

Kanchon K Dasmahapatra

List of Publications by Year in descending order

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Version: 2024-02-01

29
papers

3,212
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430754

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times ranked

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citing authors

#	ARTICLE	IF	CITATIONS
1	Complex basis of hybrid female sterility and Haldane's rule in <i>Heliconius</i> butterflies: Z-linkage and epistasis. <i>Molecular Ecology</i> , 2022, 31, 959-977.	2.0	10
2	Demographic Reconstruction of Antarctic Fur Seals Supports the Krill Surplus Hypothesis. <i>Genes</i> , 2022, 13, 541.	1.0	13
3	Anthropogenic pressures coincide with Neotropical biodiversity hotspots in a flagship butterfly group. <i>Diversity and Distributions</i> , 2022, 28, 2912-2930.	1.9	18
4	The Amazon river is a suture zone for a polyphyletic group of mimetic heliconiine butterflies. <i>Ecography</i> , 2021, 44, 177-187.	2.1	9
5	Deep Convergence, Shared Ancestry, and Evolutionary Novelty in the Genetic Architecture of <i>Heliconius</i> Mimicry. <i>Genetics</i> , 2020, 216, 765-780.	1.2	13
6	Adaptation of the carbamoyl-phosphate synthetase enzyme in an extremophile fish. <i>Royal Society Open Science</i> , 2020, 7, 201200.	1.1	5
7	Past, current, and potential future distributions of unique genetic diversity in a cold-adapted mountain butterfly. <i>Ecology and Evolution</i> , 2020, 10, 11155-11168.	0.8	15
8	Exploring the Expression of Cardiac Regulators in a Vertebrate Extremophile: The Cichlid Fish <i>Oreochromis (Alcolapia) alcalica</i> . <i>Journal of Developmental Biology</i> , 2020, 8, 22.	0.9	2
9	Cryptic speciation associated with geographic and ecological divergence in two Amazonian <i>Heliconius</i> butterflies. <i>Zoological Journal of the Linnean Society</i> , 2019, 186, 233-249.	1.0	15
10	Geographic contrasts between pre- and postzygotic barriers are consistent with reinforcement in <i>Heliconius</i> butterflies. <i>Evolution; International Journal of Organic Evolution</i> , 2019, 73, 1821-1838.	1.1	22
11	Genomic architecture and introgression shape a butterfly radiation. <i>Science</i> , 2019, 366, 594-599.	6.0	365
12	The genetic architecture of adaptation: convergence and pleiotropy in <i>Heliconius</i> wing pattern evolution. <i>Heredity</i> , 2019, 123, 138-152.	1.2	28
13	Contrasting geographic structure in evolutionarily divergent Lake Tanganyika catfishes. <i>Ecology and Evolution</i> , 2018, 8, 2688-2697.	0.8	4
14	Supergene Evolution Triggered by the Introgression of a Chromosomal Inversion. <i>Current Biology</i> , 2018, 28, 1839-1845.e3.	1.8	130
15	RAD Sequencing and a Hybrid Antarctic Fur Seal Genome Assembly Reveal Rapidly Decaying Linkage Disequilibrium, Global Population Structure and Evidence for Inbreeding. <i>Genetics</i> , 2018, 8, 2709-2722.	0.8	30
16	The Scent Chemistry of <i>Heliconius</i> Wing Androconia. <i>Journal of Chemical Ecology</i> , 2017, 43, 843-857.	0.9	36
17	Niche divergence facilitated by fine-scale ecological partitioning in a recent cichlid fish adaptive radiation. <i>Evolution; International Journal of Organic Evolution</i> , 2016, 70, 2718-2735.	1.1	38
18	Genome-wide introgression among distantly related <i>Heliconius</i> butterfly species. <i>Genome Biology</i> , 2016, 17, 25.	3.8	115

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19	Major Improvements to the <i>Heliconius melpomene</i> Genome Assembly Used to Confirm 10 Chromosome Fusion Events in 6 Million Years of Butterfly Evolution. <i>G3: Genes, Genomes, Genetics</i> , 2016, 6, 695-708.	0.8	149
20	Evolutionary Novelty in a Butterfly Wing Pattern through Enhancer Shuffling. <i>PLoS Biology</i> , 2016, 14, e1002353.	2.6	136
21	Estimation of the Spontaneous Mutation Rate in <i>Heliconius melpomene</i> . <i>Molecular Biology and Evolution</i> , 2015, 32, 239-243.	3.5	220
22	Multilocus Species Trees Show the Recent Adaptive Radiation of the Mimetic <i>Heliconius</i> Butterflies. <i>Systematic Biology</i> , 2015, 64, 505-524.	2.7	204
23	High levels of interspecific gene flow in an endemic cichlid fish adaptive radiation from an extreme lake environment. <i>Molecular Ecology</i> , 2015, 24, 3421-3440.	2.0	53
24	High-throughput sequencing reveals inbreeding depression in a natural population. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 3775-3780.	3.3	221
25	Genome-wide evidence for speciation with gene flow in <i>Heliconius</i> butterflies. <i>Genome Research</i> , 2013, 23, 1817-1828.	2.4	609
26	Genome-wide patterns of divergence and gene flow across a butterfly radiation. <i>Molecular Ecology</i> , 2013, 22, 814-826.	2.0	160
27	Genomic islands of divergence in hybridizing <i>Heliconius</i> butterflies identified by large-scale targeted sequencing. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2012, 367, 343-353.	1.8	294
28	The anatomy of a "suture zone"™ in Amazonian butterflies: a coalescent-based test for vicariant geographic divergence and speciation. <i>Molecular Ecology</i> , 2010, 19, 4283-4301.	2.0	68
29	Mitochondrial DNA barcoding detects some species that are real, and some that are not. <i>Molecular Ecology Resources</i> , 2010, 10, 264-273.	2.2	119