

Guangyi Fan

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

23
papers

492
citations

10
h-index

22
g-index

29
ext. papers

741
ext. citations

6.5
avg, IF

2.82
L-index

#	Paper	IF	Citations
23	SLR-superscaffolder: a de novo scaffolding tool for synthetic long reads using a top-to-bottom scheme. <i>BMC Bioinformatics</i> , 2021 , 22, 158	3.6	1
22	Reconstruction of the Origin of a Neo-Y Sex Chromosome and Its Evolution in the Spotted Knifejaw, <i>Oplegnathus punctatus</i> . <i>Molecular Biology and Evolution</i> , 2021 , 38, 2615-2626	8.3	7
21	Toward the massive genome of <i>Proteus anguinus</i> -illuminating longevity, regeneration, convergent evolution, and metabolic disorders. <i>Annals of the New York Academy of Sciences</i> , 2021 ,	6.5	5
20	Complete Chloroplast Genomes of 14 Mangroves: Phylogenetic and Comparative Genomic Analyses. <i>BioMed Research International</i> , 2020 , 2020, 8731857	3	7
19	Dynamics of bacteriophages in gut of giant pandas reveal a potential regulation of dietary intake on bacteriophage composition. <i>Science of the Total Environment</i> , 2020 , 734, 139424	10.2	4
18	The Chromosome Level Genome and Genome-wide Association Study for the Agronomic Traits of. <i>IScience</i> , 2020 , 23, 101538	6.1	12
17	Beaver and Naked Mole Rat Genomes Reveal Common Paths to Longevity. <i>Cell Reports</i> , 2020 , 32, 107949	10.6	10
16	African Arowana Genome Provides Insights on Ancient Teleost Evolution. <i>IScience</i> , 2020 , 23, 101662	6.1	1
15	TGS-GapCloser: A fast and accurate gap closer for large genomes with low coverage of error-prone long reads. <i>GigaScience</i> , 2020 , 9,	7.6	30
14	Initial data release and announcement of the 10,000 Fish Genomes Project (Fish10K). <i>GigaScience</i> , 2020 , 9,	7.6	12
13	Genome sequencing of deep-sea hydrothermal vent snails reveals adaptations to extreme environments. <i>GigaScience</i> , 2020 , 9,	7.6	2
12	A chromosome-level genome of black rockfish, <i>Sebastes schlegelii</i> , provides insights into the evolution of live birth. <i>Molecular Ecology Resources</i> , 2019 , 19, 1309-1321	8.4	25
11	Chromosome-level reference genome of the Siamese fighting fish <i>Betta splendens</i> , a model species for the study of aggression. <i>GigaScience</i> , 2018 , 7,	7.6	14
10	The chromosome-level genome assemblies of two rattans (<i>Calamus simplicifolius</i> and <i>Daemonorops jenkinsiana</i>). <i>GigaScience</i> , 2018 , 7,	7.6	11
9	Dynamics of Gut Microbiome in Giant Panda Cubs Reveal Transitional Microbes and Pathways in Early Life. <i>Frontiers in Microbiology</i> , 2018 , 9, 3138	5.7	15
8	Evolutionary gradient of predicted nuclear localization signals (NLS)-bearing proteins in genomes of family Planctomycetaceae. <i>BMC Microbiology</i> , 2017 , 17, 86	4.5	2
7	Draft genome of the living fossil <i>Ginkgo biloba</i> . <i>GigaScience</i> , 2016 , 5, 49	7.6	161

6	The genetic basis for ecological adaptation of the Atlantic herring revealed by genome sequencing. <i>ELife</i> , 2016 , 5,	8.9	103
5	The Asian arowana (<i>Scleropages formosus</i>) genome provides new insights into the evolution of an early lineage of teleosts. <i>Scientific Reports</i> , 2016 , 6, 24501	4.9	66
4	Bicolor angelfish (<i>Centropyge bicolor</i>) provides the first chromosome-level genome of the Pomacanthidae family. <i>GigaByte</i> , 2021, 1-13		
3	Complete chloroplast genomes of 14 mangroves: phylogenetic and genomic comparative analyses		3
2	Chromosome-level genome assembly of the humpback puffer, <i>Tetraodon palembangensis</i> . <i>GigaByte</i> , 2021, 1-12		
1	Microbiota-muscle/immune interactions in rhesus macaque under simulated microgravity revealed by integrated multi-omics analysis. <i>JCSM Rapid Communications</i> ,	2.6	1