Philine Feulner

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6261213/publications.pdf

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38 papers 2,576 citations

304743

22

h-index

315739 38 g-index

42 all docs 42 docs citations

times ranked

42

4064 citing authors

#	Article	IF	CITATIONS
1	Strong selection and high mutation supply characterize experimental <i>Chlorovirus</i> evolution. Virus Evolution, 2022, 8, veac003.	4.9	5
2	Genomic variation from an extinct species is retained in the extant radiation following speciation reversal. Nature Ecology and Evolution, 2022, 6, 461-468.	7.8	12
3	Sequencing platform shifts provide opportunities but pose challenges for combining genomic data sets. Molecular Ecology Resources, 2021, 21, 653-660.	4.8	16
4	Examining adaptive evolution of immune activity: opportunities provided by gastropods in the age of â€~omics'. Philosophical Transactions of the Royal Society B: Biological Sciences, 2021, 376, 20200158.	4.0	6
5	A de novo chromosomeâ€level genome assembly of <i>Coregonus</i> sp. " <i>Balchen</i> â€l One representative of the Swiss Alpine whitefish radiation. Molecular Ecology Resources, 2020, 20, 1093-1109.	4.8	29
6	Genome-Wide Genotype-Expression Relationships Reveal Both Copy Number and Single Nucleotide Differentiation Contribute to Differential Gene Expression between Stickleback Ecotypes. Genome Biology and Evolution, 2019, 11, 2344-2359.	2.5	16
7	The feedback between selection and demography shapes genomic diversity during coevolution. Science Advances, 2019, 5, eaax0530.	10.3	20
8	Ecological and Evolutionary Processes Shaping Viral Genetic Diversity. Viruses, 2019, 11, 220.	3.3	21
9	Genomic insights into the vulnerability of sympatric whitefish species flocks. Molecular Ecology, 2019, 28, 615-629.	3.9	30
10	Evolution as an ecosystem process: insights from genomics. Genome, 2018, 61, 298-309.	2.0	11
11	Population size changes and selection drive patterns of parallel evolution in a host–virus system. Nature Communications, 2018, 9, 1706.	12.8	29
12	A European Whitefish Linkage Map and Its Implications for Understanding Genome-Wide Synteny Between Salmonids Following Whole Genome Duplication. G3: Genes, Genomes, Genetics, 2018, 8, 3745-3755.	1.8	16
13	A Dense Linkage Map of Lake Victoria Cichlids Improved the <i>Pundamilia</i> Genome Assembly and Revealed a Major QTL for Sex-Determination. G3: Genes, Genomes, Genetics, 2018, 8, 2411-2420.	1.8	28
14	Genome evolution, structural rearrangements and speciation. Journal of Evolutionary Biology, 2017, 30, 1488-1490.	1.7	40
15	Recombination: the good, the bad and the variable. Philosophical Transactions of the Royal Society B: Biological Sciences, 2017, 372, 20170279.	4.0	39
16	Variation in recombination frequency and distribution across eukaryotes: patterns and processes. Philosophical Transactions of the Royal Society B: Biological Sciences, 2017, 372, 20160455.	4.0	306
17	Transcriptome profiling of immune tissues reveals habitatâ€specific gene expression between lake and river sticklebacks. Molecular Ecology, 2016, 25, 943-958.	3.9	49
18	Species delimitation and phylogenetic relationships in a genus of African weakly-electric fishes (Osteoglossiformes, Mormyridae, Campylomormyrus). Molecular Phylogenetics and Evolution, 2016, 101, 8-18.	2.7	24

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19	Genomics of Divergence along a Continuum of Parapatric Population Differentiation. PLoS Genetics, 2015, 11, e1004966.	3.5	135
20	Comparative histology of the adult electric organ among four species of the genus Campylomormyrus (Teleostei: Mormyridae). Journal of Comparative Physiology A: Neuroethology, Sensory, Neural, and Behavioral Physiology, 2015, 201, 357-374.	1.6	19
21	Ecological and evolutionary implications of genomic structural variations. Frontiers in Genetics, 2014, 5, 326.	2.3	13
22	Extensive Copy-Number Variation of Young Genes across Stickleback Populations. PLoS Genetics, 2014, 10, e1004830.	3.5	70
23	Genomeâ€wide patterns of standing genetic variation in a marine population of threeâ€spined sticklebacks. Molecular Ecology, 2013, 22, 635-649.	3.9	78
24	Introgression and the fate of domesticated genes in a wild mammal population. Molecular Ecology, 2013, 22, 4210-4221.	3.9	53
25	Fewer invited talks by women in evolutionary biology symposia. Journal of Evolutionary Biology, 2013, 26, 2063-2069.	1.7	120
26	Evaluating Characteristics of De Novo Assembly Software on 454 Transcriptome Data: A Simulation Approach. PLoS ONE, 2012, 7, e31410.	2.5	72
27	Genome mapping in intensively studied wild vertebrate populations. Trends in Genetics, 2010, 26, 275-284.	6.7	85
28	Adaptation genomics: the next generation. Trends in Ecology and Evolution, 2010, 25, 705-712.	8.7	589
29	Electric Organ Discharge Divergence Promotes Ecological Speciation in Sympatrically Occurring African Weakly Electric Fish (Campylomormyrus). , 2010, , 307-321.		10
30	Magic trait Electric Organ Discharge (EOD). Communicative and Integrative Biology, 2009, 2, 329-331.	1.4	36
31	Phylogeography of red deer (<i>Cervus elaphus</i>) in Europe. Journal of Biogeography, 2009, 36, 66-77.	3.0	115
32	Electrifying love: electric fish use species-specific discharge for mate recognition. Biology Letters, 2009, 5, 225-228.	2.3	82
33	TOXIC HYDROGEN SULFIDE AND DARK CAVES: PHENOTYPIC AND GENETIC DIVERGENCE ACROSS TWO ABIOTIC ENVIRONMENTAL GRADIENTS IN <i>POECILIA MEXICANA</i> Organic Evolution, 2008, 62, 2643-2659.	2.3	122
34	Adaptive radiation in the Congo River: An ecological speciation scenario for African weakly electric fish (Teleostei; Mormyridae; Campylomormyrus). Journal of Physiology (Paris), 2008, 102, 340-346.	2.1	41
35	Adaptive radiation in African weakly electric fish (Teleostei: Mormyridae: Campylomormyrus): a combined molecular and morphological approach. Journal of Evolutionary Biology, 2007, 20, 403-414.	1.7	75
36	Electrophysiological and molecular genetic evidence for sympatrically occuring cryptic species in African weakly electric fishes (Teleostei: Mormyridae: Campylomormyrus). Molecular Phylogenetics and Evolution, 2006, 39, 198-208.	2.7	67

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37	Eighteen microsatellite loci for endemic African weakly electric fish (Campylomormyrus, Mormyridae) and their cross species applicability among related taxa. Molecular Ecology Notes, 2005, 5, 446-448.	1.7	12
38	Mitochondrial DNA and microsatellite analyses of the genetic status of the presumed subspecies Cervus elaphus montanus (Carpathian red deer). Heredity, 2004, 93, 299-306.	2.6	66