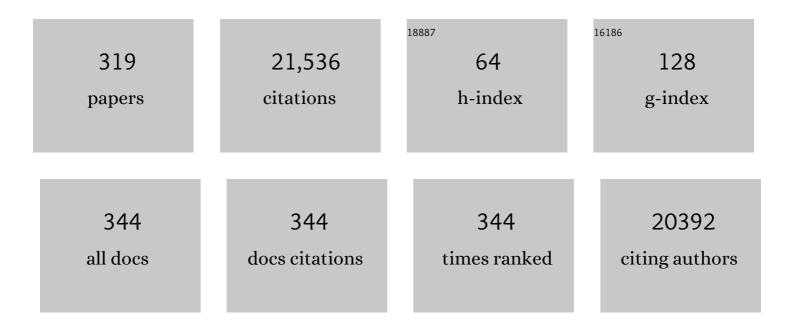
Roger K Butlin

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Ten years of demographic modelling of divergence and speciation in the sea. Evolutionary Applications, 2023, 16, 542-559.	1.5	11
2	An allozyme polymorphism is associated with a large chromosomal inversion in the marine snail <i>Littorina fabalis</i> . Evolutionary Applications, 2023, 16, 279-292.	1.5	7
3	Preface. Philosophical Transactions of the Royal Society B: Biological Sciences, 2022, 377, 20210491.	1.8	3
4	Introduction to the theme issue â€~Species' ranges in the face of changing environments'. Philosophical Transactions of the Royal Society B: Biological Sciences, 2022, 377, 20210002.	1.8	4
5	Experimental evolution of local adaptation under unidimensional and multidimensional selection. Current Biology, 2022, 32, 1310-1318.e4.	1.8	6
6	Parthenogenesis in <i>Darevskia</i> lizards: A rare outcome of common hybridization, not a common outcome of rare hybridization. Evolution; International Journal of Organic Evolution, 2022, 76, 899-914.	1.1	7
7	The rise and fall of an alien: why the successful colonizer Littorina saxatilis failed to invade the Mediterranean Sea. Biological Invasions, 2022, 24, 3169-3187.	1.2	39
8	Mutation accumulation opposes polymorphism: supergenes and the curious case of balanced lethals. Philosophical Transactions of the Royal Society B: Biological Sciences, 2022, 377, .	1.8	5
9	Inversions and parallel evolution. Philosophical Transactions of the Royal Society B: Biological Sciences, 2022, 377, .	1.8	19
10	Genomeâ€enabled discovery of candidate virulence loci in <i>Striga hermonthica</i> , a devastating parasite of African cereal crops. New Phytologist, 2022, 236, 622-638.	3.5	12
11	Longâ€ŧerm cloud forest response to climate warming revealed by insect speciation history*. Evolution; International Journal of Organic Evolution, 2021, 75, 231-244.	1.1	9
12	Deleterious mutation accumulation and the long-term fate of chromosomal inversions. PLoS Genetics, 2021, 17, e1009411.	1.5	71
13	Homage to Felsenstein 1981, or why are there so few/many species?. Evolution; International Journal of Organic Evolution, 2021, 75, 978-988.	1.1	13
14	Multisource noninvasive genetics of brown bears (Ursus arctos) in Greece reveals a highly structured population and a new matrilineal contact zone in southern Europe. Ecology and Evolution, 2021, 11, 6427-6443.	0.8	4
15	Genetic variation for adaptive traits is associated with polymorphic inversions in <i>Littorina saxatilis</i> . Evolution Letters, 2021, 5, 196-213.	1.6	42
16	Using replicate hybrid zones to understand the genomic basis of adaptive divergence. Molecular Ecology, 2021, 30, 3797-3814.	2.0	37
17	Multidimensional divergent selection, local adaptation, and speciation. Evolution; International Journal of Organic Evolution, 2021, 75, 2167-2178.	1.1	12
18	A large chromosomal inversion shapes gene expression in seaweed flies (<i>Coelopa frigida</i>). Evolution Letters, 2021, 5, 607-624.	1.6	11

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19	ls embryo abortion a postâ€zygotic barrier to gene flow between <i>Littorina</i> ecotypes?. Journal of Evolutionary Biology, 2020, 33, 342-351.	0.8	14
20	Is it time to abandon the biological species concept? No. National Science Review, 2020, 7, 1400-1401.	4.6	14
21	A developmentally descriptive method for quantifying shape in gastropod shells. Journal of the Royal Society Interface, 2020, 17, 20190721.	1.5	15
22	The evolution of strong reproductive isolation between sympatric intertidal snails. Philosophical Transactions of the Royal Society B: Biological Sciences, 2020, 375, 20190545.	1.8	23
23	Towards the completion of speciation: the evolution of reproductive isolation beyond the first barriers. Philosophical Transactions of the Royal Society B: Biological Sciences, 2020, 375, 20190528.	1.8	75
24	Instability of natural selection at candidate barrier loci underlying speciation in wood ants. Molecular Ecology, 2020, 29, 3988-3999.	2.0	13
25	Assortative mating, sexual selection, and their consequences for gene flow in <i>Littorina</i> . Evolution; International Journal of Organic Evolution, 2020, 74, 1482-1497.	1.1	23
26	Secondary contact zones of closelyâ€related <i>Erebia</i> butterflies overlap with narrow phenotypic and parasitic clines. Journal of Evolutionary Biology, 2020, 33, 1152-1163.	0.8	17
27	Transcriptomic resources for evolutionary studies in flat periwinkles and related species. Scientific Data, 2020, 7, 73.	2.4	1
28	Hybridization patterns between two marine snails, <i>Littorina fabalis</i> and <i>L. obtusata</i> . Ecology and Evolution, 2020, 10, 1158-1179.	0.8	15
29	The role of hybridisation in the origin and evolutionary persistence of vertebrate parthenogens: a case study of Darevskia lizards. Heredity, 2019, 123, 795-808.	1.2	13
30	Ecological speciation in sympatric palms: 3. Genetic map reveals genomic islands underlying species divergence in <i>Howea</i> . Evolution; International Journal of Organic Evolution, 2019, 73, 1986-1995.	1.1	13
31	Ecological speciation in sympatric palms: 4. Demographic analyses support speciation of Howea in the face of high gene flow. Evolution; International Journal of Organic Evolution, 2019, 73, 1996-2002.	1.1	14
32	Local adaptation stops where ecological gradients steepen or are interrupted. Evolutionary Applications, 2019, 12, 1449-1462.	1.5	31
33	Evolving Inversions. Trends in Ecology and Evolution, 2019, 34, 239-248.	4.2	179
34	Science policies: How should science funding be allocated? An evolutionary biologists' perspective. Journal of Evolutionary Biology, 2019, 32, 754-768.	0.8	16
35	Genomic architecture of parallel ecological divergence: Beyond a single environmental contrast. Science Advances, 2019, 5, eaav9963.	4.7	92
36	Testing an hypothesis of hybrid zone movement for toads in France. Molecular Ecology, 2019, 28, 1070-1083.	2.0	31

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37	Phylogenomics of the adaptive radiation of Triturus newts supports gradual ecological niche expansion towards an incrementally aquatic lifestyle. Molecular Phylogenetics and Evolution, 2019, 133, 120-127.	1.2	38
38	Multiple chromosomal rearrangements in a hybrid zone between <i>Littorina saxatilis</i> ecotypes. Molecular Ecology, 2019, 28, 1375-1393.	2.0	103
39	Sexes and species as rival units of niche saturation during community assembly. Global Ecology and Biogeography, 2018, 27, 593-603.	2.7	20
40	Coupling, Reinforcement, and Speciation. American Naturalist, 2018, 191, 155-172.	1.0	155
41	Conflict between heterozygote advantage and hybrid incompatibility in haplodiploids (and sex) Tj ETQq1 1 0.78	4314 rgB1 2.0	Öyerlock 1
42	Are assortative mating and genital divergence driven by reinforcement?. Evolution Letters, 2018, 2, 557-566.	1.6	16
43	Clines on the seashore: The genomic architecture underlying rapid divergence in the face of gene flow. Evolution Letters, 2018, 2, 297-309.	1.6	103
44	Inter and Intraspecific Genomic Divergence in Drosophila montana Shows Evidence for Cold Adaptation. Genome Biology and Evolution, 2018, 10, 2086-2101.	1.1	25
45	Widespread hybridization within moundâ€building wood ants in Southern Finland results in cytonuclear mismatches and potential for sexâ€specific hybrid breakdown. Molecular Ecology, 2017, 26, 4013-4026.	2.0	23
46	A genomic footprint of hybrid zone movement in crested newts. Evolution Letters, 2017, 1, 93-101.	1.6	77
47	What explains rare and conspicuous colours in a snail? A test of time-series data against models of drift, migration or selection. Heredity, 2017, 118, 21-30.	1.2	33
48	The chemical signatures underlying host plant discrimination by aphids. Scientific Reports, 2017, 7, 8498.	1.6	27
49	Interpreting the genomic landscape of speciation: a road map for finding barriers to gene flow. Journal of Evolutionary Biology, 2017, 30, 1450-1477.	0.8	399
50	A signature of dynamic biogeography: enclaves indicate past species replacement. Proceedings of the Royal Society B: Biological Sciences, 2017, 284, 20172014.	1.2	36
51	Genetic variation, selection and evolution: special issue in celebration of the 50th anniversary of the population genetics group meeting. Heredity, 2017, 118, 1-1.	1.2	5
52	Local adaptation of reproductive performance during thermal stress. Journal of Evolutionary Biology, 2017, 30, 422-429.	0.8	76
53	Targeted reâ€sequencing confirms the importance of chemosensory genes in aphid host race differentiation. Molecular Ecology, 2017, 26, 43-58.	2.0	27
54	Mechanisms of Adaptive Divergence and Speciation in Littorina saxatilis: Integrating Knowledge from Ecology and Genetics with New Data Emerging from Genomic Studies. Population Genomics, 2017, , 277-301.	0.2	20

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55	Adaptation to dislodgement risk on wave-swept rocky shores in the snail Littorina saxatilis. PLoS ONE, 2017, 12, e0186901.	1.1	34
56	Inhibition of plasmin-mediated TAFI activation may affect development but not progression of abdominal aortic aneurysms. PLoS ONE, 2017, 12, e0177117.	1.1	4
57	Targeted resequencing reveals geographical patterns of differentiation for loci implicated in parallel evolution. Molecular Ecology, 2016, 25, 3169-3186.	2.0	27
58	A Linkage Map and QTL Analysis for Pyrethroid Resistance in the Bed Bug <i>Cimex lectularius</i> . G3: Genes, Genomes, Genetics, 2016, 6, 4059-4066.	0.8	9
59	Gene expression clines reveal local adaptation and associated trade-offs at a continental scale. Scientific Reports, 2016, 6, 32975.	1.6	18
60	Shared and nonshared genomic divergence in parallel ecotypes of <i><scp>L</scp>ittorina saxatilis</i> at a local scale. Molecular Ecology, 2016, 25, 287-305.	2.0	142
61	Consequences of in-situ strategies for the conservation of plant genetic diversity. Biological Conservation, 2016, 203, 134-142.	1.9	41
62	Differential gene expression according to race and host plant in the pea aphid. Molecular Ecology, 2016, 25, 4197-4215.	2.0	59
63	Efficient screening for â€~genetic pollution' in an anthropogenic crested newt hybrid zone. Conservation Genetics Resources, 2016, 8, 553-560.	0.4	7
64	A universal mechanism generating clusters of differentiated loci during divergence-with-migration. Evolution; International Journal of Organic Evolution, 2016, 70, 1609-1621.	1.1	29
65	Ecological speciation in sympatric palms: 1. Gene expression, selection and pleiotropy. Journal of Evolutionary Biology, 2016, 29, 1472-1487.	0.8	29
66	Contrasting patterns of genetic diversity and spatial structure in an invasive symbiont-host association. Biological Invasions, 2016, 18, 3175-3191.	1.2	7
67	Tissue Culture as a Source of Replicates in Nonmodel Plants: Variation in Cold Response in <i>Arabidopsis lyrata</i> ssp. <i>petraea</i> . G3: Genes, Genomes, Genetics, 2016, 6, 3817-3823.	0.8	0
68	Ecological speciation in sympatric palms: 2. Pre―and postâ€zygotic isolation. Journal of Evolutionary Biology, 2016, 29, 2143-2156.	0.8	23
69	Deformed wing virus is a recent global epidemic in honeybees driven by <i>Varroa</i> mites. Science, 2016, 351, 594-597.	6.0	368
70	The genetic architecture of sexually selected traits in two natural populations of Drosophila montana. Heredity, 2015, 115, 565-572.	1.2	7
71	Dynamics of Copy Number Variation in Host Races of the Pea Aphid. Molecular Biology and Evolution, 2015, 32, 63-80.	3.5	55
72	Selection on outlier loci and their association with adaptive phenotypes in <i>Littorina saxatilis</i> contact zones. Journal of Evolutionary Biology, 2015, 28, 328-337.	0.8	18

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73	What can aquatic gastropods tell us about phenotypic plasticity? A review and meta-analysis. Heredity, 2015, 115, 312-321.	1.2	43
74	Localization of quantitative trait loci for diapause and other photoperiodically regulated life history traits important in adaptation to seasonally varying environments. Molecular Ecology, 2015, 24, 2809-2819.	2.0	28
75	The environmental genomics of metazoan thermal adaptation. Heredity, 2015, 114, 502-514.	1.2	61
76	Outbreeding effects in an inbreeding insect, C imex lectularius. Ecology and Evolution, 2015, 5, 409-418.	0.8	12
77	Genomics and the origin of species. Nature Reviews Genetics, 2014, 15, 176-192.	7.7	850
78	PARALLEL EVOLUTION OF LOCAL ADAPTATION AND REPRODUCTIVE ISOLATION IN THE FACE OF GENE FLOW. Evolution; International Journal of Organic Evolution, 2014, 68, 935-949.	1.1	165
79	Advances in <scp>E</scp> cological <scp>S</scp> peciation: an integrative approach. Molecular Ecology, 2014, 23, 513-521.	2.0	63
80	Humanâ€facilitated metapopulation dynamics in an emerging pest species, <i><scp>C</scp>imex lectularius</i> . Molecular Ecology, 2014, 23, 1071-1084.	2.0	52
81	Do the same genes underlie parallel phenotypic divergence in different <i><scp>L</scp>ittorina saxatilis</i> populations?. Molecular Ecology, 2014, 23, 4603-4616.	2.0	73
82	Evaluation of genetic isolation within an island flora reveals unusually widespread local adaptation and supports sympatric speciation. Philosophical Transactions of the Royal Society B: Biological Sciences, 2014, 369, 20130342.	1.8	42
83	The genetics of insect mating systems. , 2014, , 59-77.		3
84	A systematic review of phenotypic responses to between-population outbreeding. Environmental Evidence, 2013, 2, 13.	1.1	38
85	Hybridization and speciation. Journal of Evolutionary Biology, 2013, 26, 229-246.	0.8	1,735
86	A comparative analysis of the mechanisms underlying speciation on Lord Howe Island. Journal of Evolutionary Biology, 2013, 26, 733-745.	0.8	28
87	A review of dengue as an emerging disease in Pakistan. Public Health, 2013, 127, 11-17.	1.4	108
88	Genital divergence in sympatric sister snails. Journal of Evolutionary Biology, 2013, 26, 210-215.	0.8	24
89	Pulling together or pulling apart: hybridization in theory and practice. Journal of Evolutionary Biology, 2013, 26, 294-298.	0.8	24
90	Population structure of the mosquito <i>Aedes aegypti</i> (<i>Stegomyia aegypti</i>) in Pakistan. Medical and Veterinary Entomology, 2013, 27, 430-440.	0.7	23

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91	Fewer invited talks by women in evolutionary biology symposia. Journal of Evolutionary Biology, 2013, 26, 2063-2069.	0.8	120
92	Godfrey M Hewitt (1940–2013): highlights in Heredity from a career in evolutionary genetics. Heredity, 2013, 110, 405-406.	1.2	1
93	Sexual Segregation and Flexible Mating Patterns in Temperate Bats. PLoS ONE, 2013, 8, e54194.	1.1	45
94	Transcriptome Characterisation of the Ant Formica exsecta with New Insights into the Evolution of Desaturase Genes in Social Hymenoptera. PLoS ONE, 2013, 8, e68200.	1.1	14
95	Two distinct genomic regions, harbouring the period and fruitless genes, affect male courtship song in Drosophila montana. Heredity, 2012, 108, 602-608.	1.2	9
96	Challenges and pitfalls in the characterization of anonymous outlier AFLP markers in non-model species: lessons from an ocellated lizard genome scan. Heredity, 2012, 109, 340-348.	1.2	12
97	The Littorina sequence database (LSD) – an online resource for genomic data. Molecular Ecology Resources, 2012, 12, 142-148.	2.2	15
98	Weak patriline effects are present in the cuticular hydrocarbon profiles of isolated <i><scp>F</scp>ormica exsecta</i> ants but they disappear in the colony environment. Ecology and Evolution, 2012, 2, 2333-2346.	0.8	12
99	Population genetics of the malaria vector Anopheles aconitus in China and Southeast Asia. Infection, Genetics and Evolution, 2012, 12, 1958-1967.	1.0	6
100	A Genome Scan and Linkage Disequilibrium Analysis among Chromosomal Races of the Australian Grasshopper Vandiemenella viatica. PLoS ONE, 2012, 7, e47549.	1.1	8
101	Habitat Choice and Speciation. International Journal of Ecology, 2012, 2012, 1-12.	0.3	27
102	What do we need to know about speciation?. Trends in Ecology and Evolution, 2012, 27, 27-39.	4.2	358
103	Sexual selection on song and cuticular hydrocarbons in two distinct populations of <i>Drosophila montana</i> . Ecology and Evolution, 2012, 2, 80-94.	0.8	37
104	Evidence for evolutionary change associated with the recent range expansion of the British butterfly, <i>Aricia agestis</i> , in response to climate change. Molecular Ecology, 2012, 21, 267-280.	2.0	58
105	Development of conserved microsatellite markers of high crossâ€species utility in bat species (Vespertilionidae, Chiroptera, Mammalia). Molecular Ecology Resources, 2012, 12, 532-548.	2.2	29
106	LARGE-SCALE CANDIDATE GENE SCAN REVEALS THE ROLE OF CHEMORECEPTOR GENES IN HOST PLANT SPECIALIZATION AND SPECIATION IN THE PEA APHID. Evolution; International Journal of Organic Evolution, 2012, 66, 2723-2738.	1.1	99
107	Habitat Association and Seasonality in a Mosaic and Bimodal Hybrid Zone between Chorthippus brunneus and C. jacobsi (Orthoptera: Acrididae). PLoS ONE, 2012, 7, e37684.	1.1	7
108	Speciation with gene flow on Lord Howe Island. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 13188-13193.	3.3	184

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109	Taxonâ€specific PCR for DNA barcoding arthropod prey in bat faeces. Molecular Ecology Resources, 2011, 11, 236-244.	2.2	421
110	Chromosomal Speciation Revisited: Modes of Diversification in Australian Morabine Grasshoppers (Vandiemenella, viatica Species Group). Insects, 2011, 2, 49-61.	1.0	24
111	Association of Mc1r variants with ecologically relevant phenotypes in the European ocellated lizard, Lacerta lepida. Journal of Evolutionary Biology, 2011, 24, 2289-2298.	0.8	27
112	Multiple approaches to detect outliers in a genome scan for selection in ocellated lizards (Lacerta) Tj ETQq0 0 0 r	gBT /Overl 2.0	ock 10 Tf 50
113	Comparative phylogeography reveals a shared impact of pleistocene environmental change in shaping genetic diversity within nine Anopheles mosquito species across the Indo-Burma biodiversity hotspot. Molecular Ecology, 2011, 20, 4533-4549.	2.0	61
114	A framework for comparing processes of speciation in the presence of gene flow. Molecular Ecology, 2011, 20, 5123-5140.	2.0	287
115	Few parasites, and no evidence for Wolbachia infections, in a freshwater ostracod inhabiting temporary ponds. Biological Journal of the Linnean Society, 2011, 102, 208-216.	0.7	12
116	Kin assortment in juvenile shoals in wild guppy populations. Heredity, 2011, 106, 749-756.	1.2	44
117	Intron sequences of arginine kinase in an intertidal snail suggest an ecotype-specific selective sweep and a gene duplication. Heredity, 2011, 106, 808-816.	1.2	20
118	Data archiving. Heredity, 2011, 106, 709-709.	1.2	2
119	Mitochondrial DNA variation in the malaria vector Anopheles minimus across China, Thailand and Vietnam: evolutionary hypothesis, population structure and population history. Heredity, 2011, 106, 241-252.	1.2	33
120	EVALUATION OF ELEVATED PLOIDY AND ASEXUAL REPRODUCTION AS ALTERNATIVE EXPLANATIONS FOR GEOGRAPHIC PARTHENOGENESIS IN EUCYPRIS VIRENS OSTRACODS. Evolution; International Journal of Organic Evolution, 2010, 64, 986-997.	1.1	61
121	Reduction in post-invasion genetic diversity in Crangonyx pseudogracilis (Amphipoda: Crustacea): a genetic bottleneck or the work of hitchhiking vertically transmitted microparasites?. Biological Invasions, 2010, 12, 191-209.	1.2	43
122	Population genomics and speciation. Genetica, 2010, 138, 409-418.	0.5	98
123	The adaptive value of phenotypic plasticity in two ecotypes of a marine gastropod. BMC Evolutionary Biology, 2010, 10, 333.	3.2	54
124	Gems from the Heredity Archive. Heredity, 2010, 104, 121-121.	1.2	0
125	Exceptional cryptic diversity and multiple origins of parthenogenesis in a freshwater ostracod. Molecular Phylogenetics and Evolution, 2010, 54, 542-552.	1.2	114

126Why is adaptation prevented at ecological margins? New insights from individualâ€based simulations.
Ecology Letters, 2010, 13, 485-494.3.0119

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127	An ESTâ€based genome scan using 454 sequencing in the marine snail <i>Littorina saxatilis</i> . Journal of Evolutionary Biology, 2010, 23, 2004-2016.	0.8	71
128	Repeated evolution of reproductive isolation in a marine snail: unveiling mechanisms of speciation. Philosophical Transactions of the Royal Society B: Biological Sciences, 2010, 365, 1735-1747.	1.8	151
129	<i>Myotis alcathoe</i> Confirmed in the UK from Mitochondrial and Microsatellite DNA. Acta Chiropterologica, 2010, 12, 471-483.	0.2	14
130	Wind-borne insects mediate directional pollen transfer between desert fig trees 160 kilometers apart. Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 20342-20347.	3.3	178
131	Variation at range margins across multiple spatial scales: environmental temperature, population genetics and metabolomic phenotype. Proceedings of the Royal Society B: Biological Sciences, 2009, 276, 1495-1506.	1.2	52
132	Large Gene Family Expansions and Adaptive Evolution for Odorant and Gustatory Receptors in the Pea Aphid, Acyrthosiphon pisum. Molecular Biology and Evolution, 2009, 26, 2073-2086.	3.5	176
133	Social group size affects <i>Mycobacterium bovis</i> infection in European badgers (<i>Meles) Tj ETQq1 1 0.78</i>	4314 rgBT 1.3	/Overlock 10 41
134	Reâ€examination of a proposed case of stasipatric speciation: phylogeography of the Australian morabine grasshoppers (<i>Vandiemenella viatica</i> species group). Molecular Ecology, 2009, 18, 3429-3442.	2.0	23
135	How sympatric is speciation in the <i>Howea</i> palms of Lord Howe Island?. Molecular Ecology, 2009, 18, 3629-3638.	2.0	33
136	On the scent of speciation: the chemosensory system and its role in premating isolation. Heredity, 2009, 102, 77-97.	1.2	380
137	Genetics of speciation. Heredity, 2009, 102, 1-3.	1.2	21
138	GENETIC ANALYSIS OF A CHROMOSOMAL HYBRID ZONE IN THE AUSTRALIAN MORABINE GRASSHOPPERS (<i>VANDIEMENELLA</i> , <i>VIATICA</i> SPECIES GROUP). Evolution; International Journal of Organic Evolution, 2009, 63, 139-152.	1.1	41
139	Sex-specific roost movements and population dynamics of the vulnerable long-fingered bat, Myotis capaccinii. Biological Conservation, 2009, 142, 280-289.	1.9	26
140	Sites of evolutionary divergence differ between olfactory and gustatory receptors of Drosophila. Biology Letters, 2009, 5, 244-247.	1.0	15
141	Genetic population structure and introgression in Anopheles dirus mosquitoes in South-east Asia. Molecular Ecology, 2008, 10, 569-580.	2.0	53
142	Forest-obligate Sabethes mosquitoes suggest palaeoecological perturbations. Heredity, 2008, 101, 186-195.	1.2	9
143	<i>Drosophila</i> chemoreceptor gene evolution: selection, specialization and genome size. Molecular Ecology, 2008, 17, 1648-1657.	2.0	109
144	Sequence differentiation in regions identified by a genome scan for local adaptation. Molecular Ecology, 2008, 17, 3123-3135.	2.0	115

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145	Hitching a lift on the road to speciation. Molecular Ecology, 2008, 17, 4177-4180.	2.0	36
146	Signals of demographic expansion in Drosophila virilis. BMC Evolutionary Biology, 2008, 8, 59.	3.2	32
147	Identification of bat species in Greece from their echolocation calls. Acta Chiropterologica, 2008, 10, 127-143.	0.2	58
148	Seasonal Roosting Habits and Population Structure of the Long-fingered Bat Myotis capaccinii in Greece. Journal of Mammalogy, 2008, 89, 503-512.	0.6	24
149	An objective, rapid and reproducible method for scoring AFLP peakâ€height data that minimizes genotyping error. Molecular Ecology Resources, 2008, 8, 725-735.	2.2	155
150	Multiplex SNP‧CALE: a costâ€effective mediumâ€ŧhroughput single nucleotide polymorphism genotyping method. Molecular Ecology Resources, 2008, 8, 1230-1238.	2.2	65
151	Sympatric, parapatric or allopatric: the most important way to classify speciation?. Philosophical Transactions of the Royal Society B: Biological Sciences, 2008, 363, 2997-3007.	1.8	283
152	Evolution of a Complex Locus: Exon Gain, Loss and Divergence at the Gr39a Locus in Drosophila. PLoS ONE, 2008, 3, e1513.	1.1	17
153	Landscape and Land Cover Factors Influence the Presence of <i>Aedes</i> and <i>Anopheles</i> Larvae. Journal of Medical Entomology, 2007, 44, 133-144.	0.9	53
154	Landscape and Land Cover Factors Influence the Presence of <1>Aedes 1 and <1>Anopheles 1 Larvae. Journal of Medical Entomology, 2007, 44, 133-144.	0.9	39
155	Genetic diversity and molecular identification of mosquito species in the Anopheles maculatus group using the ITS2 region of rDNA. Infection, Genetics and Evolution, 2007, 7, 93-102.	1.0	105
156	Evolution of genes and genomes on the Drosophila phylogeny. Nature, 2007, 450, 203-218.	13.7	1,886
157	The extent of variation in male song, wing and genital characters among allopatric Drosophila montana populations. Journal of Evolutionary Biology, 2007, 20, 1591-1601.	0.8	35
158	Molecular identification of mosquito species in the Anopheles annularis group in southern Asia. Medical and Veterinary Entomology, 2007, 21, 30-35.	0.7	25
159	Genetic evidence that culling increases badger movement: implications for the spread of bovine tuberculosis. Molecular Ecology, 2007, 16, 4919-4929.	2.0	59
160	Differential gene flow of mitochondrial and nuclear DNA markers among chromosomal races of Australian morabine grasshoppers (Vandiemenella, viatica species group). Molecular Ecology, 2007, 16, 5044-5056.	2.0	27
161	Polymorphic microsatellite markers for chromosomal races of Australian morabine grasshoppers (Vandiemenella, viatica species group). Molecular Ecology Notes, 2007, 7, 1181-1184.	1.7	4
162	The attractiveness fragment—AFLP analysis of local adaptation and sexual selection in a caeliferan grasshopper, Chorthippus biguttulus. Die Naturwissenschaften, 2007, 94, 667-674.	0.6	9

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163	Impact of Land-use Change on Dengue and Malaria in Northern Thailand. EcoHealth, 2007, 4, 37-51.	0.9	84
164	Autumn swarming behaviour of Natterer's bats in the UK: Population size, catchment area and dispersal. Biological Conservation, 2006, 127, 215-226.	1.9	99
165	Post-modern synthesis. Trends in Ecology and Evolution, 2006, 21, 536-536.	4.2	2
166	A molecular approach to detect hybridization between bream Abramis brama, roach Rutlius rutilus and rudd Scardinius erythrophthalmus. Journal of Fish Biology, 2006, 69, 52-71.	0.7	40
167	ADAPTATION TO A STEEP ENVIRONMENTAL GRADIENT AND AN ASSOCIATED BARRIER TO GENE EXCHANGE IN LITTORINA SAXATILIS. Evolution; International Journal of Organic Evolution, 2006, 60, 268-278.	1.1	134
168	Assortative preferences and discrimination by females against hybrid male song in the grasshoppers Chorthippus brunneus and Chorthippus jacobsi (Orthoptera: Acrididae). Journal of Evolutionary Biology, 2006, 19, 1248-1256.	0.8	57
169	Molecular variation, systematics and distribution of the Anopheles fluviatilis complex in southern Asia. Medical and Veterinary Entomology, 2006, 20, 33-43.	0.7	49
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