Hamish Gordon Spencer

List of Publications by Year
in descending order

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

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

#	Article	IF	Citations
1	Developmental plasticity and human health. Nature, 2004, 430, 419-421.	27.8	1,529
2	A census of mammalian imprinting. Trends in Genetics, 2005, 21, 457-465.	6.7	612
3	Predictive adaptive responses and human evolution. Trends in Ecology and Evolution, 2005, 20, 527-533.	8.7	582
4	Metapopulation Structure Favors Plasticity over Local Adaptation. American Naturalist, 2002, 160, 271-283.	2.1	553
5	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
6	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
7	Comparative and metaâ€analytic insights into life extension via dietary restriction. Aging Cell, 2012, 11, 401-409.	6.7	182
8	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
9	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
10	Circumpolar dispersal by rafting in two subantarctic kelp-dwelling crustaceans. Marine Ecology - Progress Series, 2010, 405, 221-230.	1.9	161
11	A Theoretical Investigation of Speciation by Reinforcement. American Naturalist, 1986, 128, 241-262.	2.1	124
12	Contemporary habitat discontinuity and historic glacial ice drive genetic divergence in Chilean kelp. BMC Evolutionary Biology, 2010, 10, 203.	3.2	121
13	Phylogeographical disjunction in abundant high-dispersal littoral gastropods. Molecular Ecology, 2005, 14, 2789-2802.	3.9	105
14	Molecular systematics of the marine gastropod families Trochidae and Calliostomatidae (Mollusca:) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5	2.7	89
15	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
16	Predictive adaptive responses in perspective. Trends in Endocrinology and Metabolism, 2008, 19, 109-110.	7.1	87
17	Population-epigenetic models of selection. Theoretical Population Biology, 2012, 81, 232-242.	1.1	87
18	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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19	Identifying Cliques of Convergent Characters: Concerted Evolution in the Cormorants and Shags. Systematic Biology, 2010, 59, 433-445.	5.6	75
20	Water stratification in the marine biome restricts vertical environmental DNA (eDNA) signal dispersal. Environmental DNA, 2020, 2, 99-111.	5.8	74
21	Hop, step and gape: do the social displays of the Pelecaniformes reflect phylogeny?. Animal Behaviour, 1996, 51, 273-291.	1.9	68
22	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
23	Skeletal carbonate mineralogy of New Zealand bryozoans. Marine Geology, 1998, 151, 27-46.	2.1	63
24	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
25	Speciesâ€level biodiversity assessment using marine environmental DNA metabarcoding requires protocol optimization and standardization. Ecology and Evolution, 2019, 9, 1323-1335.	1.9	62
26	Glacial oceanographic contrasts explain phylogeography of Australian bull kelp. Molecular Ecology, 2009, 18, 2287-2296.	3.9	58
27	Untangling Long Branches: Identifying Conflicting Phylogenetic Signals Using Spectral Analysis, Neighbor-Net, and Consensus Networks. Systematic Biology, 2005, 54, 620-633.	5.6	56
28	Non-conflict theories for the evolution of genomic imprinting. Heredity, 2014, 113, 112-118.	2.6	56
29	Transoceanic genetic similarities of kelpâ€associated sea slug populations: longâ€distance dispersal via rafting?. Journal of Biogeography, 2014, 41, 2357-2370.	3.0	56
30	Passive rafting is a powerful driver of transoceanic gene flow. Biology Letters, 2013, 9, 20120821.	2.3	55
31	Genetic Conflicts, Multiple Paternity and the Evolution of Genomic Imprinting. Genetics, 1998, 148, 893-904.	2.9	55
32	The Correlation Between Relatives on the Supposition of Genomic Imprinting. Genetics, 2002, 161, 411-417.	2.9	51
33	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
34	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
35	POPULATION GENETICS AND EVOLUTION OF GENOMIC IMPRINTING. Annual Review of Genetics, 2000, 34, 457-477.	7.6	46
36	The hidden science of eugenics. Nature, 1995, 374, 302-304.	27.8	45

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37	Multigene phylogeny of the southern bull-kelp genus Durvillaea (Phaeophyceae: Fucales). Molecular Phylogenetics and Evolution, 2010, 57, 1301-1311.	2.7	45
38	Classification of the cormorants of the world. Molecular Phylogenetics and Evolution, 2014, 79, 249-257.	2.7	45
39	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
40	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
41	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
42	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
43	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
44	The Evolution of Sex-Specific Dominance in Response to Sexually Antagonistic Selection. American Naturalist, 2016, 187, 658-666.	2.1	37
45	Phylogeography of Kauri Snails and their allies from Northland, New Zealand (Mollusca: Gastropoda:) Tj ETQq1 1 0.784314 rgBT /Over	2.7	36
46	Simultaneous polyphenism and cryptic species in an intertidal limpet from New Zealand. Molecular Phylogenetics and Evolution, 2007, 45, 470-479.	2.7	36
47	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
48	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
49	Biogeography Off the Tracks. Systematic Biology, 2013, 62, 494-498.	5.6	35
50	The Evolution of Genomic Imprinting: Two Modifier-Locus Models. Theoretical Population Biology, 1997, 51, 23-35.	1.1	34
51	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
52	Exploring epiallele stability in a population-epigenetic model. Theoretical Population Biology, 2013, 83, 136-144.	1.1	33
53	For Host's Sake: The Pluses of Parasite Preservation. Trends in Ecology and Evolution, 2016, 31, 341-343.	8.7	33
54	Taxonomy and nomenclature of black nerites (Gastropoda:Neritimorpha:Nerita) from the South Pacific. Invertebrate Systematics, 2007, 21, 229.	1.3	32

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55	Patch choice with competitive asymmetries and perceptual limits: the importance of history. Animal Behaviour, 1995, 50, 497-508.	1.9	31
56	The Maintenance of Single-Locus Polymorphism. II. The Evolution of Fitnesses and Allele Frequencies. American Naturalist, 1991, 138, 1354-1371.	2.1	29
57	Effects of Genetic Drift and Gene Flow on the Selective Maintenance of Genetic Variation. Genetics, 2013, 194, 235-244.	2.9	28
58	Reinforcement, Species, and Speciation: A Reply to Butlin. American Naturalist, 1987, 130, 958-962.	2.1	27
59	Durvillaea poha sp. nov. (Fucales, Phaeophyceae): a buoyant southern bull-kelp species endemic to New Zealand. Phycologia, 2012, 51, 151-156.	1.4	27
60	Evolutionary Genetic Models of the Ovarian Time Bomb Hypothesis for the Evolution of Genomic Imprinting. Genetics, 2002, 162, 425-439.	2.9	27
61	Quantifying the Effect of Predation Risk on Foraging Bullies: No Need to Assume an IFD. Ecology, 1994, 75, 2220.	3.2	26
62	Perceptual constraints on optimal foraging: The effects of variation among foragers. Evolutionary Ecology, 1996, 10, 331-339.	1.2	26
63	Influence of Mom and Dad: Quantitative Genetic Models for Maternal Effects and Genomic Imprinting. Genetics, 2006, 173, 2297-2316.	2.9	26
64	Effects of genomic imprinting on quantitative traits. Genetica, 2009, 136, 285-293.	1.1	26
65	PHYLOGENY, BIOGEOGRAPHY, AND TAXONOMY OF AUSTRALASIAN TEALS. Auk, 2000, 117, 154.	1.4	25
66	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
67	Cladogenesis as the result of long-distance rafting events in South Pacific topshells (Gastropoda,) Tj ETQq1 1 0.784314 rgBT /Overloc	2.3	24
68	The phylogenetic position of the GalÃ¡pagos Cormorant. Molecular Phylogenetics and Evolution, 2009, 53, 94-98.	2.7	23
69	Phylogeny, Biogeography, and Taxonomy of Australasian Teals. Auk, 2000, 117, 154-163.	1.4	22
70	Evolutionary consequences of microhabitat: population-genetic structuring in kelp- vs. rock-associated chitons. Molecular Ecology, 2011, 20, 4915-4924.	3.9	22
71	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
72	Mutation-Selection Balance Under Genomic Imprinting at an Autosomal Locus. Genetics, 1997, 147, 281-287.	2.9	20

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73	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
74	The adaptive invasion of epialleles in a heterogeneous environment. Theoretical Population Biology, 2013, 88, 1-8.	1.1	19
75	The phylogenetic relationships of the extant pelicans inferred from DNA sequence data. Molecular Phylogenetics and Evolution, 2013, 66, 215-222.	2.7	19
76	Single-Locus Polymorphism in a Heterogeneous Two-Deme Model. Genetics, 2007, 176, 1625-1633.	2.9	18
77	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
78	Reply from H.G. Spencer, A.G. Clark and M.W. Feldman. Trends in Ecology and Evolution, 1999, 14, 359.	8.7	17
79	Adaptive dynamics, game theory and evolutionary population genetics. Journal of Evolutionary Biology, 2005, 18, 1191-1193.	1.7	17
80	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
81	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
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	The evolutionary potential of paramutation: A population-epigenetic model. Theoretical Population Biology, 2013, 88, 9-19.	1.1	16
84	Frequency-Dependent Selection With Dominance: A Window Onto the Behavior of the Mean Fitness. Genetics, 2004, 167, 499-512.	2.9	15
85	Evolution of Fitnesses and Allele Frequencies in a Population With Spatially Heterogeneous Selection Pressures. Genetics, 2007, 177, 1743-1751.	2.9	15
86	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
87	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
88	Strong Phylogeographic Structure in a Sedentary Seabird, the Stewart Island Shag (Leucocarbo) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 1	2.5	15
89	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
90	Genetic variation and prevalence of blood parasites do not correlate among bird species. Journal of Zoology, 2000, 252, 381-388.	1.7	14

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91	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
92	Speciation, range contraction and extinction in the endemic New Zealand King Shag complex. Molecular Phylogenetics and Evolution, 2017, 115, 197-209.	2.7	14
93	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
94	Genetic drift on networks: Ploidy and the time to fixation. Theoretical Population Biology, 2008, 74, 283-290.	1.1	12
95	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
96	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
97	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
98	Further Properties of Gavriletsâ€™ One-Locus Two-Allele Model of Maternal Selection. Genetics, 2003, 164, 1689-1692.	2.9	11
99	The evolutionary construction of molecular polymorphisms. New Zealand Journal of Botany, 1993, 31, 249-256.	1.1	10
100	Species assignation amongst morphologically cryptic larval Digenea isolated from New Zealand topshells (Gastropoda: Trochidae). Parasitology Research, 2007, 101, 433-441.	1.6	9
101	Systematics and phylogeny of a new cryptic species of Diloma Philippi (Mollusca: Gastropoda:) Tj ETQq1 1 0.784314 rgBT /Overlock 1 Invertebrate Systematics, 2009, 23, 19.	1.3	9
102	Mutation-Selection Balance at a Modifier-of-Imprinting Locus. Genetics, 1996, 144, 361-367.	2.9	9
103	The Evolution of Genomic Imprinting via Variance Minimization: An Evolutionary Genetic Model. Genetics, 2003, 165, 205-222.	2.9	9
104	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
105	Valve microstructure and phylomineralogy of New Zealand chitons. Journal of Structural Biology, 2017, 197, 250-259.	2.8	8
106	Trans-Tasman genetic connectivity in the intertidal air-breathing slug Onchidella nigricans. Marine Ecology - Progress Series, 2016, 562, 93-100.	1.9	8
107	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
108	Population Models of Genomic Imprinting. II. Maternal and Fertility Selection. Genetics, 2006, 173, 2391-2398.	2.9	7

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109	Evolution of Fitnesses in Structured Populations With Correlated Environments. Genetics, 2008, 179, 1469-1478.	2.9	7
110	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
111	Reef formation versus solitariness in two New Zealand serpulids does not involve cryptic species. Aquatic Biology, 2012, 16, 97-103.	1.4	7
112	Complex dynamics occur in a single-locus, multiallelic model of general frequency-dependent selection. Theoretical Population Biology, 2009, 76, 292-298.	1.1	6
113	Phylogeographic patterns in New Zealand and temperate Australian cantharidines (Mollusca:) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Phylogenetics and Evolution, 2016, 100, 333-344.	2.7	6
114	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
115	Beyond Equilibria: The Neglected Role of History in Ecology and Evolution. Quarterly Review of Biology, 2020, 95, 311-321.	0.1	6
116	Killing the Behavioral Zombie: Genes, Evolution, and Why Behavior Isnâ€™t Special. BioScience, 2020, 70, 515-520.	4.9	6
117	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
118	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
119	Models of Frequency-Dependent Selection with Mutation from Parental Alleles. Genetics, 2013, 195, 231-242.	2.9	5
120	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
121	Skeletal mineralogy of scaphopods: an unusual uniformity. Journal of Molluscan Studies, 2016, 82, 344-348.	1.2	4
122	A model of optimal timing for a predictive adaptive response. Journal of Developmental Origins of Health and Disease, 2021, , 1-7.	1.4	4
123	Taxonomic consistency and nomenclatural rules within oysters: Comment on Li et al. (2021). Molecular Phylogenetics and Evolution, 2022, 170, 107437.	2.7	4
124	Avoiding extinction under nonlinear environmental change: models of evolutionary rescue with plasticity. Biology Letters, 2021, 17, 20210459.	2.3	4
125	Random genetic drift and selection in a triallelic locus: a continuous diffusion model. Mathematical Biosciences, 1992, 108, 127-139.	1.9	3
126	Archival DNA reveals cryptic biodiversity within the Spotted Shag (Phalacrocorax punctatus) from New Zealand. Condor, 2019, 121, .	1.6	3

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127	The phylogenetic placement of the enigmatic Indian Cormorant, Phalacrocorax fuscicollis (Phalacrocoracidae). Molecular Phylogenetics and Evolution, 2019, 130, 227-232.	2.7	3
128	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
129	Rapid radiation of Southern Ocean shags in response to receding sea ice. Journal of Biogeography, 2022, 49, 942-953.	3.0	3
130	Measuring mating preferences: the use of Manly's beta. Heredity, 1988, 60, 305-310.	2.6	2
131	Assortative versus selective mating: Is the distinction worthwhile?. Biodemography and Social Biology, 1992, 39, 310-315.	1.0	2
132	Genomic Imprinting Leads to Less Selectively Maintained Polymorphism on X Chromosomes. Genetics, 2012, 192, 1455-1464.	2.9	2
133	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
134	Genomic imprinting: theories and data. Heredity, 2014, 113, 93-95.	2.6	2
135	The Maintenance of Single-Locus Polymorphism by Maternal Selection. G3: Genes, Genomes, Genetics, 2015, 5, 963-969.	1.8	2
136	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
137	The Selective Maintenance of Allelic Variation Under Generalized Dominance. G3: Genes, Genomes, Genetics, 2016, 6, 3725-3732.	1.8	2
138	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
139	Polymorphic microsatellite DNA markers in the mudflat topshell Diloma subrostrata (Gastropoda,) Tj ETQq1 1 0.784314 rgBT /Overlock	1.7	1
140	Developmental Origins of Health and Disease across Generations – Theory, Observation, Experiment. , 2009, , 52-64.		1
141	Case 3706 â€" Trochus (Osilinus ?) Capillaceus Philippi, 1849 (currently Cantharidus capillaceus;) Tj ETQq1 1 0.784314 rgBT /Overlock Zoological Nomenclature, 2017, 74, 8.	0.1	1
142	Graph-structured populations and the Hillâ€"Robertson effect. Royal Society Open Science, 2021, 8, 201831.	2.4	1
143	Structured Populations and the Maintenance of Sex. Lecture Notes in Computer Science, 2013, , 56-67.	1.3	1
144	A further perspective on speciation by reinforcement. Theoretical Biology Forum, 2020, 113, 63-66.	0.2	1

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145	CYTOPLASMIC INCOMPATIBILITY: WHY IS IT SEXUALLY ASYMMETRIC?. Evolution; International Journal of Organic Evolution, 1995, 49, 1277-1280.	2.3	0
146	Cytoplasmic Incompatibility: Why is it Sexually Asymmetric?. Evolution; International Journal of Organic Evolution, 1995, 49, 1277.	2.3	0
147	Onward and upward. Metascience, 1998, 7, 52-64.	0.3	0
148	Defending the group from the terror within. Metascience, 2001, 10, 192-202.	0.3	0
149	Population-genetic models of sex-limited genomic imprinting. Theoretical Population Biology, 2017, 115, 35-44.	1.1	0
150	Phylogeography of the intertidal marine bivalve Lasaea hinemoa (Mollusca: Bivalvia) in New Zealand. Molluscan Research, 2021, 41, 191-203.	0.7	0
151	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