3, 2075-2088.	0.8	34
78	Examining shifts in zooplankton community variability following biological invasion. <i>Limnology and Oceanography</i> , 2013, 58, 399-408.	1.6	7
79	Living in a hot redox soup: antioxidant defences of the hydrothermal worm <i>Alvinella pompejana</i> . <i>Aquatic Biology</i> , 2013, 18, 217-228.	0.5	28
80	An Efficient Algorithm for the Detection and Classification of Horizontal Gene Transfer Events and Identification of Mosaic Genes. <i>Studies in Classification, Data Analysis, and Knowledge Organization</i> , 2013, , 253-260.	0.1	2
81	Spatial and Temporal Variation in a Caribbean Herbivorous Fish Assemblage. <i>Journal of Coastal Research</i> , 2012, 278, 63-72.	0.1	14
82	N ₂ fixation rates and associated diversity (nifH) of microbialite and mat-forming consortia from different aquatic environments in Mexico. <i>Aquatic Microbial Ecology</i> , 2012, 67, 15-24.	0.9	26
83	Microbialite genetic diversity and composition relate to environmental variables. <i>FEMS Microbiology Ecology</i> , 2012, 82, 724-735.	1.3	46
84	Ecological data series. <i>Developments in Environmental Modelling</i> , 2012, , 711-783.	0.3	1
85	Effects of spatial scale and choice of statistical model (linear versus tree-based) on determining species-habitat relationships. <i>Canadian Journal of Fisheries and Aquatic Sciences</i> , 2012, 69, 2095-2111.	0.7	19
86	Clustering and Partitioning. <i>Developments in Paleoenvironmental Research</i> , 2012, , 167-200.	7.5	31
87	From Classical to Canonical Ordination. <i>Developments in Paleoenvironmental Research</i> , 2012, , 201-248.	7.5	112
88	Inferring Processes from Spatial Patterns: The Role of Directional and Non-Directional Forces in Shaping Fish Larvae Distribution in a Freshwater Lake System. <i>PLoS ONE</i> , 2012, 7, e50239.	1.1	29
89	Variation partitioning involving orthogonal spatial eigenfunction submodels. <i>Ecology</i> , 2012, 93, 1234-1240.	1.5	92
90	Community ecology in the age of multivariate multiscale spatial analysis. <i>Ecological Monographs</i> , 2012, 82, 257-275.	2.4	506

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91	Using species combinations in indicator value analyses. <i>Methods in Ecology and Evolution</i> , 2012, 3, 973-982.	2.2	224
92	Complex ecological data sets. <i>Developments in Environmental Modelling</i> , 2012, , 1-57.	0.3	28
93	Dimensional analysis in ecology. <i>Developments in Environmental Modelling</i> , 2012, 24, 109-142.	0.3	3
94	Multidimensional quantitative data. <i>Developments in Environmental Modelling</i> , 2012, , 143-194.	0.3	2
95	Multidimensional semiquantitative data. <i>Developments in Environmental Modelling</i> , 2012, 24, 195-218.	0.3	2
96	Multidimensional qualitative data. <i>Developments in Environmental Modelling</i> , 2012, 24, 219-264.	0.3	6
97	Ecological resemblance. <i>Developments in Environmental Modelling</i> , 2012, 24, 265-335.	0.3	23
98	Ordination in reduced space. <i>Developments in Environmental Modelling</i> , 2012, , 425-520.	0.3	32
99	Interpretation of ecological structures. <i>Developments in Environmental Modelling</i> , 2012, 24, 521-624.	0.3	14
100	Canonical analysis. <i>Developments in Environmental Modelling</i> , 2012, 24, 625-710.	0.3	48
101	Spatial analysis. <i>Developments in Environmental Modelling</i> , 2012, 24, 785-858.	0.3	9
102	Multiscale analysis. <i>Developments in Environmental Modelling</i> , 2012, 24, 859-906.	0.3	7
103	Is the Mantel correlogram powerful enough to be useful in ecological analysis? A simulation study. <i>Ecology</i> , 2012, 93, 1473-1481.	1.5	161
104	Broad-scale adaptive genetic variation in alpine plants is driven by temperature and precipitation. <i>Molecular Ecology</i> , 2012, 21, 3729-3738.	2.0	161
105	Disentangling invasion processes in a dynamic shipping-boating network. <i>Molecular Ecology</i> , 2012, 21, 4227-4241.	2.0	35
106	The variation of tree beta diversity across a global network of forest plots. <i>Global Ecology and Biogeography</i> , 2012, 21, 1191-1202.	2.7	135
107	Cascade multivariate regression tree: a novel approach for modelling nested explanatory sets. <i>Methods in Ecology and Evolution</i> , 2012, 3, 234-244.	2.2	23
108	Multiple-Table Data in R with the multitablePackage. <i>Journal of Statistical Software</i> , 2012, 51, .	1.8	3

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109	NSERC's HydroNet: A National Research Network to Promote Sustainable Hydropower and Healthy Aquatic Ecosystems. <i>Fisheries</i> , 2011, 36, 480-488.	0.6	11
110	Using phylogenetic information to predict species tolerances to toxic chemicals. , 2011, 21, 3178-3190.		54
111	Numerical Ecology with R. , 2011, , .		1,684
112	Association Measures and Matrices. , 2011, , 31-51.		7
113	Testing the significance of canonical axes in redundancy analysis. <i>Methods in Ecology and Evolution</i> , 2011, 2, 269-277.	2.2	459
114	Relationships between species feeding traits and environmental conditions in fish communities: a three-matrix approach. , 2011, 21, 363-377.		46
115	Business partner or simple catch? The economic value of the sicklefin lemon shark in French Polynesia. <i>Marine and Freshwater Research</i> , 2011, 62, 764.	0.7	67
116	Faunal changes and geographic crypticism indicate the occurrence of a biogeographic transition zone along the southern East Pacific Rise. <i>Journal of Biogeography</i> , 2011, 38, 575-594.	1.4	26
117	Scale dependency of processes structuring metacommunities of cladocerans in temporary pools of High-Andes wetlands. <i>Ecography</i> , 2011, 34, 296-305.	2.1	174
118	The role of environmental and spatial processes in structuring native and non-native fish communities across thousands of lakes. <i>Ecography</i> , 2011, 34, 762-771.	2.1	60
119	A framework for estimating niche metrics using the resemblance between qualitative resources. <i>Oikos</i> , 2011, 120, 1341-1350.	1.2	63
120	Organochlorine pollution in tropical rivers (Guadeloupe): Role of ecological factors in food web bioaccumulation. <i>Environmental Pollution</i> , 2011, 159, 1692-1701.	3.7	108
121	Modelling the effect of directional spatial ecological processes at different scales. <i>Oecologia</i> , 2011, 166, 357-368.	0.9	114
122	Diversity and composition of ectomycorrhizal community on seedling roots: the role of host preference and soil origin. <i>Mycorrhiza</i> , 2011, 21, 669-680.	1.3	54
123	The performance of the Congruence Among Distance Matrices (CADM) test in phylogenetic analysis. <i>BMC Evolutionary Biology</i> , 2011, 11, 64.	3.2	93
124	Canonical Ordination. , 2011, , 153-225.		39
125	Unconstrained Ordination. , 2011, , 115-151.		21
126	Modelling habitat associations of 14 species of holothurians from an unfished coral atoll: implications for fisheries management. <i>Aquatic Biology</i> , 2011, 14, 57-66.	0.5	20

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127	Modelling habitat associations of the common spider conch in the Cocos (Keeling) Islands. <i>Marine Ecology - Progress Series</i> , 2011, 432, 83-90.	0.9	5
128	Reefscape proxies for the conservation of Caribbean coral reef biodiversity. <i>Ciencias Marinas</i> , 2011, 37, 87-96.	0.4	20
129	Water table response to an experimental alley farming trial: dissecting the spatial and temporal structure of the data. <i>Ecological Applications</i> , 2010, 20, 1704-1720.	1.8	5
130	Weighted bootstrapping: a correction method for assessing the robustness of phylogenetic trees. <i>BMC Evolutionary Biology</i> , 2010, 10, 250.	3.2	7
131	The relations between "standard" fluvial habitat variables and turbulent flow at multiple scales in morphological units of a gravel-bed river. <i>River Research and Applications</i> , 2010, 26, 439-455.	0.7	18
132	Aggregation of Sampling Units: An Analytical Solution to Predict Variance. <i>Geographical Analysis</i> , 2010, 29, 258-266.	1.9	25
133	Improving indicator species analysis by combining groups of sites. <i>Oikos</i> , 2010, 119, 1674-1684.	1.2	1,041
134	Fire-induced taxonomic and functional changes in saproxylic beetle communities in fire sensitive regions. <i>Ecography</i> , 2010, 33, 760-771.	2.1	59
135	Utility of computer simulations in landscape genetics. <i>Molecular Ecology</i> , 2010, 19, 3549-3564.	2.0	155
136	Common factors drive adaptive genetic variation at different spatial scales in <i>Arabis alpina</i> . <i>Molecular Ecology</i> , 2010, 19, 3824-3835.	2.0	188
137	Estimating and controlling for spatial structure in the study of ecological communities. <i>Global Ecology and Biogeography</i> , 2010, 19, 174-184.	2.7	370
138	A distance-based framework for measuring functional diversity from multiple traits. <i>Ecology</i> , 2010, 91, 299-305.	1.5	2,787
139	Behavioural response of sicklefin lemon sharks <i>Negaprion acutidens</i> to underwater feeding for ecotourism purposes. <i>Marine Ecology - Progress Series</i> , 2010, 414, 257-266.	0.9	110
140	Spider, bee, and bird communities in cities are shaped by environmental control and high stochasticity. <i>Ecology</i> , 2010, 91, 3343-3353.	1.5	109
141	Multiscale codependence analysis: an integrated approach to analyze relationships across scales. <i>Ecology</i> , 2010, 91, 2952-2964.	1.5	26
142	Spatial relationships between soil moisture patterns and topographic variables at multiple scales in a humid temperate forested catchment. <i>Water Resources Research</i> , 2010, 46, .	1.7	34
143	Community surveys through space and time: testing the space-time interaction in the absence of replication. <i>Ecology</i> , 2010, 91, 262-272.	1.5	84
144	Comparison of the Mantel test and alternative approaches for detecting complex multivariate relationships in the spatial analysis of genetic data. <i>Molecular Ecology Resources</i> , 2010, 10, 831-844.	2.2	553

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145	Effects of provisioning on shark behaviour: Reply to Brunnschweiler & McKenzie (2010). <i>Marine Ecology - Progress Series</i> , 2010, 420, 285-288.	0.9	3
146	Global depression in gene expression as a response to rapid thermal changes in vent mussels. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2009, 276, 3071-3079.	1.2	49
147	Assessing the scale-specific importance of niches and other spatial processes on beta diversity: a case study from a temperate forest. <i>Oecologia</i> , 2009, 159, 377-388.	0.9	136
148	Assessing Congruence Among Ultrametric Distance Matrices. <i>Journal of Classification</i> , 2009, 26, 103-117.	1.2	17
149	Comparison of two plant functional approaches to evaluate natural restoration along an old-field deciduous forest chronosequence. <i>Journal of Vegetation Science</i> , 2009, 20, 185-198.	1.1	55
150	Independent contrasts and regression through the origin. <i>Journal of Theoretical Biology</i> , 2009, 259, 727-743.	0.8	18
151	Associations between species and groups of sites: indices and statistical inference. <i>Ecology</i> , 2009, 90, 3566-3574.	1.5	2,649
152	Shifts between biotic and physical driving forces of species organization under natural disturbance regimes. <i>Canadian Journal of Fisheries and Aquatic Sciences</i> , 2009, 66, 1282-1293.	0.7	26
153	Biogeographic relationships among deep-sea hydrothermal vent faunas at global scale. <i>Deep-Sea Research Part I: Oceanographic Research Papers</i> , 2009, 56, 1371-1378.	0.6	137
154	Partitioning beta diversity in a subtropical broad-leaved forest of China. <i>Ecology</i> , 2009, 90, 663-674.	1.5	520
155	Using the landscape morphometric context to resolve spatial patterns of submerged macrophyte communities in a fluvial lake. <i>Landscape Ecology</i> , 2008, 23, 91-105.	1.9	24
156	Beals smoothing revisited. <i>Oecologia</i> , 2008, 156, 657-669.	0.9	42
157	Explaining variation in tropical plant community composition: influence of environmental and spatial data quality. <i>Oecologia</i> , 2008, 155, 593-604.	0.9	178
158	The utility of covariances: a response to Ranta et al. <i>Oikos</i> , 2008, 117, 1912-1913.	1.2	5
159	Phylogenetic, functional, and structural components of variation in bone growth rate of amniotes. <i>Evolution & Development</i> , 2008, 10, 217-227.	1.1	83
160	Modelling directional spatial processes in ecological data. <i>Ecological Modelling</i> , 2008, 215, 325-336.	1.2	261
161	Scaling up beta diversity on Caribbean coral reefs. <i>Journal of Experimental Marine Biology and Ecology</i> , 2008, 366, 28-36.	0.7	46
162	ANALYZING OR EXPLAINING BETA DIVERSITY? COMMENT. <i>Ecology</i> , 2008, 89, 3238-3244.	1.5	81

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163	Studying beta diversity: ecological variation partitioning by multiple regression and canonical analysis. <i>Journal of Plant Ecology</i> , 2008, 1, 3-8.	1.2	405
164	Meiofaunal community structure of the deep-sea Gulf of Mexico: Variability due to the sorting methods. <i>Deep-Sea Research Part II: Topical Studies in Oceanography</i> , 2008, 55, 2627-2633.	0.6	15
165	Epibenthic megacrustaceans from the continental margin, slope and abyssal plain of the Southwestern Gulf of Mexico: Factors responsible for variability in species composition and diversity. <i>Deep-Sea Research Part II: Topical Studies in Oceanography</i> , 2008, 55, 2667-2678.	0.6	22
166	Macrofaunal density and biomass in the Campeche Canyon, Southwestern Gulf of Mexico. <i>Deep-Sea Research Part II: Topical Studies in Oceanography</i> , 2008, 55, 2679-2685.	0.6	31
167	TESTING THE SPECIES TRAITSâ€“ENVIRONMENT RELATIONSHIPS: THE FOURTHâ€“CORNER PROBLEM REVISITED. <i>Ecology</i> , 2008, 89, 3400-3412.	1.5	495
168	FORWARD SELECTION OF EXPLANATORY VARIABLES. <i>Ecology</i> , 2008, 89, 2623-2632.	1.5	1,766
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