

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3590928/publications.pdf Version: 2024-02-01



LINC CAL

#	Article	IF	CITATIONS
1	The sequence and de novo assembly of the giant panda genome. Nature, 2010, 463, 311-317.	13.7	1,058
2	The genome sequence of the orchid Phalaenopsis equestris. Nature Genetics, 2015, 47, 65-72.	9.4	413
3	<i>De Novo</i> Origination of a New Protein-Coding Gene in <i>Saccharomyces cerevisiae</i> . Genetics, 2008, 179, 487-496.	1.2	209
4	Engineering yeast for the production of breviscapine by genomic analysis and synthetic biology approaches. Nature Communications, 2018, 9, 448.	5.8	146
5	The Earth BioGenome Project 2020: Starting the clock. Proceedings of the National Academy of Sciences of the United States of America, 2022, 119, .	3.3	124
6	A de novo originated gene depresses budding yeast mating pathway and is repressed by the protein encoded by its antisense strand. Cell Research, 2010, 20, 408-420.	5.7	110
7	Apoptosis Induced by Dioscin in Hela Cells Biological and Pharmaceutical Bulletin, 2002, 25, 193-196.	0.6	102
8	Building a Genetic Manipulation Tool Box for Orchid Biology: Identification of Constitutive Promoters and Application of CRISPR/Cas9 in the Orchid, Dendrobium officinale. Frontiers in Plant Science, 2016, 7, 2036.	1.7	102
9	Biosynthesis and engineering of kaempferol in Saccharomyces cerevisiae. Microbial Cell Factories, 2017, 16, 165.	1.9	68
10	Factors affecting crop damage by wild boar and methods of mitigation in a giant panda reserve. European Journal of Wildlife Research, 2008, 54, 723-728.	0.7	58
11	Genome and Comparative Transcriptomics of African Wild Rice Oryza longistaminata Provide Insights into Molecular Mechanism of Rhizomatousness and Self-Incompatibility. Molecular Plant, 2015, 8, 1683-1686.	3.9	49
12	The origin and evolution of the diosgenin biosynthetic pathway in yam. Plant Communications, 2021, 2, 100079.	3.6	44
13	Phytochemicals, pharmacology, clinical application, patents, and products of Amomi fructus. Food and Chemical Toxicology, 2018, 119, 31-36.	1.8	42
14	The Transcriptome of the Zoanthid <i>Protopalythoa variabilis</i> (Cnidaria, Anthozoa) Predicts a Basal Repertoire of Toxin-like and Venom-Auxiliary Polypeptides. Genome Biology and Evolution, 2016, 8, 3045-3064.	1.1	37
15	High-fat diet feeding and palmitic acid increase CRC growth in β2AR-dependent manner. Cell Death and Disease, 2019, 10, 711.	2.7	33
16	Identification of candidate genes involved in isoquinoline alkaloids biosynthesis in Dactylicapnos scandens by transcriptome analysis. Scientific Reports, 2017, 7, 9119.	1.6	26
17	A Phylogenomic Analysis of the Floral Transcriptomes of Sexually Deceptive and Rewarding European Orchids, Ophrys and Gymnadenia. Frontiers in Plant Science, 2019, 10, 1553.	1.7	26
18	Chromosome-level reference genome of the Siamese fighting fish Betta splendens, a model species for the study of aggression. GigaScience, 2018, 7, .	3.3	25

Jing Cai

#	Article	IF	CITATIONS
19	Current status and conservation of the Endangered Przewalski's gazelle <i>Procapra przewalskii</i> , endemic to the Qinghai–Tibetan Plateau, China. Oryx, 2012, 46, 145-153.	0.5	23
20	Identification of long non-coding RNAs in two anthozoan species and their possible implications for coral bleaching. Scientific Reports, 2017, 7, 5333.	1.6	22
21	Paraholcoglossum and Tsiorchis, Two New Orchid Genera Established by Molecular and Morphological Analyses of the Holcoglossum Alliance. PLoS ONE, 2011, 6, e24864.	1.1	21
22	Emerging trends and new developments in monoclonal antibodies: A scientometric analysis (1980–2016). Human Vaccines and Immunotherapeutics, 2017, 13, 1388-1397.	1.4	21
23	Do local communities support the conservation of endangered Przewalski's gazelle?. European Journal of Wildlife Research, 2010, 56, 551-560.	0.7	19
24	A New Molecular Phylogeny and a New Genus, Pendulorchis, of the Aerides–Vanda Alliance (Orchidaceae: Epidendroideae). PLoS ONE, 2013, 8, e60097.	1.1	17
25	The molecular bases of floral scent evolution under artificial selection: insights from a transcriptome analysis in Brassica rapa. Scientific Reports, 2016, 6, 36966.	1.6	17
26	Comparison of volatile compounds in different parts of fresh Amomum villosum Lour. from different geographical areas using cryogenic grinding combined HS–SPME–GC–MS. Chinese Medicine, 2020, 15, 97.	1.6	16
27	Genome sequence of Apostasia ramifera provides insights into the adaptive evolution in orchids. BMC Genomics, 2021, 22, 536.	1.2	9
28	Multiple Inter-Kingdom Horizontal Gene Transfers in the Evolution of the Phosphoenolpyruvate Carboxylase Gene Family. PLoS ONE, 2012, 7, e51159.	1.1	7
29	Sequencing, annotation and comparative analysis of nine BACs of giant panda (Ailuropoda) Tj ETQq1 1 0.784314	4 rgBT /Ov	erlock 10 Tf 5
30	Sinocurculigo, a New Genus of Hypoxidaceae from China Based on Molecular and Morphological Evidence. PLoS ONE, 2012, 7, e38880.	1.1	6
31	Comparative and phylogenetic analyses of eleven complete chloroplast genomes of Dipterocarpoideae. Chinese Medicine, 2021, 16, 125.	1.6	5
32	H2S regulates low oxygen signaling via integration with the unfolded protein response in Arabidopsis thaliana. Plant and Soil, 2021, 467, 531-547.	1.8	4
33	The complete chloroplast genome of the essential medicinal herb, <i>Amomum Villosum</i> (Zingiberaceae). Mitochondrial DNA Part B: Resources, 2019, 4, 1798-1799.	0.2	2
34	Chromosomeâ€level genome assembly of <i>Welwitschia mirabilis</i> , a unique Namib Desert species. Molecular Ecology Resources, 2022, 22, 391-403.	2.2	1