

# Brent C Emerson

## List of Publications by Year in descending order

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133  
papers

7,778  
citations

57758

44  
h-index

58581

82  
g-index

138  
all docs

138  
docs citations

138  
times ranked

9938  
citing authors

#	ARTICLE	IF	CITATIONS
1	Community metabarcoding reveals the relative role of environmental filtering and spatial processes in metacommunity dynamics of soil microarthropods across a mosaic of montane forests. <i>Molecular Ecology</i> , 2023, 32, 6110-6128.	3.9	15
2	Coming of age for COI metabarcoding of whole organism community DNA: Towards bioinformatic harmonisation. <i>Molecular Ecology Resources</i> , 2022, 22, 847-861.	4.8	22
3	Dispersal ability and its consequences for population genetic differentiation and diversification. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2022, 289, 20220489.	2.6	14
4	Hidden island endemic species and their implications for cryptic speciation within soil arthropods. <i>Journal of Biogeography</i> , 2022, 49, 1367-1380.	3.0	9
5	Community assembly and metapopulogeography of soil biodiversity: Insights from haplotype-level community DNA metabarcoding within an oceanic island. <i>Molecular Ecology</i> , 2022, 31, 4078-4094.	3.9	9
6	The limited spatial scale of dispersal in soil arthropods revealed with whole community haplotype-level metabarcoding. <i>Molecular Ecology</i> , 2021, 30, 48-61.	3.9	49
7	Long-term cloud forest response to climate warming revealed by insect speciation history*. <i>Evolution; International Journal of Organic Evolution</i> , 2021, 75, 231-244.	2.3	9
8	Sharing and reporting benefits from biodiversity research. <i>Molecular Ecology</i> , 2021, 30, 1103-1107.	3.9	19
9	Flightlessness in insects enhances diversification and determines assemblage structure across whole communities. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2021, 288, 20202646.	2.6	13
10	Connecting high-throughput biodiversity inventories: Opportunities for a site-based genomic framework for global integration and synthesis. <i>Molecular Ecology</i> , 2021, 30, 1120-1135.	3.9	26
11	Genomic variation, population history and within-archipelago adaptation between island bird populations. <i>Royal Society Open Science</i> , 2021, 8, 201146.	2.4	9
12	Validated removal of nuclear pseudogenes and sequencing artefacts from mitochondrial metabarcode data. <i>Molecular Ecology Resources</i> , 2021, 21, 1772-1787.	4.8	32
13	Evaluating species origins within tropical sky-islands arthropod communities. <i>Journal of Biogeography</i> , 2021, 48, 2199-2210.	3.0	7
14	Biodiversity monitoring using environmental DNA. <i>Molecular Ecology Resources</i> , 2021, 21, 1405-1409.	4.8	15
15	A unified model of species abundance, genetic diversity, and functional diversity reveals the mechanisms structuring ecological communities. <i>Molecular Ecology Resources</i> , 2021, 21, 2782-2800.	4.8	24
16	Dispersal limitations and long-term persistence drive differentiation from haplotypes to communities within a tropical sky-island: Evidence from community metabarcoding. <i>Molecular Ecology</i> , 2021, 30, 6611-6626.	3.9	6
17	Habitat filtering and inferred dispersal ability condition across scale species turnover and rarity in Macaronesian island spider assemblages. <i>Journal of Biogeography</i> , 2021, 48, 3131-3144.	3.0	5
18	Mitochondrial Metagenomics Reveals the Ancient Origin and Phylogenetic Diversity of Soil Mites and Provides a Phylogeny of the Acari. <i>Molecular Biology and Evolution</i> , 2020, 37, 683-694.	8.9	42

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19	Climate drives community-wide divergence within species over a limited spatial scale: evidence from an oceanic island. <i>Ecology Letters</i> , 2020, 23, 305-315.	6.4	28
20	Building a Robust, Densely-Sampled Spider Tree of Life for Ecosystem Research. <i>Diversity</i> , 2020, 12, 288.	1.7	14
21	Revealing community assembly through barcoding: Mediterranean butterflies and dispersal variation. <i>Journal of Animal Ecology</i> , 2020, 89, 1992-1996.	2.8	0
22	Dispersal limitation: Evolutionary origins and consequences in arthropods. <i>Molecular Ecology</i> , 2019, 28, 3137-3140.	3.9	5
23	Unifying macroecology and macroevolution to answer fundamental questions about biodiversity. <i>Global Ecology and Biogeography</i> , 2019, 28, 1925-1936.	5.8	44
24	A topoclimate model for Quaternary insular speciation. <i>Journal of Biogeography</i> , 2019, 46, 2769-2786.	3.0	8
25	Evidence for the Pleistocene persistence of Collembola in Great Britain. <i>Journal of Biogeography</i> , 2019, 46, 1479-1493.	3.0	1
26	An integrated model of population genetics and community ecology. <i>Journal of Biogeography</i> , 2019, 46, 816-829.	3.0	37
27	Evolution of host plant use and diversification in a species complex of parasitic weevils (Coleoptera: Tj ETQq1 1 0.784314 rgBT / Overlock 10 Tf 50 62 Td	2.0	4
28	Introgressive hybridization in a Spiny-Tailed Iguana, <i>Ctenosaura pectinata</i> , and its implications for taxonomy and conservation. <i>PeerJ</i> , 2019, 7, e6744.	2.0	4
29	Anagenesis, Cladogenesis, and Speciation on Islands. <i>Trends in Ecology and Evolution</i> , 2018, 33, 488-491.	8.7	22
30	Long-term in situ persistence of biodiversity in tropical sky islands revealed by landscape genomics. <i>Molecular Ecology</i> , 2018, 27, 432-448.	3.9	39
31	Babies, Bathwater, and Straw Men? Not Quite: A Response to Meiri et al.. <i>Trends in Ecology and Evolution</i> , 2018, 33, 896-897.	8.7	2
32	Community assembly and diversification in a species-rich radiation of island weevils (Coleoptera: Tj ETQq0 0 0 rgBT / Overlock 10 Tf 50 62 Td	3.0	12
33	Statistical Evaluation of Monophyly in the "Broad-Nosed Weevils"™ through Molecular Phylogenetic Analysis Combining Mitochondrial Genome and Single-Locus Sequences (Curculionidae: Entiminae,) Tj ETQq1 1 0.784314 rgBT / Overlock 10 Tf 50 62 Td	3.0	12
34	Global Island Monitoring Scheme (GIMS): a proposal for the long-term coordinated survey and monitoring of native island forest biota. <i>Biodiversity and Conservation</i> , 2018, 27, 2567-2586.	2.6	72
35	Why the COI barcode should be the community <sc>DNA</sc> metabarcode for the metazoa. <i>Molecular Ecology</i> , 2018, 27, 3968-3975.	3.9	131
36	Lack of support for Rensch's rule in an intraspecific test using red flour beetle ( <i>Tribolium</i> ) Tj ETQq0 0 0 rgBT / Overlock 10 Tf 50 62 Td	3.0	12

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37	Comment on Rieux and Balloux: calibration from tip-dating can compromise topological accuracy and evolutionary inference. <i>Molecular Ecology</i> , 2017, 26, 2623-2624.	3.9	1
38	MtDNA metagenomics reveals large-scale invasion of belowground arthropod communities by introduced species. <i>Molecular Ecology</i> , 2017, 26, 3104-3115.	3.9	47
39	Cryptic diversity in the Azorean beetle genus <i>Tarphius</i> Erichson, 1845 (Coleoptera: Zopheridae): An integrative taxonomic approach with description of four new species. <i>Zootaxa</i> , 2017, 4236, 401.	0.5	12
40	Evidence for mega-landslides as drivers of island colonization. <i>Journal of Biogeography</i> , 2017, 44, 1053-1064.	3.0	20
41	Connecting Earth observation to high-throughput biodiversity data. <i>Nature Ecology and Evolution</i> , 2017, 1, 176.	7.8	156
42	Assessing the potential of RAD-sequencing to resolve phylogenetic relationships within species radiations: The fly genus <i>Chiastocheta</i> (Diptera: Anthomyiidae) as a case study. <i>Molecular Phylogenetics and Evolution</i> , 2017, 114, 189-198.	2.7	18
43	A roadmap for island biology: 50 fundamental questions after 50 years of <i>The Theory of Island Biogeography</i> . <i>Journal of Biogeography</i> , 2017, 44, 963-983.	3.0	167
44	A combined field survey and molecular identification protocol for comparing forest arthropod biodiversity across spatial scales. <i>Molecular Ecology Resources</i> , 2017, 17, 694-707.	4.8	30
45	Challenges, advances and perspectives in island biogeography. <i>Frontiers of Biogeography</i> , 2016, 8, .	1.8	5
46	Post-copulatory opportunities for sperm competition and cryptic female choice provide no offspring fitness benefits in externally fertilizing salmon. <i>Royal Society Open Science</i> , 2016, 3, 150709.	2.4	8
47	Evidence for multiple founding lineages and genetic admixture in the evolution of species within an oceanic island weevil (Coleoptera, Curculionidae) super-radiation. <i>Journal of Biogeography</i> , 2016, 43, 178-191.	3.0	16
48	Biodiversity in the Mexican highlands and the interaction of geology, geography and climate within the Trans-Mexican Volcanic Belt. <i>Journal of Biogeography</i> , 2015, 42, 1586-1600.	3.0	205
49	Model misspecification confounds the estimation of rates and exaggerates their time dependency. <i>Molecular Ecology</i> , 2015, 24, 6013-6020.	3.9	7
50	Lack of support for the time-dependent molecular evolution hypothesis. <i>Molecular Ecology</i> , 2015, 24, 702-709.	3.9	27
51	Host-associated genetic divergence and taxonomy in the <i>Rhinusa pilosa</i> species complex: an integrative approach. <i>Systematic Entomology</i> , 2015, 40, 268-287.	3.9	13
52	Islands as model systems in ecology and evolution: prospects fifty years after MacArthur & Wilson. <i>Ecology Letters</i> , 2015, 18, 200-217.	6.4	356
53	Sexual selection protects against extinction. <i>Nature</i> , 2015, 522, 470-473.	27.8	162
54	Gene Duplication, Population Genomics, and Species-Level Differentiation within a Tropical Mountain Shrub. <i>Genome Biology and Evolution</i> , 2014, 6, 2611-2624.	2.5	25

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55	Fission and fusion in island taxa - serendipity, or something to be expected?. <i>Molecular Ecology</i> , 2014, 23, 5132-5134.	3.9	9
56	Genetic, morphological, and dietary changes associated with novel habitat colonisation in the <i>Caryanys islandensis</i> grasshopper ( <i>Caryanys bellamyi</i> ). <i>Ecological Entomology</i> , 2014, 39, 703-715.	2.2	7
57	Experimental Removal of Sexual Selection Reveals Adaptations to Polyandry in Both Sexes. <i>Evolutionary Biology</i> , 2014, 41, 62-70.	1.1	28
58	Bulk De Novo Mitogenome Assembly from Pooled Total DNA Elucidates the Phylogeny of Weevils (Coleoptera: Curculionidae). <i>Molecular Biology and Evolution</i> , 2014, 31, 2223-2237.	8.9	195
59	Revision of <i>Mecynotarsus heydenii</i> species complex (Curculionidae): integrative taxonomy reveals multiple species exhibiting host specialization. <i>Zoologica Scripta</i> , 2014, 43, 34-51.	1.7	11
60	Quantifying surface area changes of volcanic islands driven by Pleistocene sea level cycles: biogeographical implications for the Macaronesian archipelagos. <i>Journal of Biogeography</i> , 2014, 41, 1242-1254.	3.0	73
61	Molecular characterization of trophic ecology within an island radiation of insect herbivores (Curculionidae: Entiminae: <i>Caryanys</i> ). <i>Molecular Ecology</i> , 2013, 22, 5441-5455.	3.9	32
62	Collembola, the biological species concept and the underestimation of global species richness. <i>Molecular Ecology</i> , 2013, 22, 5382-5396.	3.9	60
63	CRYPTIC CHOICE OF CONSPECIFIC SPERM CONTROLLED BY THE IMPACT OF OVARIAN FLUID ON SPERM SWIMMING BEHAVIOR. <i>Evolution; International Journal of Organic Evolution</i> , 2013, 67, 3523-3536.	2.3	92
64	Molecular phylogenetics of <i>Vanda</i> and related genera (Orchidaceae). <i>Botanical Journal of the Linnean Society</i> , 2013, 173, 549-572.	1.6	33
65	Genetic analysis of a contact zone between two lineages of the ocellated lizard ( <i>Lacerta</i> ). <i>Zoological Systematics and Evolutionary Research</i> , 2013, 51, 45-54.	1.4	21
66	A road map for molecular ecology. <i>Molecular Ecology</i> , 2013, 22, 2605-2626.	3.9	100
67	Molecular phylogenetic analysis of archival tissue reveals the origin of a disjunct southern African-Palaeartic weevil radiation. <i>Journal of Biogeography</i> , 2013, 40, 1348-1359.	3.0	11
68	Reliable, verifiable and efficient monitoring of biodiversity via metabarcoding. <i>Ecology Letters</i> , 2013, 16, 1245-1257.	6.4	514
69	Vulnerability to cavitation, hydraulic efficiency, growth and survival in an insular pine ( <i>Pinus</i> ). <i>Journal of Ecology</i> , 2013, 101, 1079-1089.	2.9	76
70	The Imprint of Geologic History on Within-Island Diversification of Woodlouse-Hunter Spiders (Araneae, Dysderidae) in the Canary Islands. <i>Journal of Heredity</i> , 2013, 104, 341-356.	2.4	27
71	PyroClean: Denoising Pyrosequences from Protein-Coding Amplicons for the Recovery of Interspecific and Intraspecific Genetic Variation. <i>PLoS ONE</i> , 2013, 8, e57615.	2.5	19
72	Phylogeographic Ancestral Inference Using the Coalescent Model on Haplotype Trees. <i>Journal of Computational Biology</i> , 2012, 19, 745-755.	1.6	13

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73	Age, origins and extinctions of the avifauna of Macaronesia: a synthesis of phylogenetic and fossil information. <i>Quaternary Science Reviews</i> , 2012, 50, 14-22.	3.0	58
74	Phylogeography and molecular phylogeny of Macaronesian island <i>Tarphius</i> (Coleoptera: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 70	3.0	22
75	Numts help to reconstruct the demographic history of the ocellated lizard ( <i>Lacerta lepida</i> ) in a secondary contact zone. <i>Molecular Ecology</i> , 2012, 21, 1005-1018.	3.9	26
76	Biodiversity soup: metabarcoding of arthropods for rapid biodiversity assessment and biomonitoring. <i>Methods in Ecology and Evolution</i> , 2012, 3, 613-623.	5.2	543
77	Phylogeny, phylogeography, phylobetadiversity and the molecular analysis of biological communities. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2011, 366, 2391-2402.	4.0	83
78	Discordant patterns of geographic variation between mitochondrial and microsatellite markers in the Mexican black iguana ( <i>Ctenosaura pectinata</i> ) in a contact zone. <i>Journal of Biogeography</i> , 2011, 38, 1394-1405.	3.0	25
79	Gene conversion rapidly generates major histocompatibility complex diversity in recently founded bird populations. <i>Molecular Ecology</i> , 2011, 20, 5213-5225.	3.9	84
80	EXPERIMENTAL EVOLUTION EXPOSES FEMALE AND MALE RESPONSES TO SEXUAL SELECTION AND CONFLICT IN <i>TRIBOLIUM CASTANEUM</i> . <i>Evolution; International Journal of Organic Evolution</i> , 2011, 65, 713-724.	2.3	76
81	Morphological, molecular and biological evidence reveal two cryptic species in <i>Mecinus janthinus</i> Germar (Coleoptera, Curculionidae), a successful biological control agent of Dalmatian toadflax, <i>Linaria dalmatica</i> (Lamiales, Plantaginaceae). <i>Systematic Entomology</i> , 2011, 36, 741-753.	3.9	46
82	Phylogeography and demographic history of <i>Lacerta lepida</i> in the Iberian Peninsula: multiple refugia, range expansions and secondary contact zones. <i>BMC Evolutionary Biology</i> , 2011, 11, 170.	3.2	62
83	A Bayesian approach to phylogeographic clustering. <i>Interface Focus</i> , 2011, 1, 909-921.	3.0	23
84	Inbreeding Promotes Female Promiscuity. <i>Science</i> , 2011, 333, 1739-1742.	12.6	84
85	Searching for Speciation Genes: Molecular Evidence for Selection Associated with Colour Morphotypes in the Caribbean Reef Fish Genus <i>Hypoplectrus</i> . <i>PLoS ONE</i> , 2011, 6, e20394.	2.5	13
86	Predominance of single paternity in the black spiny-tailed iguana: conservation genetic concerns for female-biased hunting. <i>Conservation Genetics</i> , 2010, 11, 1645-1652.	1.5	4
87	Distinguishing between hot-spots and melting-pots of genetic diversity using haplotype connectivity. <i>Algorithms for Molecular Biology</i> , 2010, 5, 19.	1.2	2
88	Host-associated genetic differentiation in a seed parasitic weevil <i>Rhinusa antirrhini</i> (Coleoptera: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 14 2286-2300.	3.9	35
89	Molecular phylogeny and Holarctic diversification of the subtribe Calathina (Coleoptera: Carabidae: Tj ETQq1 1 0.784314 rgBT /Overloc 2.7	2.7	10
90	Inbreeding depresses sperm competitiveness, but not fertilization or mating success in male <i>Tribolium castaneum</i> . <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2010, 277, 3483-3491.	2.6	62

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91	Combining contemporary and ancient DNA in population genetic and phylogeographical studies. <i>Molecular Ecology Resources</i> , 2010, 10, 760-772.	4.8	54
92	Evolution underground: shedding light on the diversification of subterranean insects. <i>Journal of Biology</i> , 2010, 9, 17.	2.7	29
93	Elevated substitution rate estimates from ancient DNA: model violation and bias of Bayesian methods. <i>Molecular Ecology</i> , 2009, 18, 4390-4397.	3.9	75
94	Isolation and characterization of polymorphic microsatellite markers in the black spiny tailed iguana ( <i>Ctenosaura pectinata</i> ) and their cross-utility in other <i>Ctenosaura</i> . <i>Molecular Ecology Resources</i> , 2009, 9, 117-119.	4.8	4
95	Genetic characterization, distribution and prevalence of avian pox and avian malaria in the Berthelot's pipit ( <i>Anthus berthelotii</i> ) in Macaronesia. <i>Parasitology Research</i> , 2008, 103, 1435-1443.	1.6	42
96	Phylogenetic relationships, biogeography and speciation in the avian genus <i>Saxicola</i> . <i>Molecular Phylogenetics and Evolution</i> , 2008, 48, 1145-1154.	2.7	46
97	Diversification in the northern neotropics: mitochondrial and nuclear DNA phylogeography of the iguana <i>Ctenosaura pectinata</i> and related species. <i>Molecular Ecology</i> , 2008, 17, 3259-3275.	3.9	71
98	Phylogenetic analysis of community assembly and structure over space and time. <i>Trends in Ecology and Evolution</i> , 2008, 23, 619-630.	8.7	559

99

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109	Characterization of polymorphic microsatellites in the castration parasite plant-ant <i>Allomerus octoarticulatus</i> cf. <i>demerarae</i> (Formicidae: Myrmicinae). <i>Molecular Ecology Notes</i> , 2006, 6, 182-184.	1.7	3
110	Characterization of microsatellite loci in <i>Brachyderes rugatus</i> , the Canary Islands pine weevil (Coleoptera: Curculionidae). <i>Molecular Ecology Notes</i> , 2006, 6, 820-822.	1.7	4
111	Chloroplast microsatellites reveal colonization and metapopulation dynamics in the Canary Island pine. <i>Molecular Ecology</i> , 2006, 15, 2691-2698.	3.9	55
112	Apparent <i>in situ</i> sympatric speciation in ecologically similar herbivorous beetles facilitated by multiple colonizations of an island. <i>Molecular Ecology</i> , 2006, 15, 2935-2947.	3.9	50
113	Testing phylogeographic predictions on an active volcanic island: <i>Brachyderes rugatus</i> (Coleoptera: Curculionidae). <i>Molecular Ecology</i> , 2006, 15, 2948-2957.	3.9	31
114	Species diversity can drive speciation. <i>Nature</i> , 2005, 434, 1015-1017.	27.8	271
115	Is speciation driven by species diversity? (Reply). <i>Nature</i> , 2005, 438, E2-E2.	27.8	12
116	DIVERSIFICATION OF THE FOREST BEETLE GENUS <i>TARPHIUS</i> ON THE CANARY ISLAND, AND THE EVOLUTIONARY ORIGINS OF ISLAND ENDEMICS. <i>Evolution; International Journal of Organic Evolution</i> , 2005, 59, 586-598.	2.3	64
117	Phylogeography. <i>Current Biology</i> , 2005, 15, R367-R371.	3.9	73
118	DIVERSIFICATION OF THE FOREST BEETLE GENUS <i>TARPHIUS</i> ON THE CANARY ISLANDS, AND THE EVOLUTIONARY ORIGINS OF ISLAND ENDEMICS. <i>Evolution; International Journal of Organic Evolution</i> , 2005, 59, 586.	2.3	67
119	Origin and Maintenance of a Broad-Spectrum Disease Resistance Locus in <i>Arabidopsis</i> . <i>Molecular Biology and Evolution</i> , 2004, 21, 1661-1672.	8.9	73
120	Molecular Phylogeny and Evolution of the Plant-Specific Seven-Transmembrane MLO Family. <i>Journal of Molecular Evolution</i> , 2003, 56, 77-88.	1.8	220
121	Revealing the demographic histories of species using DNA sequences. <i>Trends in Ecology and Evolution</i> , 2001, 16, 707-716.	8.7	182
122	The Diversification of the Genus <i>Nesotes</i> (Coleoptera: Tenebrionidae) in the Canary Islands: Evidence from mtDNA. <i>Molecular Phylogenetics and Evolution</i> , 2001, 21, 321-326.	2.7	35
123	COLONIZATION AND DIVERSIFICATION OF THE SPECIES <i>BRACHYDERES RUGATUS</i> (COLEOPTERA) ON THE CANARY ISLANDS: EVIDENCE FROM MITOCHONDRIAL DNA COII GENE SEQUENCES. <i>Evolution; International Journal of Organic Evolution</i> , 2000, 54, 911-923.	2.3	70
124	INTERPRETING COLONIZATION OF THE <i>CALATHUS</i> (COLEOPTERA: CARABIDAE) ON THE CANARY ISLANDS AND MADEIRA THROUGH THE APPLICATION OF THE PARAMETRIC BOOTSTRAP. <i>Evolution; International Journal of Organic Evolution</i> , 2000, 54, 2081-2090.	2.3	63
125	COLONIZATION AND DIVERSIFICATION OF THE SPECIES <i>BRACHYDERES RUGATUS</i> (COLEOPTERA) ON THE CANARY ISLANDS: EVIDENCE FROM MITOCHONDRIAL DNA COII GENE SEQUENCES. <i>Evolution; International Journal of Organic Evolution</i> , 2000, 54, 911.	2.3	3
126	INTERPRETING COLONIZATION OF THE <i>CALATHUS</i> (COLEOPTERA: CARABIDAE) ON THE CANARY ISLANDS AND MADEIRA THROUGH THE APPLICATION OF THE PARAMETRIC BOOTSTRAP. <i>Evolution; International Journal of Organic Evolution</i> , 2000, 54, 2081.	2.3	4



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127	Colonization and diversification: towards a phylogeographic synthesis for the Canary Islands. <i>Trends in Ecology and Evolution</i> , 2000, 15, 104-109.	8.7	363
128	Tracking colonization and diversification of insect lineages on islands: mitochondrial DNA phylogeography of <i>Tarphius canariensis</i> (Coleoptera: Colydiidae) on the Canary Islands. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2000, 267, 2199-2205.	2.6	48
129	MtDNA Phylogeography and Recent Intra-island Diversification among Canary Island <i>Calathus</i> Beetles. <i>Molecular Phylogenetics and Evolution</i> , 1999, 13, 149-158.	2.7	84
130	Biogeographic area relationships in southern New Zealand: a cladistic analysis of Lepidoptera distributions. <i>Journal of Biogeography</i> , 1997, 24, 89-99.	3.0	8
131	Phylogenetic Relationships of the <i>Prodontria</i> (Coleoptera; Scarabaeidae; Subfamily Melolonthinae), Derived from Sequence Variation in the Mitochondrial Cytochrome Oxidase II Gene. <i>Molecular Phylogenetics and Evolution</i> , 1995, 4, 433-447.	2.7	59
132	Conservation status of chafer beetles <i>Prodontria bicolorata</i> and <i>P. modesta</i> : distribution and ecological observations.. <i>New Zealand Entomologist</i> , 1994, 17, 3-6.	0.3	5
133	Speciation on islands: what are we learning?. <i>Biological Journal of the Linnean Society</i> , 0, 95, 47-52.	1.6	27