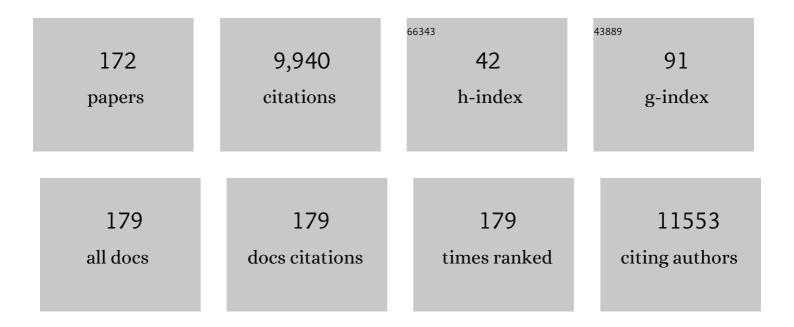
David M Lambert

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Whole-genome analyses resolve early branches in the tree of life of modern birds. Science, 2014, 346, 1320-1331.	12.6	1,583
2	Comparative genomics reveals insights into avian genome evolution and adaptation. Science, 2014, 346, 1311-1320.	12.6	895
3	An Aboriginal Australian Genome Reveals Separate Human Dispersals into Asia. Science, 2011, 334, 94-98.	12.6	675
4	A genomic history of Aboriginal Australia. Nature, 2016, 538, 207-214.	27.8	439
5	Genomic analyses inform on migration events during the peopling of Eurasia. Nature, 2016, 538, 238-242.	27.8	360
6	A recent bottleneck of Y chromosome diversity coincides with a global change in culture. Genome Research, 2015, 25, 459-466.	5.5	348
7	The prehistoric peopling of Southeast Asia. Science, 2018, 361, 88-92.	12.6	291
8	Genomic structure in Europeans dating back at least 36,200 years. Science, 2014, 346, 1113-1118.	12.6	287
9	Rates of Evolution in Ancient DNA from Adelie Penguins. Science, 2002, 295, 2270-2273.	12.6	274
10	Ground tit genome reveals avian adaptation to living at high altitudes in the Tibetan plateau. Nature Communications, 2013, 4, 2071.	12.8	229
11	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
12	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
13	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
14	A Theoretical Investigation of Speciation by Reinforcement. American Naturalist, 1986, 128, 241-262.	2.1	124
15	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
16	Nuclear DNA sequences detect species limits in ancient moa. Nature, 2003, 425, 175-178.	27.8	110
17	High mitogenomic evolutionary rates and time dependency. Trends in Genetics, 2009, 25, 482-486.	6.7	90
18	Floater males gain reproductive success through extrapair fertilizations in the stitchbird. Animal Behaviour, 1999, 58, 321-328.	1.9	83

#	Article	IF	CITATIONS
19	New developments in ancient genomics. Trends in Ecology and Evolution, 2008, 23, 386-393.	8.7	83
20	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
21	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
22	Is the black robin in genetic peril?. Molecular Ecology, 1997, 6, 21-28.	3.9	80
23	Mutation and Evolutionary Rates in Adélie Penguins from the Antarctic. PLoS Genetics, 2008, 4, e1000209.	3.5	79
24	No significant deviation from random mating of worldwide populations of Drosophila melanogaster. Nature, 1982, 300, 437-440.	27.8	78
25	Mitochondrial Phylogeny of Trematomid Fishes (Nototheniidae, Perciformes) and the Evolution of Antarctic Fish. Molecular Phylogenetics and Evolution, 1996, 5, 383-390.	2.7	78
26	Evidence for specificity of psittacine beak and feather disease viruses among avian hosts. Virology, 2003, 306, 109-115.	2.4	78
27	Gene duplication and gene conversion in class�II MHC genes of New Zealand robins (Petroicidae). Immunogenetics, 2004, 56, 178-91.	2.4	77
28	Genetic diversity and taxonomy: a reassessment of species designation in tuatara (Sphenodon: Reptilia). Conservation Genetics, 2010, 11, 1063-1081.	1.5	73
29	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
30	Gene flow on the ice: genetic differentiation among Adélie penguin colonies around Antarctica. Molecular Ecology, 2001, 10, 1645-1656.	3.9	71
31	Is a Large-Scale DNA-Based Inventory of Ancient Life Possible?. Journal of Heredity, 2005, 96, 279-284.	2.4	71
32	Kiwi genome provides insights into evolution of a nocturnal lifestyle. Genome Biology, 2015, 16, 147.	8.8	68
33	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
34	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
35	Rapid molecular evolution in a living fossil. Trends in Genetics, 2008, 24, 106-109.	6.7	60
36	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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37	Ancient DNA and conservation: lessons from the endangered kiwi of New Zealand. Molecular Ecology, 2008, 17, 2174-2184.	3.9	54
38	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
39	Functionalism, structuralism, and "Ways of seeing― Journal of Theoretical Biology, 1984, 111, 787-800.	1.7	51
40	Are Species Self-Defining?. Systematic Zoology, 1987, 36, 196.	1.6	49
41	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
42	Disentangling Immediate Adaptive Introgression from Selection on Standing Introgressed Variation in Humans. Molecular Biology and Evolution, 2018, 35, 623-630.	8.9	46
43	DNA fingerprinting in zoology: past, present, future. Investigative Genetics, 2014, 5, 3.	3.3	45
44	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
45	Ancient DNA from polynesian rats: Extraction, amplification and sequence from single small bones. Electrophoresis, 1997, 18, 1534-1537.	2.4	42
46	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
47	Spatial Attentional Bias as a Marker of Genetic Risk, Symptom Severity, and Stimulant Response in ADHD. Neuropsychopharmacology, 2008, 33, 2536-2545.	5.4	41
48	Ancient nuclear genomes enable repatriation of Indigenous human remains. Science Advances, 2018, 4, eaau5064.	10.3	41
49	Social and Sexual Monogamy in Translocated New Zealand Robin Populations Detected Using Minisatellite DNA. Auk, 1997, 114, 120-126.	1.4	40
50	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
51	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
52	Genetics ofPotamopyrgus antipodarum(Gastropoda: Prosobranchia): evidence for reproductive modes. New Zealand Journal of Zoology, 1989, 16, 435-445.	1.1	37
53	A DNA test to sex ratite birds. Molecular Ecology, 2002, 11, 851-856.	3.9	36
54	DNA science and conservation. Pacific Conservation Biology, 1995, 2, 21.	1.0	36

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55	The Use of Morphometric Measurements to Sex Yellow-eyed Penguins. Waterbirds, 2004, 27, 96-101.	0.3	35
56	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
57	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
58	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
59	A repeat complex in the mitochondrial control region of Adélie penguins from Antarctica. Genome, 2000, 43, 613-618.	2.0	30
60	The Effects of Population Bottlenecks on Multilocus DNA Variation in Robins. Journal of Heredity, 1997, 88, 179-186.	2.4	29
61	Ancient DNA Recovers the Origins of MÄori Feather Cloaks. Molecular Biology and Evolution, 2011, 28, 2741-2750.	8.9	29
62	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
63	New DNA markers for penguins. Conservation Genetics, 2002, 3, 341-344.	1.5	28
64	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
65	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
66	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
67	Reinforcement, Species, and Speciation: A Reply to Butlin. American Naturalist, 1987, 130, 958-962.	2.1	27
68	Evidence for a recent origin of penguins. Biology Letters, 2013, 9, 20130748.	2.3	27
69	Title is missing!. Conservation Genetics, 2003, 4, 265-274.	1.5	26
70	Impacts of low coverage depths and post-mortem DNA damage on variant calling: a simulation study. BMC Genomics, 2015, 16, 19.	2.8	26
71	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
72	Patterns of reproductive success determined by DNA fingerprinting in a communally breeding oceanic bird. Biological Journal of the Linnean Society, 1994, 52, 31-48.	1.6	26

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73	Towards a million-year-old genome. Nature, 2013, 499, 34-35.	27.8	25
74	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
75	DNA barcoding a unique avifauna: an important tool for evolution, systematics and conservation. BMC Evolutionary Biology, 2019, 19, 52.	3.2	24
76	UVS is rare in seabirds. Vision Research, 2011, 51, 1333-1337.	1.4	23
77	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
78	Molecular and morphological evolution in tuatara are decoupled. Trends in Genetics, 2009, 25, 16-18.	6.7	21
79	Time Dependency of Molecular Evolutionary Rates? Yes and No. Genome Biology and Evolution, 2011, 3, 1324-1328.	2.5	21
80	Co-option of the cardiac transcription factor Nkx2.5 during development of the emu wing. Nature Communications, 2017, 8, 132.	12.8	21
81	Keywords and concepts in structuralist and functionalist biology. Journal of Theoretical Biology, 1988, 133, 133-145.	1.7	20
82	Molecular ecology and biological control: the mating system of a marsupial pest. Molecular Ecology, 2000, 9, 723-733.	3.9	20
83	Scientific Prejudice, Reproductive Isolation, and Apartheid. Perspectives in Biology and Medicine, 1984, 28, 107-116.	0.5	18
84	Ancient population genomics and the study of evolution. Philosophical Transactions of the Royal Society B: Biological Sciences, 2015, 370, 20130381.	4.0	18
85	Genetic differences among pheromonally distinct New Zealand leafroller moths. Biochemical Systematics and Ecology, 1994, 22, 329-339.	1.3	17
86	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
87	Resurrecting ancient animal genomes: The extinct moa and more. BioEssays, 2012, 34, 661-669.	2.5	17
88	Archaeogenetics and human evolution: the ontogeny of a biological discipline. World Archaeology, 2019, 51, 546-559.	1.1	17
89	Molecular sexing of individual kakapo, Strigops habroptilus Aves, from faeces. Molecular Ecology, 1999, 8, 1349-1350.	3.9	16
90	Microsatellite DNA markers for tuatara (Sphenodon spp.). Conservation Genetics, 2001, 2, 183-185.	1.5	16

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91	Molecular sexing of the communally breeding pukeko: an important ecological tool. Molecular Ecology, 1996, 5, 289-293.	3.9	16
92	Ancient DNA Suggests Dwarf and â€~Giant' Emu Are Conspecific. PLoS ONE, 2011, 6, e18728.	2.5	16
93	Specific-mate recognition systems, phylogenies and asymmetrical evolution. Journal of Theoretical Biology, 1984, 109, 147-156.	1.7	15
94	Microsatellite primers for the kakapo (Strigops habroptilus) and their utility in other parrots. Conservation Genetics, 2000, 1, 93-95.	1.5	15
95	Mating system and genetic variation in the endangered New Zealand takahe. Conservation Genetics, 2002, 3, 427-434.	1.5	15
96	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
97	Birdstrikes and barcoding: can DNA methods help make the airways safer?. Molecular Ecology Resources, 2011, 11, 38-45.	4.8	15
98	Adélie penguins and temperature changes in Antarctica: a longâ€ŧerm view. Integrative Zoology, 2012, 7, 113-120.	2.6	15
99	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
100	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
101	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
102	Mitogenomic diversity in Sacred Ibis Mummies sheds light on early Egyptian practices. PLoS ONE, 2019, 14, e0223964.	2.5	14
103	Evolutionary Studies of the New Zealand Coastal Mosquito Opifex Fuscus (Hutton) I. Mating Behaviour. Behaviour, 1983, 84, 157-171.	0.8	13
104	Title is missing!. Conservation Genetics, 2000, 1, 103-113.	1.5	13
105	Nuclear microsatellite DNA markers for New Zealand kiwi (Apteryx spp.). Molecular Ecology Notes, 2006, 6, 227-229.	1.7	13
106	Ancient genomics is born. Nature, 2006, 444, 275-276.	27.8	13
107	Next generation sequencing and analysis of a conserved transcriptome of New Zealand's kiwi. BMC Evolutionary Biology, 2010, 10, 387.	3.2	13
108	Mate recognition in members of the Drosophila nasuta complex. Animal Behaviour, 1982, 30, 438-443.	1.9	12

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109	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
110	The Molecular Ecology of the Extinct New Zealand Huia. PLoS ONE, 2009, 4, e8019.	2.5	12
111	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
112	The mysterious Spotted Green Pigeon and its relation to the Dodo and its kindred. BMC Evolutionary Biology, 2014, 14, 136.	3.2	12
113	Limb patterning genes and heterochronic development of the emu wing bud. EvoDevo, 2016, 7, 26.	3.2	12
114	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
115	An evaluation of methods of blood preservation for RT-PCR from endangered species. Conservation Genetics, 2003, 4, 651-654.	1.5	11
116	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
117	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
118	A Population Genetical Study of the African Mosquito Anopheles marshallii (Theobald). Evolution; International Journal of Organic Evolution, 1983, 37, 484.	2.3	10
119	The stability of the specific-mate recognition system ofDrosophila melanogaster. Behavior Genetics, 1986, 16, 369-373.	2.1	10
120	Minisatellite DNA markers reveal hybridisation between the endangered black robin and tomtit. Electrophoresis, 1997, 18, 1682-1687.	2.4	10
121	Mutational Bias in Penguin Microsatellite DNA. Journal of Heredity, 2005, 96, 566-571.	2.4	10
122	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
123	Reconstruction and in vivo analysis of the extinct tbx5 gene from ancient wingless moa (Aves:) Tj ETQq1 1 0.784	314.rgBT 3.2	Oyerlock 10
124	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
125	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
126	DNA barcoding of animal species—response to DeSalle. BioEssays, 2008, 30, 92-93.	2.5	9

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127	Late-Holocene Penguin Occupation and Diet at King George Island Antarctic Peninsula. Antarctic Research Series, 2013, , 171-180.	0.2	9
128	Radiocarbon dating of Sacred Ibis mummies from ancient Egypt. Journal of Archaeological Science: Reports, 2015, 4, 355-361.	0.5	9
129	Ancient DNA Resolves Identity and Phylogeny of New Zealand's Extinct and Living Quail (Coturnix sp.). PLoS ONE, 2009, 4, e6400.	2.5	9
130	Selective Constraints Determine the Time Dependency of Molecular Rates for Human Nuclear Genomes. Genome Biology and Evolution, 2012, 4, 1127-1132.	2.5	8
131	The population genetics of reinforcing selection. Genetica, 1983, 62, 15-23.	1.1	7
132	The mating behavior of individuals ofDrosophila pseudoobscura from New Zealand. Experientia, 1985, 41, 950-952.	1.2	7
133	New Genetic Approach to Detecting Individuals of Rare and Endangered Species. Conservation Biology, 2008, 22, 1267-1276.	4.7	7
134	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
135	"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
136	Sexing the Critically Endangered Kakapo Strigops habroptilus. Emu, 2000, 100, 336-339.	0.6	6
137	Genetic monogamy mirrors social monogamy in the Fiordland crested penguin. New Zealand Journal of Zoology, 2000, 27, 311-316.	1.1	6
138	Genetic identification of moa remains recovered from Tiniroto, Gisborne. Journal of the Royal Society of New Zealand, 2008, 38, 231-235.	1.9	6
139	Molecular evidence for the identity of the Magenta petrel. Molecular Ecology Resources, 2009, 9, 458-461.	4.8	6
140	The Sacred Ibis debate: The first test of evolution. PLoS Biology, 2018, 16, e2005558.	5.6	6
141	Species and Neo-Darwinism. Systematic Zoology, 1990, 39, 399.	1.6	5
142	Ancient genetic variation in one of the world's rarest seabirds. Heredity, 2008, 101, 543-547.	2.6	5
143	Distance-dependent patterns of molecular divergences in tuatara mitogenomes. Scientific Reports, 2015, 5, 8703.	3.3	5
144	Highly Informative Ancient DNA â€~Snippets' for New Zealand Moa. PLoS ONE, 2013, 8, e50732.	2.5	5

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145	Evolutionary Studies of the New Zealand Coastal Mosquito Opifex Fuscus (Hutton) li. Competition for Mates. Behaviour, 1984, 88, 1-12.	0.8	4
146	Laboratory-induced changes in the mate recognition system ofDrosophila pseudoobscura. Behavior Genetics, 1986, 16, 285-294.	2.1	4
147	Genetics ofPotamopyrgus antipodarum(Gastropoda: Prosobranchia): Variation in unisexual populations. New Zealand Journal of Zoology, 1990, 17, 65-72.	1.1	4
148	Genetic continuity within, and discontinuities among, populations of leafroller moths with distinct sex-pheromones. Heredity, 1995, 75, 243-255.	2.6	4
149	Insights Into Aboriginal Australian Mortuary Practices: Perspectives From Ancient DNA. Frontiers in Ecology and Evolution, 2020, 8, .	2.2	4
150	Identification, Classification, and Growth of Moa Chicks (Aves: Dinornithiformes) from the Genus Euryapteryx. PLoS ONE, 2014, 9, e99929.	2.5	4
151	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
152	The sensitive period for yellow phenocopy induction inDrosophila melanogaster. Experientia, 1988, 44, 618-621.	1.2	3
153	A cladistic analysis of species of the molluscan genusPotamopyrgusbased on allozyme data. New Zealand Journal of Zoology, 1990, 17, 257-263.	1.1	3
154	Genetic Relatedness and Alloparental Behaviour in a Captive Group of Spider Monkeys (Ateles) Tj ETQq0 0 0 rgB ⁻	[Qverlocl	۶ 10 Tf 50 38
155	Complex Species Status for Extinct Moa (Aves: Dinornithiformes) from the Genus Euryapteryx. PLoS ONE, 2014, 9, e90212.	2.5	3
156	Organisms not Species Evolve: A Reply to Ghiselin. Systematic Zoology, 1990, 39, 79.	1.6	2
157	Face of the past reconstructed. Nature, 2010, 463, 739-740.	27.8	2
158	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
159	Data sharing: do scientists know best?. Nature, 2017, 548, 281-281.	27.8	2
160	Are BOLD searches scientific? A response to Federhen (2011). Molecular Ecology Resources, 2011, 11, 939-940.	4.8	1
161	Complete mitochondrial genomes of Tuatara endemic to different islands of New Zealand. Mitochondrial DNA, 2015, 26, 25-26.	0.6	1
162	A different paradigm for the colonisation of Sahul. Archaeology in Oceania, 2020, 55, 182-191.	0.7	1

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163	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
164	Industrial Melanism: A Classic Example of Another Kind?. BioScience, 1999, 49, 1021.	4.9	1
165	The †̃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
166	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	0
167	A novel restriction fragment length polymorphism for petrels or tube-nosed seabirds. Molecular Ecology, 2000, 9, 1915-1917.	3.9	0
168	Non-Darwinian Evolution. , 2013, , 87-89.		0
169	What's happening in New Zealand Conservation?. Pacific Conservation Biology, 1995, 2, 1.	1.0	0
170	Editorial. Preface to a new century of «Theoretical Biology Forum». Theoretical Biology Forum, 2018, 111, 9-10.	0.2	0
171	Editorial. Theoretical biology forum. The next century. Theoretical Biology Forum, 2019, 112, 9-10.	0.2	0
172	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