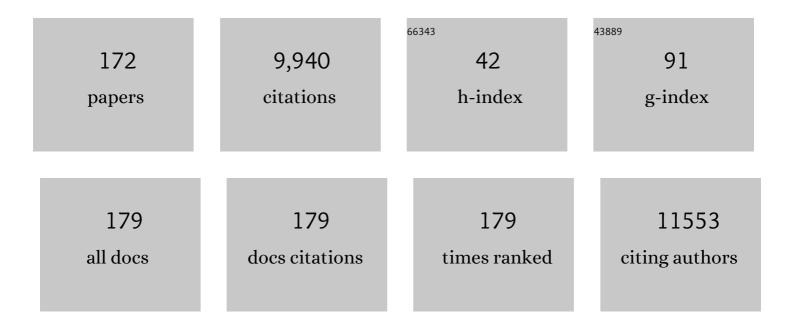
David M Lambert

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Second asymptomatic carotid surgery trial (ACST-2): a randomised comparison of carotid artery stenting versus carotid endarterectomy. Lancet, The, 2021, 398, 1065-1073.	13.7	133
2	Insights Into Aboriginal Australian Mortuary Practices: Perspectives From Ancient DNA. Frontiers in Ecology and Evolution, 2020, 8, .	2.2	4
3	A different paradigm for the colonisation of Sahul. Archaeology in Oceania, 2020, 55, 182-191.	0.7	1
4	Scientific and personal reflections on an iconoclastic thinker in evolutionary biology: hugh edward ha Idane paterson. Theoretical Biology Forum, 2020, 113, 59-62.	0.2	0
5	DNA barcoding a unique avifauna: an important tool for evolution, systematics and conservation. BMC Evolutionary Biology, 2019, 19, 52.	3.2	24
6	Mitogenomic diversity in Sacred Ibis Mummies sheds light on early Egyptian practices. PLoS ONE, 2019, 14, e0223964.	2.5	14
7	Archaeogenetics and human evolution: the ontogeny of a biological discipline. World Archaeology, 2019, 51, 546-559.	1.1	17
8	Editorial. Theoretical biology forum. The next century. Theoretical Biology Forum, 2019, 112, 9-10.	0.2	0
9	Disentangling Immediate Adaptive Introgression from Selection on Standing Introgressed Variation in Humans. Molecular Biology and Evolution, 2018, 35, 623-630.	8.9	46
10	Ancient nuclear genomes enable repatriation of Indigenous human remains. Science Advances, 2018, 4, eaau5064.	10.3	41
11	The Sacred Ibis debate: The first test of evolution. PLoS Biology, 2018, 16, e2005558.	5.6	6
12	The prehistoric peopling of Southeast Asia. Science, 2018, 361, 88-92.	12.6	291
13	Editorial. Preface to a new century of «Theoretical Biology Forum». Theoretical Biology Forum, 2018, 111, 9-10.	0.2	0
14	Eggshell palaeogenomics: Palaeognath evolutionary history revealed through ancient nuclear and mitochondrial DNA from Madagascan elephant bird (Aepyornis sp.) eggshell. Molecular Phylogenetics and Evolution, 2017, 109, 151-163.	2.7	65
15	Co-option of the cardiac transcription factor Nkx2.5 during development of the emu wing. Nature Communications, 2017, 8, 132.	12.8	21
16	Data sharing: do scientists know best?. Nature, 2017, 548, 281-281.	27.8	2
17	East Asian allopatry and north Eurasian sympatry in Longâ€ŧailed Tit lineages despite similar population dynamics during the late Pleistocene. Zoologica Scripta, 2016, 45, 115-126.	1.7	14
18	Limb patterning genes and heterochronic development of the emu wing bud. EvoDevo, 2016, 7, 26.	3.2	12

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19	A genomic history of Aboriginal Australia. Nature, 2016, 538, 207-214.	27.8	439
20	Genomic analyses inform on migration events during the peopling of Eurasia. Nature, 2016, 538, 238-242.	27.8	360
21	A zoogeographical boundary between the Palaearctic and Sinoâ€Japanese realms documented by consistent north/south phylogeographical divergences in three woodland birds in eastern China. Journal of Biogeography, 2016, 43, 2099-2112.	3.0	12
22	Ancient mtDNA sequences from the First Australians revisited. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 6892-6897.	7.1	26
23	Distance-dependent patterns of molecular divergences in tuatara mitogenomes. Scientific Reports, 2015, 5, 8703.	3.3	5
24	A Concentrated Hydrochloric Acidâ€based Method for Complete Recovery of <scp>DNA</scp> from Bone. Journal of Forensic Sciences, 2015, 60, 1553-1557.	1.6	7
25	Radiocarbon dating of Sacred Ibis mummies from ancient Egypt. Journal of Archaeological Science: Reports, 2015, 4, 355-361.	0.5	9
26	Impacts of low coverage depths and post-mortem DNA damage on variant calling: a simulation study. BMC Genomics, 2015, 16, 19.	2.8	26
27	Kiwi genome provides insights into evolution of a nocturnal lifestyle. Genome Biology, 2015, 16, 147.	8.8	68
28	A recent bottleneck of Y chromosome diversity coincides with a global change in culture. Genome Research, 2015, 25, 459-466.	5.5	348
29	Ancient population genomics and the study of evolution. Philosophical Transactions of the Royal Society B: Biological Sciences, 2015, 370, 20130381.	4.0	18
30	Complete mitochondrial genomes of Tuatara endemic to different islands of New Zealand. Mitochondrial DNA, 2015, 26, 25-26.	0.6	1
31	The mysterious Spotted Green Pigeon and its relation to the Dodo and its kindred. BMC Evolutionary Biology, 2014, 14, 136.	3.2	12
32	Whole-genome analyses resolve early branches in the tree of life of modern birds. Science, 2014, 346, 1320-1331.	12.6	1,583
33	Comparative genomics reveals insights into avian genome evolution and adaptation. Science, 2014, 346, 1311-1320.	12.6	895
34	Two Antarctic penguin genomes reveal insights into their evolutionary history and molecular changes related to the Antarctic environment. GigaScience, 2014, 3, 27.	6.4	72
35	Second generation DNA sequencing of the mitogenome of the Chinstrap penguin and comparative genomics of Antarctic penguins. Mitochondrial DNA, 2014, 25, 271-272.	0.6	2
36	DNA fingerprinting in zoology: past, present, future. Investigative Genetics, 2014, 5, 3.	3.3	45

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37	Genomic structure in Europeans dating back at least 36,200 years. Science, 2014, 346, 1113-1118.	12.6	287
38	Reconstruction and in vivo analysis of the extinct tbx5 gene from ancient wingless moa (Aves:) Tj ETQq0 0 0 rgBT	/Qverlock	10 Tf 50 70

39	Population genetic structure and taxonomy of the common dolphin (<i>Delphinus</i> sp.) at its southernmost range limit: New Zealand waters. Marine Mammal Science, 2014, 30, 44-63.	1.8	14
40	Identification, Classification, and Growth of Moa Chicks (Aves: Dinornithiformes) from the Genus Euryapteryx. PLoS ONE, 2014, 9, e99929.	2.5	4
41	Complex Species Status for Extinct Moa (Aves: Dinornithiformes) from the Genus Euryapteryx. PLoS ONE, 2014, 9, e90212.	2.5	3
42	Using ancient DNA to enhance museum collections: a case study of rare kiwi (Apteryxspp.) specimens. Journal of the Royal Society of New Zealand, 2013, 43, 119-127.	1.9	11
43	Evidence for a recent origin of penguins. Biology Letters, 2013, 9, 20130748.	2.3	27
44	Gene flow maintains genetic diversity and colonization potential in recently rangeâ€expanded populations of an <scp>O</scp> riental bird, the <scp>L</scp> ightâ€vented <scp>B</scp> ulbul (<i><scp>P</scp>ycnonotus sinensis</i> , <scp> A</scp> ves: <scp>P</scp> ycnonotidae). Diversity and Distributions, 2013, 19, 1248-1262.	4.1	28
45	Towards a million-year-old genome. Nature, 2013, 499, 34-35.	27.8	25
46	Ground tit genome reveals avian adaptation to living at high altitudes in the Tibetan plateau. Nature Communications, 2013, 4, 2071.	12.8	229
47	Methylphenidate Side Effect Profile Is Influenced by Genetic Variation in the Attention-Deficit/Hyperactivity Disorder-Associated CES1 Gene. Journal of Child and Adolescent Psychopharmacology, 2013, 23, 655-664.	1.3	29
48	Late-Holocene Penguin Occupation and Diet at King George Island Antarctic Peninsula. Antarctic Research Series, 2013, , 171-180.	0.2	9
49	Highly Informative Ancient DNA â€~Snippets' for New Zealand Moa. PLoS ONE, 2013, 8, e50732.	2.5	5
50	Non-Darwinian Evolution. , 2013, , 87-89.		0
51	King penguin population on Macquarie Island recovers ancient DNA diversity after heavy exploitation in historic times. Biology Letters, 2012, 8, 586-589.	2.3	12
52	Selective Constraints Determine the Time Dependency of Molecular Rates for Human Nuclear Genomes. Genome Biology and Evolution, 2012, 4, 1127-1132.	2.5	8
53	Resurrecting ancient animal genomes: The extinct moa and more. BioEssays, 2012, 34, 661-669.	2.5	17
54	Ultraviolet visual sensitivity in three avian lineages: paleognaths, parrots, and passerines. Journal of Comparative Physiology A: Neuroethology, Sensory, Neural, and Behavioral Physiology, 2012, 198, 495-510.	1.6	59

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55	Adélie penguins and temperature changes in Antarctica: a longâ€ŧerm view. Integrative Zoology, 2012, 7, 113-120.	2.6	15
56	Ancient DNA Analyses Reveal Contrasting Phylogeographic Patterns amongst Kiwi (Apteryx spp.) and a Recently Extinct Lineage of Spotted Kiwi. PLoS ONE, 2012, 7, e42384.	2.5	33
57	Birdstrikes and barcoding: can DNA methods help make the airways safer?. Molecular Ecology Resources, 2011, 11, 38-45.	4.8	15
58	Are BOLD searches scientific? A response to Federhen (2011). Molecular Ecology Resources, 2011, 11, 939-940.	4.8	1
59	UVS is rare in seabirds. Vision Research, 2011, 51, 1333-1337.	1.4	23
60	An Aboriginal Australian Genome Reveals Separate Human Dispersals into Asia. Science, 2011, 334, 94-98.	12.6	675
61	Time Dependency of Molecular Evolutionary Rates? Yes and No. Genome Biology and Evolution, 2011, 3, 1324-1328.	2.5	21
62	Ancient DNA Recovers the Origins of MÄori Feather Cloaks. Molecular Biology and Evolution, 2011, 28, 2741-2750.	8.9	29
63	Ancient DNA Suggests Dwarf and â€~Giant' Emu Are Conspecific. PLoS ONE, 2011, 6, e18728.	2.5	16
64	Genetic diversity and taxonomy: a reassessment of species designation in tuatara (Sphenodon: Reptilia). Conservation Genetics, 2010, 11, 1063-1081.	1.5	73
65	Next generation sequencing and analysis of a conserved transcriptome of New Zealand's kiwi. BMC Evolutionary Biology, 2010, 10, 387.	3.2	13
66	Face of the past reconstructed. Nature, 2010, 463, 739-740.	27.8	2
67	Ancient DNA reveals extreme egg morphology and nesting behavior in New Zealand's extinct moa. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 16201-16206.	7.1	49
68	Conserved primers for DNA barcoding historical and modern samples from New Zealand and Antarctic birds. Molecular Ecology Resources, 2010, 10, 431-438.	4.8	43
69	The Molecular Ecology of the Extinct New Zealand Huia. PLoS ONE, 2009, 4, e8019.	2.5	12
70	Molecular and morphological evolution in tuatara are decoupled. Trends in Genetics, 2009, 25, 16-18.	6.7	21
71	High mitogenomic evolutionary rates and time dependency. Trends in Genetics, 2009, 25, 482-486.	6.7	90
72	Molecular evidence for the identity of the Magenta petrel. Molecular Ecology Resources, 2009, 9, 458-461.	4.8	6

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73	Ancient DNA Resolves Identity and Phylogeny of New Zealand's Extinct and Living Quail (Coturnix sp.). PLoS ONE, 2009, 4, e6400.	2.5	9
74	Microsatellite DNA loci identify individuals and provide no evidence for multiple paternity in wild tuatara (Sphenodon: Reptilia). Conservation Genetics, 2008, 9, 1039-1043.	1.5	9
75	High mitochondrial and nuclear genetic diversity in one of the world's most endangered seabirds, the Chatham Island Taiko (Pterodroma magentae). Conservation Genetics, 2008, 9, 1293-1301.	1.5	34
76	DNA barcoding of animal species—response to DeSalle. BioEssays, 2008, 30, 92-93.	2.5	9
77	Excess of unpaired males in one of the World's most endangered seabirds, the Chatham Island taiko Pterodroma magentae. Journal of Avian Biology, 2008, 39, 359-363.	1.2	14
78	Ancient genetic variation in one of the world's rarest seabirds. Heredity, 2008, 101, 543-547.	2.6	5
79	New Genetic Approach to Detecting Individuals of Rare and Endangered Species. Conservation Biology, 2008, 22, 1267-1276.	4.7	7
80	Ancient DNA and conservation: lessons from the endangered kiwi of New Zealand. Molecular Ecology, 2008, 17, 2174-2184.	3.9	54
81	Rapid molecular evolution in a living fossil. Trends in Genetics, 2008, 24, 106-109.	6.7	60
82	New developments in ancient genomics. Trends in Ecology and Evolution, 2008, 23, 386-393.	8.7	83
83	Genetic identification of moa remains recovered from Tiniroto, Gisborne. Journal of the Royal Society of New Zealand, 2008, 38, 231-235.	1.9	6
84	Mutation and Evolutionary Rates in Adélie Penguins from the Antarctic. PLoS Genetics, 2008, 4, e1000209.	3.5	79
85	Spatial Attentional Bias as a Marker of Genetic Risk, Symptom Severity, and Stimulant Response in ADHD. Neuropsychopharmacology, 2008, 33, 2536-2545.	5.4	41
86	Excess of unpaired males in one of the World's most endangered seabirds, the Chatham Island taiko Pterodroma magentae. Journal of Avian Biology, 2008, .	1.2	1
87	The relationships and origins of the New Zealand wattlebirds (Passeriformes, Callaeatidae) from DNA sequence analyses. Molecular Phylogenetics and Evolution, 2007, 43, 480-492.	2.7	28
88	Nuclear microsatellite DNA markers for New Zealand kiwi (Apteryx spp.). Molecular Ecology Notes, 2006, 6, 227-229.	1.7	13
89	Unusual electrophoretic mobility of a DNA fragment of the universal â€~non-ratite' sexing marker CHD allows sexing of New Zealand's endangered kiwi ratite Apteryx spp Ibis, 2006, 148, 167-168.	1.9	10

90 Ancient genomics is born. Nature, 2006, 444, 275-276.

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91	A molecular phylogeny of New Zealand's Petroica (Aves: Petroicidae) species based on mitochondrial DNA sequences. Molecular Phylogenetics and Evolution, 2006, 40, 844-855.	2.7	38
92	Characterization of variable microsatellite loci in Forbes' parakeet (Cyanoramphus forbesi) and their use in other parrots. Conservation Genetics, 2006, 6, 651-654.	1.5	15
93	Serial population bottlenecks and genetic variation: Translocated populations of the New Zealand Saddleback (Philesturnus carunculatus rufusater). Conservation Genetics, 2005, 6, 1-14.	1.5	39
94	From The Cover: Microevolution and mega-icebergs in the Antarctic. Proceedings of the National Academy of Sciences of the United States of America, 2005, 102, 16717-16722.	7.1	52
95	Mutational Bias in Penguin Microsatellite DNA. Journal of Heredity, 2005, 96, 566-571.	2.4	10
96	Reconstructing the tempo and mode of evolution in an extinct clade of birds with ancient DNA: The giant moas of New Zealand. Proceedings of the National Academy of Sciences of the United States of America, 2005, 102, 8257-8262.	7.1	82
97	Is a Large-Scale DNA-Based Inventory of Ancient Life Possible?. Journal of Heredity, 2005, 96, 279-284.	2.4	71
98	The Use of Morphometric Measurements to Sex Yellow-eyed Penguins. Waterbirds, 2004, 27, 96-101.	0.3	35
99	Genetic drift outweighs balancing selection in shaping post-bottleneck major histocompatibility complex variation in New Zealand robins (Petroicidae). Molecular Ecology, 2004, 13, 3709-3721.	3.9	153
100	Effect of Extra-Pair Paternity on Effective Population Size in a Reintroduced Population of the Endangered Hihi, and Potential for Behavioural Management. Conservation Genetics, 2004, 5, 381-393.	1.5	28
101	Gene duplication and gene conversion in class�II MHC genes of New Zealand robins (Petroicidae). Immunogenetics, 2004, 56, 178-91.	2.4	77
102	Title is missing!. Conservation Genetics, 2003, 4, 265-274.	1.5	26
103	An evaluation of methods of blood preservation for RT-PCR from endangered species. Conservation Genetics, 2003, 4, 651-654.	1.5	11
104	Evidence for specificity of psittacine beak and feather disease viruses among avian hosts. Virology, 2003, 306, 109-115.	2.4	78
105	Nuclear DNA sequences detect species limits in ancient moa. Nature, 2003, 425, 175-178.	27.8	110
106	Ancient DNA Enables Timing of the Pleistocene Origin and Holocene Expansion of Two Adelie Penguin Lineages in Antarctica. Molecular Biology and Evolution, 2003, 21, 240-248.	8.9	82
107	Rates of Evolution in Ancient DNA from Adelie Penguins. Science, 2002, 295, 2270-2273.	12.6	274
108	A DNA test to sex ratite birds. Molecular Ecology, 2002, 11, 851-856.	3.9	36

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109	New DNA markers for penguins. Conservation Genetics, 2002, 3, 341-344.	1.5	28
110	Mating system and genetic variation in the endangered New Zealand takahe. Conservation Genetics, 2002, 3, 427-434.	1.5	15
111	Cene flow on the ice: genetic differentiation among Adélie penguin colonies around Antarctica. Molecular Ecology, 2001, 10, 1645-1656.	3.9	71
112	A comparison of five methods for assignment of sex in the takahe (Aves: Porphyrio mantelli). Journal of Zoology, 2001, 253, 281-292.	1.7	9
113	Microsatellite DNA markers for tuatara (Sphenodon spp.). Conservation Genetics, 2001, 2, 183-185.	1.5	16
114	Sexing the Critically Endangered Kakapo Strigops habroptilus. Emu, 2000, 100, 336-339.	0.6	6
115	Molecular ecology and biological control: the mating system of a marsupial pest. Molecular Ecology, 2000, 9, 723-733.	3.9	20
116	A novel restriction fragment length polymorphism for petrels or tube-nosed seabirds. Molecular Ecology, 2000, 9, 1915-1917.	3.9	0
117	Microsatellite primers for the kakapo (Strigops habroptilus) and their utility in other parrots. Conservation Genetics, 2000, 1, 93-95.	1.5	15
118	Title is missing!. Conservation Genetics, 2000, 1, 103-113.	1.5	13
119	ASW : a gene with conserved avian W-linkage and female specific expression in chick embryonic gonad. Development Genes and Evolution, 2000, 210, 243-249.	0.9	112
120	A repeat complex in the mitochondrial control region of Adélie penguins from Antarctica. Genome, 2000, 43, 613-618.	2.0	30
121	Genetic monogamy mirrors social monogamy in the Fiordland crested penguin. New Zealand Journal of Zoology, 2000, 27, 311-316.	1.1	6
122	Molecular sexing of individual kakapo, Strigops habroptilus Aves, from faeces. Molecular Ecology, 1999, 8, 1349-1350.	3.9	16
123	Floater males gain reproductive success through extrapair fertilizations in the stitchbird. Animal Behaviour, 1999, 58, 321-328.	1.9	83
124	Industrial Melanism: A Classic Example of Another Kind?. BioScience, 1999, 49, 1021.	4.9	1
125	"Wife-Sharing" in the Tasmanian Native Hen (Gallinula mortierii): Is It Caused by a Male-Biased Sex Ratio?. Auk, 1998, 115, 528-532.	1.4	6
126	Patterns of prehistoric human mobility in Polynesia indicated by mtDNA from the Pacific rat. Proceedings of the National Academy of Sciences of the United States of America, 1998, 95, 15145-15150.	7.1	152

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127	Phospholipase Cγ ₁ in Bovine Rod Outer Segments: Immunolocalization and Lightâ€Dependent Binding to Membranes. Journal of Neurochemistry, 1998, 70, 171-178.	3.9	24
128	Genetic Relatedness and Alloparental Behaviour in a Captive Group of Spider Monkeys (Ateles) Tj ETQq0 0 0 rgBT	Qverlock	10 Tf 50 70
129	The Effects of Population Bottlenecks on Multilocus DNA Variation in Robins. Journal of Heredity, 1997, 88, 179-186.	2.4	29
130	Social and Sexual Monogamy in Translocated New Zealand Robin Populations Detected Using Minisatellite DNA. Auk, 1997, 114, 120-126.	1.4	40
131	Captive management and molecular sexing of endangered avian species: An application to the black stilt Himantopus novaezelandiae and hybrids. Biological Conservation, 1997, 82, 81-86.	4.1	17
132	Is the black robin in genetic peril?. Molecular Ecology, 1997, 6, 21-28.	3.9	80
133	Ancient DNA from polynesian rats: Extraction, amplification and sequence from single small bones. Electrophoresis, 1997, 18, 1534-1537.	2.4	42
134	Minisatellite DNA markers reveal hybridisation between the endangered black robin and tomtit. Electrophoresis, 1997, 18, 1682-1687.	2.4	10
135	Kinship and genetic divergence among populations of tuatara Sphenodon punctatus as revealed by minisatellite DNA profiling. Molecular Ecology, 1996, 5, 651-658.	3.9	12
136	Mitochondrial Phylogeny of Trematomid Fishes (Nototheniidae, Perciformes) and the Evolution of Antarctic Fish. Molecular Phylogenetics and Evolution, 1996, 5, 383-390.	2.7	78
137	Molecular sexing of the communally breeding pukeko: an important ecological tool. Molecular Ecology, 1996, 5, 289-293.	3.9	16
138	Genetic continuity within, and discontinuities among, populations of leafroller moths with distinct sex-pheromones. Heredity, 1995, 75, 243-255.	2.6	4
139	DNA science and conservation. Pacific Conservation Biology, 1995, 2, 21.	1.0	36
140	What's happening in New Zealand Conservation?. Pacific Conservation Biology, 1995, 2, 1.	1.0	0
141	Genetic differences among pheromonally distinct New Zealand leafroller moths. Biochemical Systematics and Ecology, 1994, 22, 329-339.	1.3	17
142	The Effects of Blood Sampling on the Behavior and Survival of the Endangered Chatham Island Black Robin (Petroica traversi). Conservation Biology, 1994, 8, 857-862.	4.7	22
143	Single- and multilocus DNA fingerprinting of communally breeding pukeko: do copulations or dominance ensure reproductive success?. Proceedings of the National Academy of Sciences of the United States of America, 1994, 91, 9641-9645.	7.1	63

144Patterns of reproductive success determined by DNA fingerprinting in a communally breeding oceanic
bird. Biological Journal of the Linnean Society, 1994, 52, 31-48.1.626

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145	Sex-Specific Restriction Fragments and Sex Ratios Revealed by DNA Fingerprinting in the Brown Skua. Journal of Heredity, 1992, 83, 350-355.	2.4	41
146	Species and Neo-Darwinism. Systematic Zoology, 1990, 39, 399.	1.6	5
147	Genetics ofPotamopyrgus antipodarum(Gastropoda: Prosobranchia): Variation in unisexual populations. New Zealand Journal of Zoology, 1990, 17, 65-72.	1.1	4
148	Organisms not Species Evolve: A Reply to Ghiselin. Systematic Zoology, 1990, 39, 79.	1.6	2
149	A cladistic analysis of species of the molluscan genusPotamopyrgusbased on allozyme data. New Zealand Journal of Zoology, 1990, 17, 257-263.	1.1	3
150	Genetics ofPotamopyrgus antipodarum(Gastropoda: Prosobranchia): evidence for reproductive modes. New Zealand Journal of Zoology, 1989, 16, 435-445.	1.1	37
151	The sensitive period for yellow phenocopy induction inDrosophila melanogaster. Experientia, 1988, 44, 618-621.	1.2	3
152	Keywords and concepts in structuralist and functionalist biology. Journal of Theoretical Biology, 1988, 133, 133-145.	1.7	20
153	A genetic analysis of populations ofGalaxias maculatusfrom the Bay of Plenty: Implications for natal river return. New Zealand Journal of Marine and Freshwater Research, 1988, 22, 321-326.	2.0	31
154	Are Species Self-Defining?. Systematic Zoology, 1987, 36, 196.	1.6	49
155	Reinforcement, Species, and Speciation: A Reply to Butlin. American Naturalist, 1987, 130, 958-962.	2.1	27
156	A Theoretical Investigation of Speciation by Reinforcement. American Naturalist, 1986, 128, 241-262.	2.1	124
157	Laboratory-induced changes in the mate recognition system ofDrosophila pseudoobscura. Behavior Genetics, 1986, 16, 285-294.	2.1	4
158	The stability of the specific-mate recognition system ofDrosophila melanogaster. Behavior Genetics, 1986, 16, 369-373.	2.1	10
159	Cytogenetics of New Zealand blackflies of the genus Austrosimulium (Diptera: Simuliidae) II. Heterozygote deficiency and non-random association of inversion heterozygotes. Genetica, 1985, 66, 203-211.	1.1	Ο
160	The mating behavior of individuals ofDrosophila pseudoobscura from New Zealand. Experientia, 1985, 41, 950-952.	1.2	7
161	Scientific Prejudice, Reproductive Isolation, and Apartheid. Perspectives in Biology and Medicine, 1984, 28, 107-116.	0.5	18
162	Evolutionary Studies of the New Zealand Coastal Mosquito Opifex Fuscus (Hutton) li. Competition for Mates. Behaviour, 1984, 88, 1-12.	0.8	4

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163	Specific-mate recognition systems, phylogenies and asymmetrical evolution. Journal of Theoretical Biology, 1984, 109, 147-156.	1.7	15
164	Functionalism, structuralism, and "Ways of seeing― Journal of Theoretical Biology, 1984, 111, 787-800.	1.7	51
165	The population genetics of reinforcing selection. Genetica, 1983, 62, 15-23.	1.1	7
166	Cytogenetics of New Zealand blackflies of the genus <i>Austrosimulium</i> (Diptera: Simuliidae) 1. The cytogenetics of <i>Austrosimulium australense</i> . New Zealand Journal of Zoology, 1983, 10, 271-280.	1.1	3
167	Evolutionary Studies of the New Zealand Coastal Mosquito Opifex Fuscus (Hutton) I. Mating Behaviour. Behaviour, 1983, 84, 157-171.	0.8	13
168	A Population Genetical Study of the African Mosquito Anopheles marshallii (Theobald). Evolution; International Journal of Organic Evolution, 1983, 37, 484.	2.3	10
169	Mate recognition in members of the Drosophila nasuta complex. Animal Behaviour, 1982, 30, 438-443.	1.9	12
170	The â€~good old days' of evolutionary biology? A review of the SYSTANZ meeting on evolution, Greta Point, Wellington, 3 September 1982. New Zealand Journal of Zoology, 1982, 9, 519-520.	1.1	0
171	No significant deviation from random mating of worldwide populations of Drosophila melanogaster. Nature, 1982, 300, 437-440.	27.8	78
172	The Chromosomes of Four Species of the nasuta complex of Drosophila: I. Chromosome maps and inversion polymorphism. Journal of Heredity, 1976, 67, 92-98.	2.4	10