Hamish Gordon Spencer

List of Publications by Year
in descending order

151	8,147	76326	51608
papers	citations	40 h-index	86 g-index

162	162	162	8766
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Rapid radiation of Southern Ocean shags in response to receding sea ice. Journal of Biogeography, 2022, 49, 942-953.	3.0	3
2	Taxonomic consistency and nomenclatural rules within oysters: Comment on Li et al. (2021). Molecular Phylogenetics and Evolution, 2022, 170, 107437.	2.7	4
3	Epigenetic induction may speed up or slow down speciation with gene flow. Evolution; International Journal of Organic Evolution, 2022, 76, 1170-1182.	2.3	8
4	Seven snail species hidden in one: Biogeographic diversity in an apparently widespread periwinkle in the Southern Ocean. Journal of Biogeography, 2022, 49, 1521-1534.	3.0	9
5	Graph-structured populations and the Hillâ€“Robertson effect. Royal Society Open Science, 2021, 8, 201831.	2.4	1
6	Phylogeography of the intertidal marine bivalve Lasaea hinemoa (Mollusca: Bivalvia) in New Zealand. Molluscan Research, 2021, 41, 191-203.	0.7	0
7	A model of optimal timing for a predictive adaptive response. Journal of Developmental Origins of Health and Disease, 2021, , 1-7.	1.4	4
8	Avoiding extinction under nonlinear environmental change: models of evolutionary rescue with plasticity. Biology Letters, 2021, 17, 20210459.	2.3	4
9	Water stratification in the marine biome restricts vertical environmental DNA (eDNA) signal dispersal. Environmental DNA, 2020, 2, 99-111.	5.8	74
10	Beyond Equilibria: The Neglected Role of History in Ecology and Evolution. Quarterly Review of Biology, 2020, 95, 311-321.	0.1	6
11	Killing the Behavioral Zombie: Genes, Evolution, and Why Behavior Isnâ€™t Special. BioScience, 2020, 70, 515-520.	4.9	6
12	Population structure of the New Zealand whelk, Cominella glandiformis (Gastropoda: Buccinidae), suggests sporadic dispersal of a direct developer. Biological Journal of the Linnean Society, 2020, 130, 49-60.	1.6	2
13	Catalogue of New Zealand land, freshwater and estuarine molluscan taxa named by Frederick Wollaston Hutton between 1879 and 1904 . Zootaxa, 2020, 4865, 1-73.	0.5	0
14	A further perspective on speciation by reinforcement. Theoretical Biology Forum, 2020, 113, 63-66.	0.2	1
15	Archival DNA reveals cryptic biodiversity within the Spotted Shag (Phalacrocorax punctatus) from New Zealand. Condor, 2019, 121, .	1.6	3
16	Sorting out the Snakebirds: The species status, phylogeny, and biogeography of the Darters (Aves:) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50	1.4	6
17	Speciesâ€level biodiversity assessment using marine environmental DNA metabarcoding requires protocol optimization and standardization. Ecology and Evolution, 2019, 9, 1323-1335.	1.9	62
18	Environmental DNA (eDNA) metabarcoding reveals strong discrimination among diverse marine habitats connected by water movement. Molecular Ecology Resources, 2019, 19, 426-438.	4.8	180

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19	Systematic revision of Nacella (Patellogastropoda: Nacellidae) based on a complete phylogeny of the genus, with the description of a new species from the southern tip of South America. Zoological Journal of the Linnean Society, 2019, 186, 303-336.	2.3	12
20	The phylogenetic placement of the enigmatic Indian Cormorant, Phalacrocorax fuscicollis (Phalacrocoracidae). Molecular Phylogenetics and Evolution, 2019, 130, 227-232.	2.7	3
21	Unexpected absence of island endemics: Longâ€distance dispersal in higher latitude subâ€Antarctic Siphonaria (Gastropoda: Euthyneura) species. Journal of Biogeography, 2018, 45, 874-884.	3.0	34
22	The evolution of epigenetically mediated adaptive transgenerational plasticity in a subdivided population. Evolution; International Journal of Organic Evolution, 2018, 72, 2773-2780.	2.3	18
23	Population-genetic models of sex-limited genomic imprinting. Theoretical Population Biology, 2017, 115, 35-44.	1.1	0
24	Case 3706 â€" Trochus (Osilinus ?) Capillaceus Philippi, 1849 (currently Cantharidus capillaceus;) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 Zoological Nomenclature, 2017, 74, 8.	0.1	1
25	Following the Antarctic Circumpolar Current: patterns and processes in the biogeography of the limpet Nacella (Mollusca: Patellogastropoda) across the Southern Ocean. Journal of Biogeography, 2017, 44, 861-874.	3.0	41
26	Valve microstructure and phylomineralogy of New Zealand chitons. Journal of Structural Biology, 2017, 197, 250-259.	2.8	8
27	Speciation, range contraction and extinction in the endemic New Zealand King Shag complex. Molecular Phylogenetics and Evolution, 2017, 115, 197-209.	2.7	14
28	The Selective Maintenance of Allelic Variation Under Generalized Dominance. G3: Genes, Genomes, Genetics, 2016, 6, 3725-3732.	1.8	2
29	Genetic and morphological evidence for two species ofLeucocarboshag (Aves, Pelecaniformes,) Tj ETQq1 1 0.784314 rgBT /Ov Society, 2016, 177, 676-694.	2.3	17
30	Host and ecology both play a role in shaping distribution of digenean parasites of New Zealand whelks (Gastropoda: Buccinidae: Cominella). Parasitology, 2016, 143, 1143-1156.	1.5	7
31	Phylogeographic patterns in New Zealand and temperate Australian cantharidines (Mollusca:) Tj ETQq1 1 0.784314 rgBT /Overlock 1 Phylogenetics and Evolution, 2016, 100, 333-344.	2.7	6
32	The Evolution of Sex-Specific Dominance in Response to Sexually Antagonistic Selection. American Naturalist, 2016, 187, 658-666.	2.1	37
33	For Host's Sake: The Pluses of Parasite Preservation. Trends in Ecology and Evolution, 2016, 31, 341-343.	8.7	33
34	Skeletal mineralogy of scaphopods: an unusual uniformity. Journal of Molluscan Studies, 2016, 82, 344-348.	1.2	4
35	Trans-Tasman genetic connectivity in the intertidal air-breathing slug Onchidella nigricans. Marine Ecology - Progress Series, 2016, 562, 93-100.	1.9	8
36	The Maintenance of Single-Locus Polymorphism by Maternal Selection. G3: Genes, Genomes, Genetics, 2015, 5, 963-969.	1.8	2

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37	New Zealand screw shellsMaoricolpus roseus(Gastropoda: Turritellidae): two species, two subspecies or a single variable species?. Molluscan Research, 2015, 35, 123-127.	0.7	2
38	Phylogeography of the whelk genusCominella(Gastropoda: Buccinidae) suggests long-distance counter-current dispersal of a direct developer. Biological Journal of the Linnean Society, 2015, 115, 315-332.	1.6	17
39	Non-conflict theories for the evolution of genomic imprinting. Heredity, 2014, 113, 112-118.	2.6	56
40	HOW STABLE â€˜SHOULDâ€™ EPIGENETIC MODIFICATIONS BE? INSIGHTS FROM ADAPTIVE PLASTICITY AND BET HEDGING. Evolution; International Journal of Organic Evolution, 2014, 68, 632-643.	2.3	164
41	Classification of the cormorants of the world. Molecular Phylogenetics and Evolution, 2014, 79, 249-257.	2.7	45
42	Transoceanic genetic similarities of kelpâ€associated sea slug populations: longâ€distance dispersal via rafting?. Journal of Biogeography, 2014, 41, 2357-2370.	3.0	56
43	Genomic imprinting: theories and data. Heredity, 2014, 113, 93-95.	2.6	2
44	Strong Phylogeographic Structure in a Sedentary Seabird, the Stewart Island Shag (Leucocarbo) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 46	2.5	15
45	The evolutionary potential of paramutation: A population-epigenetic model. Theoretical Population Biology, 2013, 88, 9-19.	1.1	16
46	The Two Faces of Robert FitzRoy, Captain of HMS Beagle and Governor of New Zealand. Quarterly Review of Biology, 2013, 88, 219-225.	0.1	2
47	Exploring epiallele stability in a population-epigenetic model. Theoretical Population Biology, 2013, 83, 136-144.	1.1	33
48	Biogeography Off the Tracks. Systematic Biology, 2013, 62, 494-498.	5.6	35
49	Comments on some taxonomic changes affecting marine Bivalvia of the New Zealand region recently introduced in Huber'sCompendium of bivalves, with some additional taxonomic changes. Molluscan Research, 2013, 33, 40-49.	0.7	4
50	The adaptive invasion of epialleles in a heterogeneous environment. Theoretical Population Biology, 2013, 88, 1-8.	1.1	19
51	The phylogenetic relationships of the extant pelicans inferred from DNA sequence data. Molecular Phylogenetics and Evolution, 2013, 66, 215-222.	2.7	19
52	Effects of Genetic Drift and Gene Flow on the Selective Maintenance of Genetic Variation. Genetics, 2013, 194, 235-244.	2.9	28
53	Passive rafting is a powerful driver of transoceanic gene flow. Biology Letters, 2013, 9, 20120821.	2.3	55
54	Models of Frequency-Dependent Selection with Mutation from Parental Alleles. Genetics, 2013, 195, 231-242.	2.9	5

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55	Genetic Affinities between Trans-Oceanic Populations of Non-Buoyant Macroalgae in the High Latitudes of the Southern Hemisphere. PLoS ONE, 2013, 8, e69138.	2.5	63
56	Structured Populations and the Maintenance of Sex. Lecture Notes in Computer Science, 2013, , 56-67.	1.3	1
57	Genomic Imprinting Leads to Less Selectively Maintained Polymorphism on X Chromosomes. Genetics, 2012, 192, 1455-1464.	2.9	2
58	Durvillaea poha sp. nov. (Fucales, Phaeophyceae): a buoyant southern bull-kelp species endemic to New Zealand. Phycologia, 2012, 51, 151-156.	1.4	27
59	Phylogenetic relationships elucidate colonization patterns in the intertidal grazers Osilinus Philippi, 1847 and Phorcus Risso, 1826 (Gastropoda: Trochidae) in the northeastern Atlantic Ocean and Mediterranean Sea. Molecular Phylogenetics and Evolution, 2012, 62, 35-45.	2.7	42
60	Comparative and metaâ€analytic insights into life extension via dietary restriction. Aging Cell, 2012, 11, 401-409.	6.7	182
61	Population-epigenetic models of selection. Theoretical Population Biology, 2012, 81, 232-242.	1.1	87
62	Reef formation versus solitariness in two New Zealand serpulids does not involve cryptic species. Aquatic Biology, 2012, 16, 97-103.	1.4	7
63	Evolutionary consequences of microhabitat: population-genetic structuring in kelp- vs. rock-associated chitons. Molecular Ecology, 2011, 20, 4915-4924.	3.9	22
64	Contrasting population makeup of two intertidal gastropod species that differ in dispersal opportunities. Journal of Experimental Marine Biology and Ecology, 2011, 396, 224-232.	1.5	14
65	Comparison of populationâ€genetic structuring in congeneric kelpâ€versus rockâ€associated snails: a test of a dispersalâ€byâ€rafting hypothesis. Ecology and Evolution, 2011, 1, 169-180.	1.9	19
66	Quantitative Genetics of Genomic Imprinting: A Comparison of Simple Variance Derivations, the Effects of Inbreeding, and Response to Selection. G3: Genes, Genomes, Genetics, 2011, 1, 131-142.	1.8	12
67	Circumpolar dispersal by rafting in two subantarctic kelp-dwelling crustaceans. Marine Ecology - Progress Series, 2010, 405, 221-230.	1.9	161
68	Contemporary habitat discontinuity and historic glacial ice drive genetic divergence in Chilean kelp. BMC Evolutionary Biology, 2010, 10, 203.	3.2	121
69	Multigene phylogeny of the southern bull-kelp genus Durvillaea (Phaeophyceae: Fucales). Molecular Phylogenetics and Evolution, 2010, 57, 1301-1311.	2.7	45
70	Molecular systematics of the marine gastropod families Trochidae and Calliostomatidae (Mollusca:) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5	2.7	89
71	Identifying Cliques of Convergent Characters: Concerted Evolution in the Cormorants and Shags. Systematic Biology, 2010, 59, 433-445.	5.6	75
72	Kelp genes reveal effects of subantarctic sea ice during the Last Glacial Maximum. Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 3249-3253.	7.1	247

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73	Complex dynamics occur in a single-locus, multiallelic model of general frequency-dependent selection. Theoretical Population Biology, 2009, 76, 292-298.	1.1	6
74	Effects of genomic imprinting on quantitative traits. Genetica, 2009, 136, 285-293.	1.1	26
75	Glacial oceanographic contrasts explain phylogeography of Australian bull kelp. Molecular Ecology, 2009, 18, 2287-2296.	3.9	58
76	GENETIC AND MORPHOLOGICAL ANALYSES OF THE SOUTHERN BULL KELP DURVILLAEA ANTARCTICA (PHAEOPHYCEAE: DURVILLAEALES) IN NEW ZEALAND REVEAL CRYPTIC SPECIES¹. Journal of Phycology, 2009, 45, 436-443.	2.3	68
77	The phylogenetic position of the GalÃ¡pagos Cormorant. Molecular Phylogenetics and Evolution, 2009, 53, 94-98.	2.7	23
78	Systematics and phylogeny of a new cryptic species of Diloma Philippi (Mollusca: Gastropoda:) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 54 Invertebrate Systematics, 2009, 23, 19.	1.3	9
79	Developmental Origins of Health and Disease across Generations – Theory, Observation, Experiment. , 2009, , 52-64.		1
80	Genetic drift on networks: Ploidy and the time to fixation. Theoretical Population Biology, 2008, 74, 283-290.	1.1	12
81	Predictive adaptive responses in perspective. Trends in Endocrinology and Metabolism, 2008, 19, 109-110.	7.1	87
82	The Generation and Maintenance of Genetic Variation by Frequency-Dependent Selection: Constructing Polymorphisms Under the Pairwise Interaction Model. Genetics, 2008, 180, 1547-1557.	2.9	16
83	Evolution of Fitnesses in Structured Populations With Correlated Environments. Genetics, 2008, 179, 1469-1478.	2.9	7
84	An Asymmetric Model of Heterozygote Advantage at Major Histocompatibility Complex Genes: Degenerate Pathogen Recognition and Intersection Advantage. Genetics, 2008, 178, 1473-1489.	2.9	15
85	â€œIt's Ok, We're Not Cousins by Bloodâ€: The Cousin Marriage Controversy in Historical Perspective. PLoS Biology, 2008, 6, e320.	5.6	36
86	Frequency-Dependent Selection and the Maintenance of Genetic Variation: Exploring the Parameter Space of the Multiallelic Pairwise Interaction Model. Genetics, 2007, 176, 1729-1740.	2.9	35
87	Single-Locus Polymorphism in a Heterogeneous Two-Deme Model. Genetics, 2007, 176, 1625-1633.	2.9	18
88	Evolution of Fitnesses and Allele Frequencies in a Population With Spatially Heterogeneous Selection Pressures. Genetics, 2007, 177, 1743-1751.	2.9	15
89	Taxonomy and nomenclature of black nerites (Gastropoda:Neritimorpha:Nerita) from the South Pacific. Invertebrate Systematics, 2007, 21, 229.	1.3	32
90	Simultaneous polyphenism and cryptic species in an intertidal limpet from New Zealand. Molecular Phylogenetics and Evolution, 2007, 45, 470-479.	2.7	36

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91	Species assignation amongst morphologically cryptic larval Digenea isolated from New Zealand topshells (Gastropoda: Trochidae). Parasitology Research, 2007, 101, 433-441.	1.6	9
92	Response to Wells: Phenotypic responses to early environmental cues can be adaptive in adults. Trends in Ecology and Evolution, 2006, 21, 425-426.	8.7	11
93	Polymorphic microsatellite DNA markers in the mudflat topshell Diloma subrostrata (Gastropoda,) Tj ETQq1 1 0.784314 rgBT /Overlock	1.7	1
94	Phylogeography of Kauri Snails and their allies from Northland, New Zealand (Mollusca: Gastropoda:) Tj ETQq0 0 0 rgBT /Overlock 10 Tf	2.7	36
95	Population Models of Genomic Imprinting. II. Maternal and Fertility Selection. Genetics, 2006, 173, 2391-2398.	2.9	7
96	A Chip off the Old Block: A Model for the Evolution of Genomic Imprinting via Selection for Parental Similarity. Genetics, 2006, 174, 931-935.	2.9	12
97	Influence of Mom and Dad: Quantitative Genetic Models for Maternal Effects and Genomic Imprinting. Genetics, 2006, 173, 2297-2316.	2.9	26
98	The phylogeny and taxonomy of austral monodontine topshells (Mollusca: Gastropoda: Trochidae), inferred from DNA sequencesâ˜†. Molecular Phylogenetics and Evolution, 2005, 37, 474-483.	2.7	41
99	Adaptive dynamics, game theory and evolutionary population genetics. Journal of Evolutionary Biology, 2005, 18, 1191-1193.	1.7	17
100	Phylogeographical disjunction in abundant high-dispersal littoral gastropods. Molecular Ecology, 2005, 14, 2789-2802.	3.9	105
101	CLADOGENESIS AS THE RESULT OF LONGâ€DISTANCE RAFTING EVENTS IN SOUTH PACIFIC TOPSHELLS (GASTROPODA, TROCHIDAE). Evolution; International Journal of Organic Evolution, 2005, 59, 1701-1711.	2.3	87
102	A census of mammalian imprinting. Trends in Genetics, 2005, 21, 457-465.	6.7	612
103	Untangling Long Branches: Identifying Conflicting Phylogenetic Signals Using Spectral Analysis, Neighbor-Net, and Consensus Networks. Systematic Biology, 2005, 54, 620-633.	5.6	56
104	Environmental influences during development and their later consequences for health and disease: implications for the interpretation of empirical studies. Proceedings of the Royal Society B: Biological Sciences, 2005, 272, 671-677.	2.6	366
105	CLADOGENESIS AS THE RESULT OF LONG-DISTANCE RAFTING EVENTS IN SOUTH PACIFIC TOPSHELLS (GASTROPODA, TROCHIDAE). Evolution; International Journal of Organic Evolution, 2005, 59, 1701.	2.3	5
106	Predictive adaptive responses and human evolution. Trends in Ecology and Evolution, 2005, 20, 527-533.	8.7	582
107	Cladogenesis as the result of long-distance rafting events in South Pacific topshells (Gastropoda,) Tj ETQq1 1 0.784314 rgBT /Overlock	2.3	24
108	Frequency-Dependent Selection With Dominance: A Window Onto the Behavior of the Mean Fitness. Genetics, 2004, 167, 499-512.	2.9	15

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109	The Effect of Genetic Conflict on Genomic Imprinting and Modification of Expression at a Sex-Linked Locus. Genetics, 2004, 166, 565-579.	2.9	21
110	Developmental plasticity and human health. Nature, 2004, 430, 419-421.	27.8	1,529
111	Host specificity and molecular phylogeny of larval Digenea isolated from New Zealand and Australian topshells (Gastropoda: Trochidae). International Journal for Parasitology, 2004, 34, 557-568.	3.1	48
112	Phylogenies of the Frigatebirds (Fregatidae) and Tropicbirds (Phaethonidae), two divergent groups of the traditional order Pelecaniformes, inferred from mitochondrial DNA sequences. Molecular Phylogenetics and Evolution, 2004, 31, 31-38.	2.7	42
113	Further Properties of Gavriletsâ€™ One-Locus Two-Allele Model of Maternal Selection. Genetics, 2003, 164, 1689-1692.	2.9	11
114	The Evolution of Genomic Imprinting via Variance Minimization: An Evolutionary Genetic Model. Genetics, 2003, 165, 205-222.	2.9	9
115	Metapopulation Structure Favors Plasticity over Local Adaptation. American Naturalist, 2002, 160, 271-283.	2.1	553
116	The Correlation Between Relatives on the Supposition of Genomic Imprinting. Genetics, 2002, 161, 411-417.	2.9	51
117	Evolutionary Genetic Models of the Ovarian Time Bomb Hypothesis for the Evolution of Genomic Imprinting. Genetics, 2002, 162, 425-439.	2.9	27
118	Defending the group from the terror within. Metascience, 2001, 10, 192-202.	0.3	0
119	The Phylogenetic Relationships of the Shags and Cormorants: Can Sequence Data Resolve a Disagreement between Behavior and Morphology?. Molecular Phylogenetics and Evolution, 2000, 17, 345-359.	2.7	47
120	Genetic variation and prevalence of blood parasites do not correlate among bird species. Journal of Zoology, 2000, 252, 381-388.	1.7	14
121	Metazoan parasite species richness and genetic variation among freshwater fish species: cause or consequence?. International Journal for Parasitology, 2000, 30, 697-703.	3.1	24
122	Phylogeny, Biogeography, and Taxonomy of Australasian Teals. Auk, 2000, 117, 154-163.	1.4	22
123	POPULATION GENETICS AND EVOLUTION OF GENOMIC IMPRINTING. Annual Review of Genetics, 2000, 34, 457-477.	7.6	46
124	PHYLOGENY, BIOGEOGRAPHY, AND TAXONOMY OF AUSTRALASIAN TEALS. Auk, 2000, 117, 154.	1.4	25
125	The Long and Short of It: Branch Lengths and the Problem of Placing the New Zealand Short-Tailed Bat, Mystacina. Molecular Phylogenetics and Evolution, 1999, 13, 405-416.	2.7	42
126	Genetic conflicts and the evolutionary origin of genomic imprinting. Trends in Ecology and Evolution, 1999, 14, 197-201.	8.7	83

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127	Reply from H.G. Spencer, A.G. Clark and M.W. Feldman. Trends in Ecology and Evolution, 1999, 14, 359.	8.7	17
128	Population Models of Genomic Imprinting. I. Differential Viability in the Sexes and the Analogy With Genetic Dominance. Genetics, 1999, 153, 1949-1958.	2.9	15
129	Case 3095. Mystacina Gray, 1843, Chalinolobus Peters, 1866, M. tuberculata Gray, 1843 and Vespertilio tuberculatus J.R. Forster, 1844 (currently C. tuberculatus) (Mammalia, Chiroptera): proposed conservation of usage of the names. Bulletin of Zoological Nomenclature, 1999, 56, 250-254.	0.1	3
130	Onward and upward. Metascience, 1998, 7, 52-64.	0.3	0
131	Skeletal carbonate mineralogy of New Zealand bryozoans. Marine Geology, 1998, 151, 27-46.	2.1	63
132	Distribution of seabirds in coastal waters off Otago, New Zealand. New Zealand Journal of Marine and Freshwater Research, 1998, 32, 203-213.	2.0	5
133	The failure of a scientific critique: David Heron, Karl Pearson and Mendelian eugenics. British Journal for the History of Science, 1998, 31, 441-452.	0.7	14
134	Genetic Conflicts, Multiple Paternity and the Evolution of Genomic Imprinting. Genetics, 1998, 148, 893-904.	2.9	55
135	The Evolution of Genomic Imprinting: Two Modifier-Locus Models. Theoretical Population Biology, 1997, 51, 23-35.	1.1	34
136	Mutation-Selection Balance Under Genomic Imprinting at an Autosomal Locus. Genetics, 1997, 147, 281-287.	2.9	20
137	Hop, step and gape: do the social displays of the Pelecaniformes reflect phylogeny?. Animal Behaviour, 1996, 51, 273-291.	1.9	68
138	Perceptual constraints on optimal foraging: The effects of variation among foragers. Evolutionary Ecology, 1996, 10, 331-339.	1.2	26
139	Mutation-Selection Balance at a Modifier-of-Imprinting Locus. Genetics, 1996, 144, 361-367.	2.9	9
140	CYTOPLASMIC INCOMPATIBILITY: WHY IS IT SEXUALLY ASYMMETRIC?. Evolution; International Journal of Organic Evolution, 1995, 49, 1277-1280.	2.3	0
141	The hidden science of eugenics. Nature, 1995, 374, 302-304.	27.8	45
142	Cytoplasmic Incompatibility: Why is it Sexually Asymmetric?. Evolution; International Journal of Organic Evolution, 1995, 49, 1277.	2.3	0
143	Patch choice with competitive asymmetries and perceptual limits: the importance of history. Animal Behaviour, 1995, 50, 497-508.	1.9	31
144	Quantifying the Effect of Predation Risk on Foraging Bullies: No Need to Assume an IFD. Ecology, 1994, 75, 2220.	3.2	26

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145	The evolutionary construction of molecular polymorphisms. New Zealand Journal of Botany, 1993, 31, 249-256.	1.1	10
146	Assortative versus selective mating: Is the distinction worthwhile?. Biodemography and Social Biology, 1992, 39, 310-315.	1.0	2
147	Random genetic drift and selection in a triallelic locus: a continuous diffusion model. Mathematical Biosciences, 1992, 108, 127-139.	1.9	3
148	The Maintenance of Single-Locus Polymorphism. II. The Evolution of Fitnesses and Allele Frequencies. American Naturalist, 1991, 138, 1354-1371.	2.1	29
149	Measuring mating preferences: the use of Manly's beta. Heredity, 1988, 60, 305-310.	2.6	2
150	Reinforcement, Species, and Speciation: A Reply to Butlin. American Naturalist, 1987, 130, 958-962.	2.1	27
151	A Theoretical Investigation of Speciation by Reinforcement. American Naturalist, 1986, 128, 241-262.	2.1	124