

Christopher W Wheat

List of Publications by Year in descending order

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Version: 2024-02-01

67
papers

4,235
citations

186265
28
h-index

133252
59
g-index

77
all docs

77
docs citations

77
times ranked

6411
citing authors

#	ARTICLE	IF	CITATIONS
1	A genetic switch for male UV iridescence in an incipient species pair of sulphur butterflies. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2022, 119, .	7.1	22
2	Local adaptation of life cycles in a butterfly is associated with variation in several circadian clock genes. <i>Molecular Ecology</i> , 2022, 31, 1461-1475.	3.9	8
3	Alternative splicing in seasonal plasticity and the potential for adaptation to environmental change. <i>Nature Communications</i> , 2022, 13, 755.	12.8	18
4	The unresolved phylogenomic tree of butterflies and moths (Lepidoptera): Assessing the potential causes and consequences. <i>Systematic Entomology</i> , 2022, 47, 531-550.	3.9	14
5	A large and diverse autosomal haplotype is associated with sex-linked colour polymorphism in the guppy. <i>Nature Communications</i> , 2022, 13, 1233.	12.8	3
6	Extensive transcriptomic profiling of pupal diapause in a butterfly reveals a dynamic phenotype. <i>Molecular Ecology</i> , 2022, 31, 1269-1280.	3.9	7
7	Planned cull endangers Swedish wolf population. <i>Science</i> , 2022, 377, 162-162.	12.6	5
8	A Population Genomic Investigation of Immune Cell Diversity and Phagocytic Capacity in a Butterfly. <i>Genes</i> , 2021, 12, 279.	2.4	5
9	The Genome of the Margined White Butterfly (<i>Pieris macdunnoughii</i>): Sex Chromosome Insights and the Power of Polishing with PoolSeq Data. <i>Genome Biology and Evolution</i> , 2021, 13, .	2.5	7
10	Chromosome Level Assembly of the Comma Butterfly (<i>Polygonia c-album</i>). <i>Genome Biology and Evolution</i> , 2021, 13, .	2.5	8
11	Genomic insights into the conservation status of the world's last remaining Sumatran rhinoceros populations. <i>Nature Communications</i> , 2021, 12, 2393.	12.8	39
12	The Genome Assembly and Annotation of the Apollo Butterfly (<i>Parnassius apollo</i>), a Flagship Species for Conservation Biology. <i>Genome Biology and Evolution</i> , 2021, 13, .	2.5	19
13	High-Quality Genome Assembly and Comprehensive Transcriptome of the Painted Lady Butterfly (<i>Vanessa cardui</i>). <i>Genome Biology and Evolution</i> , 2021, 13, .	2.5	10
14	Museomics of a rare taxon: placing Whalleyanidae in the Lepidoptera Tree of Life. <i>Systematic Entomology</i> , 2021, 46, 926-937.	3.9	17
15	Population genomics of the critically endangered <i>Phaenocarpa</i> . <i>Cell Genomics</i> , 2021, 1, 100002.	6.5	106
16	<i>Drosophila</i> Evolution over Space and Time (DEST): A New Population Genomics Resource. <i>Molecular Biology and Evolution</i> , 2021, 38, 5782-5805.	8.9	37
17	Conserved ancestral tropical niche but different continental histories explain the latitudinal diversity gradient in brush-footed butterflies. <i>Nature Communications</i> , 2021, 12, 5717.	12.8	33
18	From Inquilines to Gall Inducers: Genomic Signature of a Life-Style Transition in <i>Synergus</i> Gall Wasps. <i>Genome Biology and Evolution</i> , 2020, 12, 2060-2073.	2.5	9

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19	Butterfly dichromatism primarily evolved via Darwin's, not Wallace's, model. <i>Evolution Letters</i> , 2020, 4, 545-555.	3.3	16
20	Changes in gene expression during female reproductive development in a color polymorphic insect. <i>Evolution; International Journal of Organic Evolution</i> , 2020, 74, 1063-1081.	2.3	13
21	Morphology does not covary with predicted behavioral correlations of the domestication syndrome in dogs. <i>Evolution Letters</i> , 2020, 4, 189-199.	3.3	13
22	Differential Expression of Immune Genes between Two Closely Related Beetle Species with Different Immunocompetence following Attack by <i>Asecodes parviclava</i> . <i>Genome Biology and Evolution</i> , 2020, 12, 522-534.	2.5	6
23	Physiological Tradeoffs of Immune Response Differs by Infection Type in <i>Pieris napi</i> . <i>Frontiers in Physiology</i> , 2020, 11, 576797.	2.8	4
24	A complete time-calibrated multi-gene phylogeny of the European butterflies. <i>ZooKeys</i> , 2020, 938, 97-124.	1.1	61
25	Unprecedented reorganization of holocentric chromosomes provides insights into the enigma of lepidopteran chromosome evolution. <i>Science Advances</i> , 2019, 5, eaau3648.	10.3	66
26	Colour lightness of butterfly assemblages across North America and Europe. <i>Scientific Reports</i> , 2019, 9, 1760.	3.3	32
27	Exploring a Pool-Seq approach for gaining population genomic insights in nonmodel species. <i>Ecology and Evolution</i> , 2019, 9, 11448-11463.	1.9	23
28	Diapause: Circadian Clock Genes Are at It Again. <i>Current Biology</i> , 2019, 29, R1245-R1246.	3.9	20
29	A transposable element insertion is associated with an alternative life history strategy. <i>Nature Communications</i> , 2019, 10, 5757.	12.8	41
30	Climate-induced phenology shifts linked to range expansions in species with multiple reproductive cycles per year. <i>Nature Communications</i> , 2019, 10, 4455.	12.8	82
31	Local adaptation of photoperiodic plasticity maintains life cycle variation within latitudes in a butterfly. <i>Ecology</i> , 2019, 100, e02550.	3.2	46
32	Metabolome dynamics of diapause in the butterfly <i>Pieris napi</i> : distinguishing maintenance, termination and post-diapause phases. <i>Journal of Experimental Biology</i> , 2018, 221, .	1.7	25
33	Strong phenotypic plasticity limits potential for evolutionary responses to climate change. <i>Nature Communications</i> , 2018, 9, 1005.	12.8	137
34	Physiological differences between female limited, alternative life history strategies: The Alba phenotype in the butterfly <i>Colias croceus</i> . <i>Journal of Insect Physiology</i> , 2018, 107, 257-264.	2.0	6
35	Embracing Colonizations: A New Paradigm for Species Association Dynamics. <i>Trends in Ecology and Evolution</i> , 2018, 33, 4-14.	8.7	94
36	Genetics of dispersal. <i>Biological Reviews</i> , 2018, 93, 574-599.	10.4	182

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37	Unifying host-associated diversification processes using butterfly-plant networks. <i>Nature Communications</i> , 2018, 9, 5155.	12.8	35
38	Microevolutionary selection dynamics acting on immune genes of the green-veined white butterfly, <i>Pieris napi</i> . <i>Molecular Ecology</i> , 2018, 27, 2807-2822.	3.9	9
39	The molecular genetic basis of herbivory between butterflies and their host plants. <i>Nature Ecology and Evolution</i> , 2018, 2, 1418-1427.	7.8	56
40	Genetic variation underlying local adaptation of diapause induction along a cline in a butterfly. <i>Molecular Ecology</i> , 2018, 27, 3613-3626.	3.9	67
41	Female fecundity variation affects reproducibility of experiments on host plant preference and acceptance in a phytophagous insect. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2017, 284, 20162643.	2.6	9
42	Timing of diapause termination in relation to variation in winter climate. <i>Physiological Entomology</i> , 2017, 42, 232-238.	1.5	53
43	A high-coverage draft genome of the mycalesine butterfly <i>Bicyclus anynana</i> . <i>GigaScience</i> , 2017, 6, 1-7.	6.4	55
44	Structural complexity and molecular heterogeneity of a butterfly ejaculate reflect a complex history of selection. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, E5406-E5413.	7.1	37
45	Investigating the genomic basis of discrete phenotypes using a Pool-Seq only approach: New insights into the genetics underlying colour variation in diverse taxa. <i>Molecular Ecology</i> , 2017, 26, 4990-5002.	3.9	27
46	Sex-linked inheritance of diapause induction in the butterfly <i>Pieris napi</i> . <i>Physiological Entomology</i> , 2017, 42, 257-265.	1.5	33
47	Bioinformatic processing of RAD-seq data dramatically impacts downstream population genetic inference. <i>Methods in Ecology and Evolution</i> , 2017, 8, 907-917.	5.2	253
48	Transcriptome sequencing reveals high isoform diversity in the ant <i>Formica exsecta</i> . <i>PeerJ</i> , 2017, 5, e3998.	2.0	7
49	Energy and lipid metabolism during direct and diapause development in a pierid butterfly. <i>Journal of Experimental Biology</i> , 2016, 219, 3049-3060.	1.7	64
50	Evolutionary history of host use, rather than plant phylogeny, determines gene expression in a generalist butterfly. <i>BMC Evolutionary Biology</i> , 2016, 16, 59.	3.2	36
51	PCR primers for 30 novel gene regions in the nuclear genomes of Lepidoptera. <i>ZooKeys</i> , 2016, 596, 129-141.	1.1	24
52	The butterfly plant arms-race escalated by gene and genome duplications. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 8362-8366.	7.1	458
53	Genomics and the challenging translation into conservation practice. <i>Trends in Ecology and Evolution</i> , 2015, 30, 78-87.	8.7	469
54	Pgi: the ongoing saga of a candidate gene. <i>Current Opinion in Insect Science</i> , 2014, 4, 42-47.	4.4	12

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55	The Glanville fritillary genome retains an ancient karyotype and reveals selective chromosomal fusions in Lepidoptera. <i>Nature Communications</i> , 2014, 5, 4737.	12.8	196
56	Revised systematics and higher classification of pierid butterflies (Lepidoptera: Pieridae) based on molecular data. <i>Zoologica Scripta</i> , 2014, 43, 641-650.	1.7	61
57	Critiquing blind dating: the dangers of over-confident date estimates in comparative genomics. <i>Trends in Ecology and Evolution</i> , 2013, 28, 636-642.	8.7	24
58	Mechanisms of macroevolution: polyphagous plasticity in butterfly larvae revealed by RNA-seq. <i>Molecular Ecology</i> , 2013, 22, 4884-4895.	3.9	101
59	Phylogenomic Insights into the Cambrian Explosion, the Colonization of Land and the Evolution of Flight in Arthropoda. <i>Systematic Biology</i> , 2013, 62, 93-109.	5.6	75
60	Timing and Patterns in the Taxonomic Diversification of Lepidoptera (Butterflies and Moths). <i>PLoS ONE</i> , 2013, 8, e80875.	2.5	197
61	Quantitative RNA-Seq analysis in non-model species: assessing transcriptome assemblies as a scaffold and the utility of evolutionary divergent genomic reference species. <i>BMC Genomics</i> , 2012, 13, 361.	2.8	79
62	Phosphoglucose isomerase (Pgi) performance and fitness effects among Arthropods and its potential role as an adaptive marker in conservation genetics. <i>Conservation Genetics</i> , 2010, 11, 387-397.	1.5	30
63	Microevolutionary dynamics of a macroevolutionary key innovation in a Lepidopteran herbivore. <i>BMC Evolutionary Biology</i> , 2010, 10, 60.	3.2	17
64	INTEGRATING EVOLUTIONARY AND FUNCTIONAL APPROACHES TO INFER ADAPTATION AT SPECIFIC LOCI. <i>Evolution; International Journal of Organic Evolution</i> , 2010, 64, 2489-2509.	2.3	103
65	A mitochondrial-DNA-based phylogeny for some evolutionary-genetic model species of <i>Colias</i> butterflies (Lepidoptera, Pieridae). <i>Molecular Phylogenetics and Evolution</i> , 2008, 47, 893-902.	2.7	37
66	X-tox: An atypical defensin derived family of immune-related proteins specific to Lepidoptera. <i>Developmental and Comparative Immunology</i> , 2008, 32, 575-584.	2.3	24
67	The genetic basis of a plant-insect coevolutionary key innovation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 20427-20431.	7.1	325