Chris D Jiggins

List of Publications by Year
in descending order

153 papers	15,743 citations	20817 60 h-index	23533 111 g-index
188 all docs	188 docs citations	188 times ranked	11518 citing authors

#	Article	IF	Citations
1	Genomics and the origin of species. Nature Reviews Genetics, 2014, 15, 176-192.	16.3	850
2	Reproductive isolation caused by colour pattern mimicry. Nature, 2001, 411, 302-305.	27.8	611
3	Genome-wide evidence for speciation with gene flow in<i>Heliconius</i>butterflies. Genome Research, 2013, 23, 1817-1828.	5.5	609
4	Bimodal hybrid zones and speciation. Trends in Ecology and Evolution, 2000, 15, 250-255.	8.7	538
5	Evaluating the Use of ABBAâ€“BABA Statistics to Locate Introgressed Loci. Molecular Biology and Evolution, 2015, 32, 244-257.	8.9	532
6	Chromosomal rearrangements maintain a polymorphic supergene controlling butterfly mimicry. Nature, 2011, 477, 203-206.	27.8	509
7	The biology of color. Science, 2017, 357, .	12.6	509
8	<i>optix</i> Drives the Repeated Convergent Evolution of Butterfly Wing Pattern Mimicry. Science, 2011, 333, 1137-1141.	12.6	431
9	Speciation by hybridization in Heliconius butterflies. Nature, 2006, 441, 868-871.	27.8	412
10	Genomic architecture and introgression shape a butterfly radiation. Science, 2019, 366, 594-599.	12.6	365
11	Adaptive Introgression across Species Boundaries in Heliconius Butterflies. PLoS Genetics, 2012, 8, e1002752.	3.5	319
12	Genomic islands of divergence in hybridizing <i>Heliconius</i> butterflies identified by large-scale targeted sequencing. Philosophical Transactions of the Royal Society B: Biological Sciences, 2012, 367, 343-353.	4.0	294
13	Linkage Mapping and Comparative Genomics Using Next-Generation RAD Sequencing of a Non-Model Organism. PLoS ONE, 2011, 6, e19315.	2.5	270
14	Recombination rate variation shapes barriers to introgression across butterfly genomes. PLoS Biology, 2019, 17, e2006288.	5.6	253
15	Diversification of complex butterfly wing patterns by repeated regulatory evolution of a <i>Wnt</i> ligand. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 12632-12637.	7.1	244
16	A Conserved Supergene Locus Controls Colour Pattern Diversity in Heliconius Butterflies. PLoS Biology, 2006, 4, e303.	5.6	242
17	Parallel Evolution of <i>Bacillus thuringiensis</i> Toxin Resistance in Lepidoptera. Genetics, 2011, 189, 675-679.	2.9	239
18	Limited performance of DNA barcoding in a diverse community of tropical butterflies. Proceedings of the Royal Society B: Biological Sciences, 2007, 274, 2881-2889.	2.6	233

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19	Estimation of the Spontaneous Mutation Rate in Heliconius melpomene. Molecular Biology and Evolution, 2015, 32, 239-243.	8.9	220
20	The gene cortex controls mimicry and crypsis in butterflies and moths. Nature, 2016, 534, 106-110.	27.8	212
21	Multilocus Species Trees Show the Recent Adaptive Radiation of the Mimetic Heliconius Butterflies. Systematic Biology, 2015, 64, 505-524.	5.6	204
22	Disruptive sexual selection against hybrids contributes to speciation between Heliconius cydno and Heliconius melpomene. Proceedings of the Royal Society B: Biological Sciences, 2001, 268, 1849-1854.	2.6	189
23	Interpreting the genomic landscape of introgression. Current Opinion in Genetics and Development, 2017, 47, 69-74.	3.3	186
24	Complex modular architecture around a simple toolkit of wing pattern genes. Nature Ecology and Evolution, 2017, 1, 52.	7.8	179
25	Genomeâ€wide patterns of divergence and gene flow across a butterfly radiation. Molecular Ecology, 2013, 22, 814-826.	3.9	160
26	Phylogenetic Discordance at the Species Boundary: Comparative Gene Genealogies Among Rapidly Radiating Heliconius Butterflies. Molecular Biology and Evolution, 2002, 19, 2176-2190.	8.9	156
27	Female Behaviour Drives Expression and Evolution of Gustatory Receptors in Butterflies. PLoS Genetics, 2013, 9, e1003620.	3.5	154
28	Major Improvements to the <i>Heliconius melpomene</i> Genome Assembly Used to Confirm 10 Chromosome Fusion Events in 6ÂMillion Years of Butterfly Evolution. G3: Genes, Genomes, Genetics, 2016, 6, 695-708.	1.8	149
29	Disruptive ecological selection on a mating cue. Proceedings of the Royal Society B: Biological Sciences, 2012, 279, 4907-4913.	2.6	143
30	Do pollen feeding, pupal-mating and larval gregariousness have a single origin in Heliconius butterflies? Inferences from multilocus DNA sequence data. Biological Journal of the Linnean Society, 2007, 92, 221-239.	1.6	138
31	Macroevolutionary shifts of <i>WntA</i> function potentiate butterfly wing-pattern diversity. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, 10701-10706.	7.1	137
32	Evolutionary Novelty in a Butterfly Wing Pattern through Enhancer Shuffling. PLoS Biology, 2016, 14, e1002353.	5.6	136
33	A golden age for evolutionary genetics? Genomic studies of adaptation in natural populations. Trends in Genetics, 2010, 26, 484-492.	6.7	127
34	Ecological Speciation in Mimetic Butterflies. BioScience, 2008, 58, 541-548.	4.9	119
35	Towards the identification of the loci of adaptive evolution. Methods in Ecology and Evolution, 2015, 6, 445-464.	5.2	115
36	Evolution of the Insect Yellow Gene Family. Molecular Biology and Evolution, 2011, 28, 257-272.	8.9	114

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37	Population genomics of parallel hybrid zones in the mimetic butterflies, <i>H. melpomene</i> and <i>H. erato</i>. Genome Research, 2014, 24, 1316-1333.	5.5	114
38	Polyphyly and gene flow between non-sibling Heliconius species. BMC Biology, 2006, 4, 11.	3.8	113
39	Mimicry: developmental genes that contribute to speciation. Evolution & Development, 2003, 5, 269-280.	2.0	112
40	A Genetic Linkage Map of the Mimetic Butterfly Heliconius melpomene. Genetics, 2005, 171, 557-570.	2.9	111
41	Hybrid Sterility, Haldane's Rule and Speciation in <i>Heliconius cydno</i> and <i>H. melpomene</i>. Genetics, 2002, 161, 1517-1526.	2.9	111
42	Mis-Spliced Transcripts of Nicotinic Acetylcholine Receptor Î±6 Are Associated with Field Evolved Spinosad Resistance in Plutella xylostella (L.). PLoS Genetics, 2010, 6, e1000802.	3.5	110
43	Hybrid trait speciation and<i>Heliconius</i>butterflies. Philosophical Transactions of the Royal Society B: Biological Sciences, 2008, 363, 3047-3054.	4.0	108
44	Pervasive genetic associations between traits causing reproductive isolation in<i>Heliconius</i>butterflies. Proceedings of the Royal Society B: Biological Sciences, 2011, 278, 511-518.	2.6	106
45	Wing patterning gene redefines the mimetic history of <i>Heliconius</i> butterflies. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 19666-19671.	7.1	104
46	Synteny and Chromosome Evolution in the Lepidoptera: Evidence From Mapping in <i>Heliconius melpomene</i>. Genetics, 2007, 177, 417-426.	2.9	101
47	MATE PREFERENCE ACROSS THE SPECIATION CONTINUUM IN A CLADE OF MIMETIC BUTTERFLIES. Evolution; International Journal of Organic Evolution, 2011, 65, 1489-1500.	2.3	101
48	Genomic Hotspots for Adaptation: The Population Genetics of MÃ¼llerian Mimicry in Heliconius erato. PLoS Genetics, 2010, 6, e1000796.	3.5	99
49	SEX-LINKED HYBRID STERILITY IN A BUTTERFLY. Evolution; International Journal of Organic Evolution, 2001, 55, 1631-1638.	2.3	98
50	Patterns of pollen feeding and habitat preference among Heliconius species. Ecological Entomology, 2002, 27, 448-456.	2.2	97
51	Genomic Hotspots for Adaptation: The Population Genetics of MÃ¼llerian Mimicry in the Heliconius melpomene Clade. PLoS Genetics, 2010, 6, e1000794.	3.5	97
52	ASSORTATIVE MATING PREFERENCES AMONG HYBRIDS OFFERS A ROUTE TO HYBRID SPECIATION. Evolution; International Journal of Organic Evolution, 2009, 63, 1660-1665.	2.3	96
53	patternize: An R package for quantifying colour pattern variation. Methods in Ecology and Evolution, 2018, 9, 390-398.	5.2	96
54	Natural Selection and Genetic Diversity in the Butterfly <i>Heliconius melpomene</i>. Genetics, 2016, 203, 525-541.	2.9	94

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55	ButterflyBase: a platform for lepidopteran genomics. Nucleic Acids Research, 2007, 36, D582-D587.	14.5	90
56	Genetic Evidence for Hybrid Trait Speciation in Heliconius Butterflies. PLoS Genetics, 2010, 6, e1000930.	3.5	90
57	No evidence for maintenance of a sympatric<i>Heliconius</i>species barrier by chromosomal inversions. Evolution Letters, 2017, 1, 138-154.	3.3	90
58	The genomics of coloration provides insights into adaptive evolution. Nature Reviews Genetics, 2020, 21, 461-475.	16.3	88
59	Convergent Evolution in the Genetic Basis of MuÌˆllerian Mimicry in Heliconius Butterflies. Genetics, 2008, 180, 1567-1577.	2.9	79
60	Genetic dissection of assortative mating behavior. PLoS Biology, 2019, 17, e2005902.	5.6	79
61	Male sex pheromone components in <i> Heliconius</i> butterflies released by the androconia affect female choice. PeerJ, 2017, 5, e3953.	2.0	79
62	The genetic basis of an adaptive radiation: warning colour in two Heliconius species. Proceedings of the Royal Society B: Biological Sciences, 1997, 264, 1167-1175.	2.6	78
63	What can hybrid zones tell us about speciation? The case of<i>Heliconius erato</i>and<i>H. himera</i>(Lepidoptera: Nymphalidae). Biological Journal of the Linnean Society, 1996, 59, 221-242.	1.6	76
64	Evolution of novel mimicry rings facilitated by adaptive introgression in tropical butterflies. Molecular Ecology, 2017, 26, 5160-5172.	3.9	70
65	Patterns of Z chromosome divergence among <i>Heliconius</i> species highlight the importance of historical demography. Molecular Ecology, 2018, 27, 3852-3872.	3.9	69
66	Waiting in the wings: what can we learn about gene co-option from the diversification of butterfly wing patterns?. Philosophical Transactions of the Royal Society B: Biological Sciences, 2017, 372, 20150485.	4.0	67
67	THE PHYLOGENETIC PATTERN OF SPECIATION AND WING PATTERN CHANGE IN NEOTROPICALITHOMIABUTTERFLIES (LEPIDOPTERA: NYMPHALIDAE). Evolution; International Journal of Organic Evolution, 2006, 60, 1454-1466.	2.3	64
68	Adaptive Introgression across Semipermeable Species Boundaries between Local Helicoverpa zea and Invasive Helicoverpa armigera Moths. Molecular Biology and Evolution, 2020, 37, 2568-2583.	8.9	64
69	Maintaining mimicry diversity: optimal warning colour patterns differ among microhabitats in Amazonian clearwing butterflies. Proceedings of the Royal Society B: Biological Sciences, 2017, 284, 20170744.	2.6	60
70	Selective sweeps on novel and introgressed variation shape mimicry loci in a butterfly adaptive radiation. PLoS Biology, 2020, 18, e3000597.	5.6	60
71	A hybrid zone provides evidence for incipient ecological speciation in<i>Heliconius</i>butterflies. Molecular Ecology, 2008, 17, 4699-4712.	3.9	57
72	Insights into invasive species from wholeâ€genome resequencing. Molecular Ecology, 2021, 30, 6289-6308.	3.9	56

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73	Interplay between Developmental Flexibility and Determinism in the Evolution of Mimetic Heliconius Wing Patterns. Current Biology, 2019, 29, 3996-4009.e4.	3.9	55
74	Molecular systematics of the butterfly genus Ithomia (Lepidoptera: Ithomiinae): a composite phylogenetic hypothesis based on seven genes. Molecular Phylogenetics and Evolution, 2005, 34, 625-644.	2.7	54
75	Two sisters in the same dress: Heliconius cryptic species. BMC Evolutionary Biology, 2008, 8, 324.	3.2	54
76	Sex Chromosome Dosage Compensation in<i>Heliconius</i>Butterflies: Global yet Still Incomplete?. Genome Biology and Evolution, 2015, 7, 2545-2559.	2.5	54
77	Mutualistic Mimicry and Filtering by Altitude Shape the Structure of Andean Butterfly Communities. American Naturalist, 2014, 183, 26-39.	2.1	52
78	Into the Andes: multiple independent colonizations drive montane diversity in the Neotropical clearwing butterflies Godyridina. Molecular Ecology, 2016, 25, 5765-5784.	3.9	52
79	Highly conserved gene order and numerous novel repetitive elements in genomic regions linked to wing pattern variation in Heliconius butterflies. BMC Genomics, 2008, 9, 345.	2.8	51
80	North Andean origin and diversification of the largest ithomiine butterfly genus. Scientific Reports, 2017, 7, 45966.	3.3	48
81	Peace in Colombia is a critical moment for Neotropical connectivity and conservation: Save the northern Andesâ€"Amazon biodiversity bridge. Conservation Letters, 2019, 12, e12594.	5.7	46
82	Haplotype tagging reveals parallel formation of hybrid races in two butterfly species. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	7.1	46
83	Characterisation and expression of microRNAs in developing wings of the neotropical butterfly Heliconius melpomene. BMC Genomics, 2011, 12, 62.	2.8	44
84	Whole-chromosome hitchhiking driven by a male-killing endosymbiont. PLoS Biology, 2020, 18, e3000610.	5.6	44
85	Shared and divergent expression domains on mimetic<i>Heliconius</i>wings. Evolution & Development, 2009, 11, 498-512.	2.0	43
86	Microclimate buffering and thermal tolerance across elevations in a tropical butterfly. Journal of Experimental Biology, 2020, 223, .	1.7	41
87	Comparative genomics of the mimicry switch in<i>Papilio dardanus</i>. Proceedings of the Royal Society B: Biological Sciences, 2014, 281, 20140465.	2.6	40
88	Cortex cis-regulatory switches establish scale colour identity and pattern diversity in Heliconius. ELife, 2021, 10, .	6.0	40
89	The transcriptome response of <i>Heliconius melpomene</i> larvae to a novel host plant. Molecular Ecology, 2016, 25, 4850-4865.	3.9	39
90	Genome-wide analysis of ionotropic receptors provides insight into their evolution in Heliconius butterflies. BMC Genomics, 2016, 17, 254.	2.8	38

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91	Genomic tools and cDNA derived markers for butterflies. Molecular Ecology, 2005, 14, 2883-2897.	3.9	37
92	Evaluating female remating rates in light of spermatophore degradation in <i>Heliconius</i> butterflies: pupalâ€mating monandry versus adultâ€mating polyandry. Ecological Entomology, 2012, 37, 257-268.	2.2	37
93	Convergent, modular expression of ebony and tan in the mimetic wing patterns of Heliconius butterflies. Development Genes and Evolution, 2011, 221, 297-308.	0.9	36
94	Evolution: Mimicry meets the mitochondrion. Current Biology, 1996, 6, 937-940.	3.9	35
95	Colour pattern specification in the Mocker swallowtail Papilio dardanus : the transcription factor invected is a candidate for the mimicry locus H. Proceedings of the Royal Society B: Biological Sciences, 2008, 275, 1181-1188.	2.6	35
96	Male pheromone composition depends on larval but not adult diet in <i>Heliconius melpomene</i>. Ecological Entomology, 2019, 44, 397-405.	2.2	35
97	Rapidly Shifting Sex Ratio across a Species Range. Current Biology, 2009, 19, 1628-1631.	3.9	34
98	Avoidance of an aposematically coloured butterfly by wild birds in a tropical forest. Ecological Entomology, 2016, 41, 627-632.	2.2	34
99	Evolution of a mimicry supergene from a multilocus architecture. Proceedings of the Royal Society B: Biological Sciences, 2012, 279, 316-325.	2.6	33
100	The appearance of mimetic <i>Heliconius</i> butterflies to predators and conspecifics. Evolution; International Journal of Organic Evolution, 2018, 72, 2156-2166.	2.3	33
101	Conserved ancestral tropical niche but different continental histories explain the latitudinal diversity gradient in brush-footed butterflies. Nature Communications, 2021, 12, 5717.	12.8	33
102	ESTIMATING THE MATING BEHAVIOR OF A PAIR OF HYBRIDIZING <i>HELICONIUS</i> SPECIES IN THE WILD. Evolution; International Journal of Organic Evolution, 1998, 52, 503-510.	2.3	32
103	Sexually dimorphic gene expression and transcriptome evolution provide mixed evidence for a fastâ€Z effect in <i>Heliconius</i>. Journal of Evolutionary Biology, 2019, 32, 194-204.	1.7	31
104	Species specificity and intraspecific variation in the chemical profiles of <i>Heliconius</i> butterflies across a large geographic range. Ecology and Evolution, 2020, 10, 3895-3918.	1.9	31
105	Rampant Genome-Wide Admixture across the<i>Heliconius</i>Radiation. Genome Biology and Evolution, 2021, 13, .	2.5	31
106	Gene flow and the genealogical history of Heliconius heurippa. BMC Evolutionary Biology, 2008, 8, 132.	3.2	30
107	Butterfly Learning and the Diversification of Plant Leaf Shape. Frontiers in Ecology and Evolution, 2016, 4, .	2.2	29
108	A novel terpene synthase controls differences in anti-aphrodisiac pheromone production between closely related Heliconius butterflies. PLoS Biology, 2021, 19, e3001022.	5.6	29

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109	The Evolution of Sex Ratio Distorter Suppression Affects a 25 cM Genomic Region in the Butterfly Hypolimnas bolina. PLoS Genetics, 2014, 10, e1004822.	3.5	27
110	Altitude and lifeâ€history shape the evolution of<i>Heliconius</i>wings. Evolution; International Journal of Organic Evolution, 2019, 73, 2436-2450.	2.3	27
111	An introgressed wing pattern acts as a mating cue. Evolution; International Journal of Organic Evolution, 2015, 69, 1619-1629.	2.3	25
112	Divergence of chemosensing during the early stages of speciation. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 16438-16447.	7.1	25
113	Pollen feeding proteomics: Salivary proteins of the passion flower butterfly, Heliconius melpomene. Insect Biochemistry and Molecular Biology, 2015, 63, 7-13.	2.7	24
114	Visual mate preference evolution during butterfly speciation is linked to neural processing genes. Nature Communications, 2020, 11, 4763.	12.8	24
115	Deep mitochondrial divergence within a Heliconius butterfly species is not explained by cryptic speciation or endosymbiotic bacteria. BMC Evolutionary Biology, 2011, 11, 358.	3.2	23
116	Conservation and flexibility in the gene regulatory landscape of heliconiine butterfly wings. EvoDevo, 2019, 10, 15.	3.2	22
117	Estimating the Mating Behavior of a Pair of Hybridizing Heliconius Species in the Wild. Evolution; International Journal of Organic Evolution, 1998, 52, 503.	2.3	21
118	A major locus controls a biologically active pheromone component in <i>Heliconius melpomene</i>. Evolution; International Journal of Organic Evolution, 2020, 74, 349-364.	2.3	19
119	Genetic evidence for a sibling species of Heliconius charithonia (Lepidoptera; Nymphalidae). Biological Journal of the Linnean Society, 1998, 64, 57-67.	1.6	17
120	Behavioral and Physiological Differences between Two Parapatric Heliconius Species1. Biotropica, 1999, 31, 661-668.	1.6	17
121	THE PHYLOGENETIC PATTERN OF SPECIATION AND WING PATTERN CHANGE IN NEOTROPICAL ITHOMIA BUTTERFLIES (LEPIDOPTERA: NYMPHALIDAE). Evolution; International Journal of Organic Evolution, 2006, 60, 1454.	2.3	17
122	Ecologically relevant cryptic species in the highly polymorphic Amazonian butterfly Mechanitis mazaeusâ€ƒs.l. (Lepidoptera: Nymphalidae; Ithomiini). Biological Journal of the Linnean Society, 2012, 106, 540-560.	1.6	17
123	Adaptive dynamics: is speciation too easy?. Trends in Ecology and Evolution, 2000, 15, 225-226.	8.7	16
124	Assessing genotype-phenotype associations in three dorsal colour morphs in the meadow spittlebug Philaenus spumarius (L.) (Hemiptera: Aphrophoridae) using genomic and transcriptomic resources. BMC Genetics, 2016, 17, 144.	2.7	14
125	SEX-LINKED HYBRID STERILITY IN A BUTTERFLY. Evolution; International Journal of Organic Evolution, 2001, 55, 1631.	2.3	13
126	Deep Convergence, Shared Ancestry, and Evolutionary Novelty in the Genetic Architecture of <i>Heliconius</i> Mimicry. Genetics, 2020, 216, 765-780.	2.9	13

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127	Suppression of<i>Wolbachia</i>-mediated male-killing in the butterfly<i>Hypolimnas bolina</i>involves a single genomic region. PeerJ, 2019, 7, e7677.	2.0	13
128	The evolutionary genetics of highly divergent alleles of the mimicry locus in Papilio dardanus. BMC Evolutionary Biology, 2014, 14, 140.	3.2	12
129	Hybridization and transgressive exploration of colour pattern and wing morphology in <i>Heliconius</i> butterflies. Journal of Evolutionary Biology, 2020, 33, 942-956.	1.7	12
130	Phenotypic plasticity in chemical defence of butterflies allows usage of diverse host plants. Biology Letters, 2021, 17, 20200863.	2.3	12
131	MÃ¼llerian Mimicry: Sharing the Load Reduces the Legwork. Current Biology, 2009, 19, R687-R689.	3.9	11
132	A new subspecies in a Heliconius butterfly adaptive radiation (Lepidoptera: Nymphalidae). Zoological Journal of the Linnean Society, 2017, 180, 805-818.	2.3	11
133	Identification and Composition of Clasper Scent Gland Components of the Butterfly <i>Heliconius erato</i> and Its Relation to Mimicry. ChemBioChem, 2021, 22, 3300-3313.	2.6	10
134	Can genomics shed light on the origin of species?. PLoS Biology, 2019, 17, e3000394.	5.6	9
135	Population structure, adaptation and divergence of the meadow spittlebug, <i>Philaenus spumarius</i> (Hemiptera, Aphrophoridae), revealed by genomic and morphological data. PeerJ, 2021, 9, e11425.	2.0	9
136	Clustering of loci controlling species differences in male chemical bouquets of sympatric <i>Heliconius</i> butterflies. Ecology and Evolution, 2021, 11, 89-107.	1.9	9
137	Neighboring genes shaping a single adaptive mimetic trait. Evolution & Development, 2014, 16, 3-12.	2.0	8
138	Genomics of altitudeâ€associated wing shape in two tropical butterflies. Molecular Ecology, 2021, 30, 6387-6402.	3.9	8
139	Partial Complementarity of the Mimetic Yellow Bar Phenotype in Heliconius Butterflies. PLoS ONE, 2012, 7, e48627.	2.5	7
140	Evolutionary and ecological processes influencing chemical defense variation in an aposematic and mimetic <i>Heliconius</i> butterfly. PeerJ, 2021, 9, e11523.	2.0	7
141	A Narrow <i>Heliconius cydno</i> (Nymphalidae; Heliconiini) Hybrid Zone With Differences in Morph Sex Ratios¹. Biotropica, 2005, 37, 119-128.	1.6	6
142	A large deletion at the cortex locus eliminates butterfly wing patterning. G3: Genes, Genomes, Genetics, 2022, 12, .	1.8	6
143	Signatures of selection in loci governing major colour patterns in Heliconius butterflies and related species. BMC Evolutionary Biology, 2010, 10, 368.	3.2	5
144	A flamboyant behavioral polymorphism is controlled by a lethal supergene. Nature Genetics, 2016, 48, 7-8.	21.4	4

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145	Estimating the age of<i>Heliconius</i>butterflies from calibrated photographs. PeerJ, 2017, 5, e3821.	2.0	4
146	Reply from J.R. Bridle, C.D. Jiggins and T. Tregenza. Trends in Ecology and Evolution, 2000, 15, 420.	8.7	2
147	A Peppered Icon Enters the Genomic Era. BioScience, 2011, 61, 655-656.	4.9	2
148	What Can We Learn About Adaptation from the Wing Pattern Genetics of Heliconius Butterflies?. , 2017, , 173-188.		2
149	Plasticity in flower size as an adaptation to variation in pollinator specificity. Ecological Entomology, 2020, 45, 1367-1372.	2.2	2
150	Reply from C.D. Jiggins and J. Mallet. Trends in Ecology and Evolution, 2000, 15, 469.	8.7	1
151	Condition dependence in biosynthesized chemical defenses of an aposematic and mimetic <i>Heliconius</i> butterfly. Ecology and Evolution, 2022, 12, .	1.9	1
152	A gut feeling for isolation. Nature, 2013, 500, 412-413.	27.8	0
153	Functional genomics of supergene-controlled behavior in the white-throated sparrow. Faculty Reviews, 2021, 10, 75.	3.9	0