Pierre de Villemereuil

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Common garden experiments to study local adaptation need to account for population structure. Journal of Ecology, 2022, 110, 1005-1009.	4.0	12
2	Phenotypic plasticity drives phenological changes in a Mediterranean blue tit population. Journal of Evolutionary Biology, 2022, 35, 347-359.	1.7	9
3	Disturbance and indirect effects of climate warming support a plant invader in mountains. Oikos, 2022, 2022, .	2.7	3
4	Do leaf nitrogen resorption dynamics align with the slowâ€ f ast continuum? A test at the intraspecific level. Functional Ecology, 2022, 36, 1315-1328.	3.6	6
5	Finding the adaptive needles in a populationâ€structured haystack: A case study in a New Zealand mollusc. Journal of Animal Ecology, 2022, 91, 1209-1221.	2.8	3
6	Genetic variance in fitness indicates rapid contemporary adaptive evolution in wild animals. Science, 2022, 376, 1012-1016.	12.6	69
7	Heritability of a resting heart rate in a 20-year follow-up family cohort with GWAS data: Insights from the STANISLAS cohort. European Journal of Preventive Cardiology, 2021, 28, 1334-1341.	1.8	12
8	Predicting population genetic change in an autocorrelated random environment: Insights from a large automated experiment. PLoS Genetics, 2021, 17, e1009611.	3.5	8
9	Genetic Variations and Differential DNA Methylation to Face Contrasted Climates in Small Ruminants: An Analysis on Traditionally-Managed Sheep and Goats. Frontiers in Genetics, 2021, 12, 745284.	2.3	4
10	Climate dependent heating efficiency in the common lizard. Ecology and Evolution, 2020, 10, 8007-8017.	1.9	9
11	Consequences of space sharing on individual phenotypes in the New Zealand hihi. Evolutionary Ecology, 2020, 34, 821-839.	1.2	5
12	Polygenic basis for adaptive morphological variation in a threatened Aotearoa New Zealand bird, the hihi (<i>Notiomystis cincta</i>). Proceedings of the Royal Society B: Biological Sciences, 2020, 287, 20200948.	2.6	23
13	Hitchhiking consequences for genetic and morphological patterns: the influence of kelp-rafting on a brooding chiton. Biological Journal of the Linnean Society, 2020, 130, 756-770.	1.6	6
14	Accounting for stochasticity in demographic compensation along the elevational range of an alpine plant. Ecology Letters, 2020, 23, 870-880.	6.4	5
15	Fluctuating optimum and temporally variable selection on breeding date in birds and mammals. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 31969-31978.	7.1	69
16	On the relevance of Bayesian statistics and MCMC for animal models. Journal of Animal Breeding and Genetics, 2019, 136, 339-340.	2.0	13
17	Can threatened species adapt in a restored habitat? No expected evolutionary response in lay date for the New Zealand hihi. Evolutionary Applications, 2019, 12, 482-497.	3.1	17
18	Directional selection on body size but no apparent survival cost to being large in wild New Zealand giraffe weevils. Evolution; International Journal of Organic Evolution, 2019, 73, 762-776.	2.3	10

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19	Little Adaptive Potential in a Threatened Passerine Bird. Current Biology, 2019, 29, 889-894.e3.	3.9	53
20	A General Method for Simultaneously Accounting for Phylogenetic and Species Sampling Uncertainty via Rubin's Rules in Comparative Analysis. Systematic Biology, 2019, 68, 632-641.	5.6	33
21	Convergent genomic signatures of domestication in sheep and goats. Nature Communications, 2018, 9, 813.	12.8	220
22	Patterns of phenotypic plasticity and local adaptation in the wide elevation range of the alpine plant <i>Arabis alpina</i> . Journal of Ecology, 2018, 106, 1952-1971.	4.0	65
23	Quantitative genetic methods depending on the nature of the phenotypic trait. Annals of the New York Academy of Sciences, 2018, 1422, 29-47.	3.8	56
24	Fixedâ€effect variance and the estimation of repeatabilities and heritabilities: issues and solutions. Journal of Evolutionary Biology, 2018, 31, 621-632.	1.7	73
25	Perturbations on the uniform distribution of p-values can lead to misleading inferences from null-hypothesis testing. Trends in Neuroscience and Education, 2017, 8-9, 18-27.	3.1	2
26	General Methods for Evolutionary Quantitative Genetic Inference from Generalized Mixed Models. Genetics, 2016, 204, 1281-1294.	2.9	156
27	Kin recognition or phenotype matching?. New Phytologist, 2016, 209, 13-14.	7.3	16
28	Common garden experiments in the genomic era: new perspectives and opportunities. Heredity, 2016, 116, 249-254.	2.6	252
29	A new F _{ST} â€based method to uncover local adaptation using environmental variables. Methods in Ecology and Evolution, 2015, 6, 1248-1258.	5.2	164
30	Detecting adaptive evolution based on association with ecological gradients: Orientation matters!. Heredity, 2015, 115, 22-28.	2.6	76
31	Dispersal evolution and resource matching in a spatially and temporally variable environment. Journal of Theoretical Biology, 2015, 370, 184-196.	1.7	2
32	Genome scan methods against more complex models: when and how much should we trust them?. Molecular Ecology, 2014, 23, 2006-2019.	3.9	265
33	General Quantitative Genetic Methods for Comparative Biology. , 2014, , 287-303.		109
34	Bayesian approaches to the quantitative genetic analysis of natural populations. , 2014, , 228-253.		25
35	Comparing parent–offspring regression with frequentist and Bayesian animal models to estimate heritability in wild populations: a simulation study for Gaussian and binary traits. Methods in Ecology and Evolution, 2013, 4, 260-275.	5.2	139
36	Bayesian models for comparative analysis integrating phylogenetic uncertainty. BMC Evolutionary Biology, 2012, 12, 102.	3.2	87

#	Article	IF	CITATIONS
37	Consumer functional responses under intra- and inter-specific interference competition. Ecological Modelling, 2011, 222, 419-426.	2.5	46