

# Benjamin M Fitzpatrick

## List of Publications by Year in descending order

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Version: 2024-02-01

70  
papers

4,292  
citations

126907

33  
h-index

123424

61  
g-index

71  
all docs

71  
docs citations

71  
times ranked

5391  
citing authors

#	ARTICLE	IF	CITATIONS
1	Amphibious mudskipper populations are genetically connected along coastlines, but differentiated across water. <i>Journal of Biogeography</i> , 2022, 49, 767-779.	3.0	4
2	Breeding behaviour predicts patterns of natural hybridization in North American minnows (Cyprinidae). <i>Journal of Evolutionary Biology</i> , 2021, 34, 486-500.	1.7	6
3	Genetic data reveal fine-scale ecological segregation between larval plethodontid salamanders in replicate contact zones. <i>Evolutionary Ecology</i> , 2021, 35, 309-322.	1.2	5
4	GEOGRAPHIC AND INDIVIDUAL DETERMINANTS OF IMPORTANT AMPHIBIAN PATHOGENS IN HELLBENDERS (CRYPTOBRANCHIUS ALLEGANIENSIS) IN TENNESSEE AND ARKANSAS, USA. <i>Journal of Wildlife Diseases</i> , 2020, 56, 803-814.	0.8	4
5	Extensive Cryptic Diversity Within the <i>Physalaemus cuvieri</i> – <i>Physalaemus ephippifer</i> Species Complex (Amphibia, Anura) Revealed by Cytogenetic, Mitochondrial, and Genomic Markers. <i>Frontiers in Genetics</i> , 2019, 10, 719.	2.3	9
6	Morphological Polymorphism Associated with Alternative Reproductive Tactics in a Plethodontid Salamander. <i>American Naturalist</i> , 2019, 193, 608-618.	2.1	10
7	A hierarchical Bayesian model to incorporate uncertainty into methods for diversity partitioning. <i>Ecology</i> , 2018, 99, 947-956.	3.2	10
8	Isolation by distance, local adaptation, and fortuitous coincidence of geo-political boundaries with spatial-genetic clusters in southern Bog Turtles. <i>Global Ecology and Conservation</i> , 2018, 16, e00474.	2.1	3
9	Population Viability of Nonnative Mediterranean House Geckos ( <i>Hemidactylus turcicus</i> ) at an Urban Site Near the Northern Invasion Front. <i>Journal of Herpetology</i> , 2018, 52, 215.	0.5	9
10	Pairwise beta diversity resolves an underappreciated source of confusion in calculating species turnover. <i>Ecology</i> , 2017, 98, 933-939.	3.2	40
11	Genome scale assessment of a species translocation program. <i>Conservation Genetics</i> , 2017, 18, 1191-1199.	1.5	17
12	Co-occurrence and Hybridization between <i>Necturus maculosus</i> and a Heretofore Unknown <i>Necturus</i> in the Southern Appalachians. <i>Journal of Herpetology</i> , 2017, 51, 559.	0.5	1
13	Hybridization and the species problem in conservation. <i>Environmental Epigenetics</i> , 2015, 61, 206-216.	1.8	74
14	Extending the Concept of Diversity Partitioning to Characterize Phenotypic Complexity. <i>American Naturalist</i> , 2015, 186, 348-361.	2.1	27
15	Symbiote transmission and maintenance of extra-genomic associations. <i>Frontiers in Microbiology</i> , 2014, 5, 46.	3.5	35
16	Similarity and differentiation between bacteria associated with skin of salamanders ( <i>Plethodon</i> )	2.7	64
17	iteRates: An R Package for Implementing a Parametric Rate Comparison on Phylogenetic Trees. <i>Evolutionary Bioinformatics</i> , 2014, 10, EBO.S16487.	1.2	4
18	Doomed before they are described? The need for conservation assessments of cryptic species complexes using an amblyopsid cavefish ( <i>Amblyopsidae</i> : <i>Typhlichthys</i> ) as a case study. <i>Biodiversity and Conservation</i> , 2013, 22, 1799-1820.	2.6	58

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19	Transgressive Hybrids as Hopeful Monsters. <i>Evolutionary Biology</i> , 2013, 40, 310-315.	1.1	128
20	A PARAMETRIC METHOD FOR ASSESSING DIVERSIFICATION-RATE VARIATION IN PHYLOGENETIC TREES. <i>Evolution; International Journal of Organic Evolution</i> , 2013, 67, 368-377.	2.3	11
21	Lethal Effects of Water Quality on Threatened California Salamanders but Not on Co-occurring Hybrid Salamanders. <i>Conservation Biology</i> , 2013, 27, 95-102.	4.7	18
22	EFFECTS OF CLIMATIC AND GEOLOGICAL PROCESSES DURING THE PLEISTOCENE ON THE EVOLUTIONARY HISTORY OF THE NORTHERN CAVEFISH, <i>AMBLYOPSIS SPELAEA</i> (TELEOSTEI: AMBLYOPSIDAE). <i>Evolution; International Journal of Organic Evolution</i> , 2013, 67, 1011-1025.	2.3	33
23	Tests of two methods for identifying founder effects in metapopulations reveal substantial type II error. <i>Genetica</i> , 2013, 141, 119-131.	1.1	9
24	Can genetic data confirm or refute historical records? The island invasion of the small Indian mongoose ( <i>Herpestes auropunctatus</i> ). <i>Biological Invasions</i> , 2013, 15, 2243-2251.	2.4	18
25	EVIDENCE FOR REPEATED LOSS OF SELECTIVE CONSTRAINT IN RHODOPSIN OF AMBLYOPSID CAVEFISHES (TELEOSTEI: AMBLYOPSIDAE). <i>Evolution; International Journal of Organic Evolution</i> , 2013, 67, 732-748.	2.3	82
26	Alternative forms for genomic clines. <i>Ecology and Evolution</i> , 2013, 3, 1951-1966.	1.9	64
27	Successive virgin births of viable male progeny in the checkered gartersnake, <i>Thamnophis marcianus</i> . <i>Biological Journal of the Linnean Society</i> , 2012, 107, 566-572.	1.6	20
28	Estimating ancestry and heterozygosity of hybrids using molecular markers. <i>BMC Evolutionary Biology</i> , 2012, 12, 131.	3.2	119
29	Underappreciated Consequences of Phenotypic Plasticity for Ecological Speciation. <i>International Journal of Ecology</i> , 2012, 2012, 1-12.	0.8	87
30	From genes to ecosystems. , 2012, , 269-286.		10
31	Hybridization between two gartersnake species ( <i>Thamnophis</i> ) of conservation concern: a threat or an important natural interaction?. <i>Conservation Genetics</i> , 2012, 13, 649-663.	1.5	13
32	Genetic analysis of an endemic archipelagic lizard reveals sympatric cryptic lineages and taxonomic discordance. <i>Conservation Genetics</i> , 2012, 13, 953-963.	1.5	4
33	DELIMITING SPECIES USING MULTILOCUS DATA: DIAGNOSING CRYPTIC DIVERSITY IN THE SOUTHERN CAVEFISH, <i>TYPHLICHTHYS SUBTERRANEUS</i> (TELEOSTEI: AMBLYOPSIDAE). <i>Evolution; International Journal of Organic Evolution</i> , 2012, 66, 846-866.	2.3	143
34	What can DNA tell us about biological invasions?. <i>Biological Invasions</i> , 2012, 14, 245-253.	2.4	133
35	Unexpected Shallow Genetic Divergence in Turks Island Boas ( <i>Epicrates c. chrysogaster</i> ) Reveals Single Evolutionarily Significant Unit for Conservation. <i>Herpetologica</i> , 2011, 67, 477-486.	0.4	14
36	Genetic variation and community change – selection, evolution, and feedbacks. <i>Functional Ecology</i> , 2011, 25, 408-419.	3.6	47

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37	Population genetics of the Honduran spiny-tailed iguana <i>Ctenosaura melanosterna</i> : implications for conservation and management. <i>Endangered Species Research</i> , 2011, 14, 113-126.	2.4	7
38	Rapid spread of invasive genes into a threatened native species. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 3606-3610.	7.1	178
39	Gene trees, species and species trees in the <i>Ctenosaura palearis</i> clade. <i>Conservation Genetics</i> , 2010, 11, 1767-1781.	1.5	14
40	Analysis of genetic diversity in flowering dogwood natural stands using microsatellites: the effects of dogwood anthracnose. <i>Genetica</i> , 2010, 138, 1047-1057.	1.1	20
41	Retention of low-fitness genotypes over six decades of admixture between native and introduced tiger salamanders. <i>BMC Evolutionary Biology</i> , 2010, 10, 147.	3.2	37
42	Relatedness and genetic structure in a socially polymorphic population of the spider <i>Anelosimus studiosus</i> . <i>Molecular Ecology</i> , 2010, 19, 810-818.	3.9	24
43	Geography disentangles introgression from ancestral polymorphism in Lake Malawi cichlids. <i>Molecular Ecology</i> , 2010, 19, 940-951.	3.9	65
44	Patterns of differential introgression in a salamander hybrid zone: inferences from genetic data and ecological niche modelling. <i>Molecular Ecology</i> , 2010, 19, 4265-4282.	3.9	46
45	Population differences in behaviour are explained by shared within-population trait correlations. <i>Journal of Evolutionary Biology</i> , 2010, 23, 748-756.	1.7	68
46	Invasive hybrid tiger salamander genotypes impact native amphibians. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 11166-11171.	7.1	100
47	Gene flow between an endangered endemic iguana, and its wide spread relative, on the island of Utila, Honduras: when is hybridization a threat?. <i>Conservation Genetics</i> , 2009, 10, 1247-1254.	1.5	22
48	Frequency-dependent selection by wild birds promotes polymorphism in model salamanders. <i>BMC Ecology</i> , 2009, 9, 12.	3.0	46
49	Power and sample size for nested analysis of molecular variance. <i>Molecular Ecology</i> , 2009, 18, 3961-3966.	3.9	82
50	Pattern, process and geographic modes of speciation. <i>Journal of Evolutionary Biology</i> , 2009, 22, 2342-2347.	1.7	142
51	Rapid fixation of non-native alleles revealed by genome-wide SNP analysis of hybrid tiger salamanders. <i>BMC Evolutionary Biology</i> , 2009, 9, 176.	3.2	75
52	Hybrid Dysfunction: Population Genetic and Quantitative Genetic Perspectives. <i>American Naturalist</i> , 2008, 171, 491-498.	2.1	21
53	Dobzhansky's Muller model of hybrid dysfunction supported by poor burst-speed performance in hybrid tiger salamanders. <i>Journal of Evolutionary Biology</i> , 2008, 21, 342-351.	1.7	15
54	What, if anything, is sympatric speciation?. <i>Journal of Evolutionary Biology</i> , 2008, 21, 1452-1459.	1.7	188

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55	Recent divergence with gene flow in Tennessee cave salamanders ( <i>Plethodontidae</i> : <i>Gyrinophilus</i> ) inferred from gene genealogies. <i>Molecular Ecology</i> , 2008, 17, 2258-2275.	3.9	218
56	Distinctiveness in the face of gene flow: hybridization between specialist and generalist gartersnakes. <i>Molecular Ecology</i> , 2008, 17, 4107-4117.	3.9	42
57	Hybrid vigor between native and introduced salamanders raises new challenges for conservation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 15793-15798.	7.1	141
58	Sympatric Speciation: Models and Empirical Evidence. <i>Annual Review of Ecology, Evolution, and Systematics</i> , 2007, 38, 459-487.	8.3	624
59	INTRODUCTION HISTORY AND HABITAT VARIATION EXPLAIN THE LANDSCAPE GENETICS OF HYBRID TIGER SALAMANDERS. , 2007, 17, 598-608.		55
60	ASSORTATIVE MATING IN POISON-DART FROGS BASED ON AN ECOLOGICALLY IMPORTANT TRAIT. <i>Evolution; International Journal of Organic Evolution</i> , 2007, 61, 2253-2259.	2.3	141
61	THE GEOGRAPHY OF MAMMALIAN SPECIATION: MIXED SIGNALS FROM PHYLOGENIES AND RANGE MAPS. <i>Evolution; International Journal of Organic Evolution</i> , 2006, 60, 601.	2.3	16
62	THE GEOGRAPHY OF MAMMALIAN SPECIATION: MIXED SIGNALS FROM PHYLOGENIES AND RANGE MAPS. <i>Evolution; International Journal of Organic Evolution</i> , 2006, 60, 601-615.	2.3	161
63	RATES OF EVOLUTION OF HYBRID INVIABILITY IN BIRDS AND MAMMALS. <i>Evolution; International Journal of Organic Evolution</i> , 2004, 58, 1865.	2.3	6
64	ENVIRONMENT-DEPENDENT ADMIXTURE DYNAMICS IN A TIGER SALAMANDER HYBRID ZONE. <i>Evolution; International Journal of Organic Evolution</i> , 2004, 58, 1282.	2.3	8
65	RATES OF EVOLUTION OF HYBRID INVIABILITY IN BIRDS AND MAMMALS. <i>Evolution; International Journal of Organic Evolution</i> , 2004, 58, 1865-1870.	2.3	127
66	ENVIRONMENT-DEPENDENT ADMIXTURE DYNAMICS IN A TIGER SALAMANDER HYBRID ZONE. <i>Evolution; International Journal of Organic Evolution</i> , 2004, 58, 1282-1293.	2.3	48
67	Morphology and escape performance of tiger salamander larvae ( <i>Ambystoma tigrinum mavortium</i> ). <i>The Journal of Experimental Zoology</i> , 2003, 297A, 147-159.	1.4	34
68	HYBRIDIZATION BETWEEN A RARE, NATIVE TIGER SALAMANDER ( <i>AMBYSTOMA CALIFORNIENSE</i> ) AND ITS INTRODUCED CONGENER. , 2003, 13, 1263-1275.		109
69	MOLECULAR CORRELATES OF REPRODUCTIVE ISOLATION. <i>Evolution; International Journal of Organic Evolution</i> , 2002, 56, 191.	2.3	7
70	MOLECULAR CORRELATES OF REPRODUCTIVE ISOLATION. <i>Evolution; International Journal of Organic Evolution</i> , 2002, 56, 191-198.	2.3	64