He Zhang

List of Publications by Year in descending order

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Version: 2024-02-01

430874 677142 3,488 23 18 22 citations h-index g-index papers 27 27 27 5143 citing authors all docs docs citations times ranked

#	Article	IF	CITATIONS
1	Genome sequence of cultivated Upland cotton (GossypiumÂhirsutum TM-1) provides insights into genome evolution. Nature Biotechnology, 2015, 33, 524-530.	17.5	1,064
2	Pearl millet genome sequence provides a resource to improve agronomic traits in arid environments. Nature Biotechnology, 2017, 35, 969-976.	17.5	356
3	Structural genomic changes underlie alternative reproductive strategies in the ruff (Philomachus) Tj ETQq $1\ 1\ 0.78$	84314 rgB ⁷ 21.4	T /Overlock
4	The Sinocyclocheilus cavefish genome provides insights into cave adaptation. BMC Biology, 2016, 14, 1.	3.8	292
5	The highâ€quality genome of <i>Brassica napus</i> cultivar â€~ <scp>ZS</scp> 11' reveals the introgression history in semiâ€winter morphotype. Plant Journal, 2017, 92, 452-468.	5.7	233
6	Draft genome of the living fossil Ginkgo biloba. GigaScience, 2016, 5, 49.	6.4	232
7	Allele-aware chromosome-level genome assembly and efficient transgene-free genome editing for the autotetraploid cultivated alfalfa. Nature Communications, 2020, 11 , 2494.	12.8	224
8	The genetic basis for ecological adaptation of the Atlantic herring revealed by genome sequencing. ELife, 2016, 5, .	6.0	143
9	Resequencing 545 ginkgo genomes across the world reveals the evolutionary history of the living fossil. Nature Communications, 2019, 10, 4201.	12.8	99
10	The Asian arowana (Scleropages formosus) genome provides new insights into the evolution of an early lineage of teleosts. Scientific Reports, 2016, 6, 24501.	3.3	89
11	A chromosome-level assembly of the Atlantic herring genomeâ€"detection of a supergene and other signals of selection. Genome Research, 2019, 29, 1919-1928.	5.5	84
12	The genetic architecture of floral traits in the woody plant Prunus mume. Nature Communications, 2018, 9, 1702.	12.8	73
13	Initial data release and announcement of the 10,000 Fish Genomes Project (Fish10K). GigaScience, 2020, 9, .	6.4	47
14	Deletion and tandem duplications of biosynthetic genes drive the diversity of triterpenoids in Aralia elata. Nature Communications, 2022, 13, 2224.	12.8	34
15	Draft genome sequence of the Tibetan medicinal herb Rhodiola crenulata. GigaScience, 2017, 6, 1-5.	6.4	33
16	<i>Arabidopsis</i> DXO1 possesses deNADding and exonuclease activities and its mutation affects defenseâ€related and photosynthetic gene expression. Journal of Integrative Plant Biology, 2020, 62, 967-983.	8.5	29
17	Genome sequences reveal global dispersal routes and suggest convergent genetic adaptations in seahorse evolution. Nature Communications, 2021, 12, 1094.	12.8	29
18	The first chromosomeâ€level genome for a marine mammal as a resource to study ecology and evolution. Molecular Ecology Resources, 2019, 19, 944-956.	4.8	27

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19	Chromosome-level reference genome of the Siamese fighting fish Betta splendens, a model species for the study of aggression. GigaScience, 2018, 7, .	6.4	25
20	A survey of the sperm whale (<i>Physeter catodon</i>) commensal microbiome. PeerJ, 2019, 7, e7257.	2.0	15
21	<i>De Novo</i> Biosynthesis of Oleanane-Type Ginsenosides in <i>Saccharomyces cerevisiae</i> Using Two Types of Glycosyltransferases from <i>Panax ginseng</i> Journal of Agricultural and Food Chemistry, 2022, 70, 2231-2240.	5.2	14
22	Chromosome-level genome assembly of the humpback puffer, Tetraodon palembangensis. GigaByte, 0, 2021, 1-12.	0.0	0
23	Bicolor angelfish (Centropyge bicolor) provides the first chromosome-level genome of the Pomacanthidae family. GigaByte, 0, 2021, 1-13.	0.0	0