Dominik R Laetsch

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/1382946/publications.pdf

Version: 2024-02-01

25 papers 1,870 citations

567281 15 h-index 677142 22 g-index

40 all docs 40 docs citations

times ranked

40

3283 citing authors

#	Article	IF	CITATIONS
1	BlobTools: Interrogation of genome assemblies. F1000Research, 0, 6, 1287.	1.6	600
2	No evidence for extensive horizontal gene transfer in the genome of the tardigrade <i>Hypsibius dujardini</i> . Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 5053-5058.	7.1	214
3	Comparative genomics of the tardigrades Hypsibius dujardini and Ramazzottius varieornatus. PLoS Biology, 2017, 15, e2002266.	5.6	170
4	The genome of the yellow potato cyst nematode, Globodera rostochiensis, reveals insights into the basis of parasitism and virulence. Genome Biology, 2016, 17, 124.	8.8	156
5	Comparative Genomics of Apomictic Root-Knot Nematodes: Hybridization, Ploidy, and Dynamic Genome Change. Genome Biology and Evolution, 2017, 9, 2844-2861.	2.5	98
6	The determinants of genetic diversity in butterflies. Nature Communications, 2019, 10, 3466.	12.8	80
7	Secretion of an Argonaute protein by a parasitic nematode and the evolution of its siRNA guides. Nucleic Acids Research, 2019, 47, 3594-3606.	14.5	75
8	KinFin: Software for Taxon-Aware Analysis of Clustered Protein Sequences. G3: Genes, Genomes, Genetics, 2017, 7, 3349-3357.	1.8	72
9	The phylogenetics of Anguillicolidae (Nematoda: Anguillicoloidea), swimbladder parasites of eels. BMC Evolutionary Biology, 2012, 12, 60.	3.2	56
10	The submarine volcano eruption at the island of El Hierro: physical-chemical perturbation and biological response. Scientific Reports, 2012, 2, 486.	3.3	52
11	Characterisation of the British honey bee metagenome. Nature Communications, 2018, 9, 4995.	12.8	51
12	Signatures of balancing selection in toll-like receptor (TLRs) genes – novel insights from a free-living rodent. Scientific Reports, 2018, 8, 8361.	3. 3	38
13	The Pleistocene species pump past its prime: Evidence from European butterfly sister species. Molecular Ecology, 2021, 30, 3575-3589.	3.9	35
14	Sexâ€specific expression and DNA methylation in a species with extreme sexual dimorphism and paternal genome elimination. Molecular Ecology, 2021, 30, 5687-5703.	3.9	29
15	Males That Silence Their Father's Genes: Genomic Imprinting of a Complete Haploid Genome. Molecular Biology and Evolution, 2021, 38, 2566-2581.	8.9	23
16	Massive encapsulation of larval Anguillicoloides crassus in the intestinal wall of Japanese eels. Parasites and Vectors, 2009, 2, 48.	2.5	19
17	Improving the annotation of the Heterorhabditis bacteriophora genome. GigaScience, 2018, 7, .	6.4	18
18	MolluscDB: a genome and transcriptome database for molluscs. Philosophical Transactions of the Royal Society B: Biological Sciences, 2021, 376, 20200157.	4.0	17

#	Article	IF	CITATIONS
19	The Evolution of Interdependence in a Four-Way Mealybug Symbiosis. Genome Biology and Evolution, 2021, 13, .	2.5	9
20	The genome sequence of the lesser marbled fritillary, <i>Brenthis ino</i> , and evidence for a segregating neo-Z chromosome. G3: Genes, Genomes, Genetics, 2022, 12, .	1.8	8
21	The genome sequence of the large tortoiseshell, Nymphalis polychloros (Linnaeus, 1758). Wellcome Open Research, 2021, 6, 238.	1.8	3
22	The genome sequence of the small copper, Lycaena phlaeas (Linnaeus, 1760). Wellcome Open Research, 0, 6, 294.	1.8	1
23	The genome sequence of the heath fritillary, Melitaea athalia (Rottemburg, 1775). Wellcome Open Research, 2021, 6, 304.	1.8	1
24	The genome sequence of the marbled white butterfly, Melanargia galathea (Linnaeus, 1758). Wellcome Open Research, 0, 7, 123.	1.8	1
25	The genome sequence of the grizzled skipper, Pyrgus malvae (Linnaeus, 1758). Wellcome Open Research, 0, 7, 114.	1.8	0