

Andrew J Veale

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6895643/publications.pdf>

Version: 2024-02-01

28
papers

902
citations

687363

13
h-index

580821

25
g-index

30
all docs

30
docs citations

30
times ranked

1692
citing authors

#	ARTICLE	IF	CITATIONS
1	Genomics and the challenging translation into conservation practice. <i>Trends in Ecology and Evolution</i> , 2015, 30, 78-87.	8.7	469
2	An ancient selective sweep linked to reproductive life history evolution in sockeye salmon. <i>Scientific Reports</i> , 2017, 7, 1747.	3.3	44
3	Genomic Changes Associated with Reproductive and Migratory Ecotypes in Sockeye Salmon (<i>Oncorhynchus nerka</i>). <i>Genome Biology and Evolution</i> , 2017, 9, 2921-2939.	2.5	38
4	Using genetic techniques to quantify reinvasion, survival and <i>in situ</i> breeding rates during control operations. <i>Molecular Ecology</i> , 2013, 22, 5071-5083.	3.9	32
5	Phylogeography of the snakeskin chiton <i>Sypharochiton pelliserpentis</i> (Mollusca: Polyplacophora) around New Zealand: are seasonal near-shore upwelling events a dynamic barrier to gene flow?. <i>Biological Journal of the Linnean Society</i> , 2011, 104, 552-563.	1.6	30
6	Ecological gradients drive insect wing loss and speciation: The role of the alpine treeline. <i>Molecular Ecology</i> , 2019, 28, 3141-3150.	3.9	27
7	The population genetic structure of the waratah anemone (<i>Actinia tenebrosa</i>) around New Zealand. <i>New Zealand Journal of Marine and Freshwater Research</i> , 2012, 46, 523-536.	2.0	26
8	Sockeye salmon repatriation leads to population re-establishment and rapid introgression with native kokanee. <i>Evolutionary Applications</i> , 2016, 9, 1301-1311.	3.1	24
9	Genetic population assignment reveals a long-distance incursion to an island by a stoat (<i>Mustela</i>) Tj ETQq1 1 0.784314 rgBT /Overlock 2.4 23	2.4	23
10	The genomic ancestry, landscape genetics and invasion history of introduced mice in New Zealand. <i>Royal Society Open Science</i> , 2018, 5, 170879.	2.4	21
11	Genomics detects population structure within and between ocean basins in a circumpolar seabird: The white-chinned petrel. <i>Molecular Ecology</i> , 2019, 28, 4552-4572.	3.9	21
12	Comparative transcriptomic analysis of a wing-dimorphic stonefly reveals candidate wing loss genes. <i>EvoDevo</i> , 2019, 10, 21.	3.2	18
13	Genotyping-by-sequencing supports a genetic basis for wing reduction in an alpine New Zealand stonefly. <i>Scientific Reports</i> , 2018, 8, 16275.	3.3	17
14	Sap flow of the southern conifer, <i>Agathis australis</i> during wet and dry summers. <i>Trees - Structure and Function</i> , 2016, 30, 19-33.	1.9	15
15	Unwelcome visitors: employing forensic methodologies to inform the stoat (<i>Mustela erminea</i>) incursion response plan on Kapiti Island. <i>New Zealand Journal of Zoology</i> , 2014, 41, 1-9.	1.1	12
16	What can the geographic distribution of mtDNA haplotypes tell us about the invasion of New Zealand by house mice <i>Mus musculus</i> ?. <i>Biological Invasions</i> , 2016, 18, 1551-1565.	2.4	12
17	The complete mitochondrial genomes of two chiton species (<i>Sypharochiton</i>) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 107 To (sequencing. <i>Mitochondrial DNA</i> , 2016, 27, 537-538.	0.6	12
18	Observations of stoats (<i>Mustela erminea</i>) swimming. <i>New Zealand Journal of Zoology</i> , 2013, 40, 166-169.	1.1	11

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19	Swimming capabilities of stoats and the threat to inshore sanctuaries. <i>Biological Invasions</i> , 2014, 16, 987-995.	2.4	9
20	An invasive non-native mammal population conserves genetic diversity lost from its native range. <i>Molecular Ecology</i> , 2015, 24, 2156-2163.	3.9	9
21	Using te reo Māori and ta re Moriori in taxonomy. <i>New Zealand Journal of Ecology</i> , 2019, 43, .	1.1	9
22	The effects of mice on stoats in southern beech forests. <i>Austral Ecology</i> , 2015, 40, 32-39.	1.5	7
23	Population Connectivity and Traces of Mitochondrial Introgression in New Zealand Black-Billed Gulls (<i>Larus bulleri</i>). <i>Genes</i> , 2018, 9, 544.	2.4	6
24	Measuring connectivity of invasive stoat populations to inform conservation management. <i>Wildlife Research</i> , 2014, 41, 395.	1.4	4
25	First complete mitochondrial genome of a Gripopterygid stonefly from the sub-order Antarcoperlaria: <i>Zelandoperla fenestrata</i> . <i>Mitochondrial DNA Part B: Resources</i> , 2019, 4, 886-888.	0.4	2
26	Small mustelids in New Zealand: invasion ecology in a different world. , 2018, , .		1
27	Genetic diversity and differentiation in the leaf litter weevil <i>Geochus politus</i> across an urban-rural gradient. <i>New Zealand Journal of Ecology</i> , 0, , .	1.1	1
28	The introduction and diversity of commensal rodents in 19th century Australasia. <i>Biological Invasions</i> , 0, , 1.	2.4	1