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List of Publications by Year in descending order

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687363 552781 27 726 13 26 citations h-index g-index papers 32 32 32 653 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Different genome-wide transcriptome responses of Nocardioides simplex VKM Ac-2033D to phytosterol and cortisone 21-acetate. BMC Biotechnology, 2021, 21, 7.	3.3	12
2	Comparative Analysis of Plastid Genomes in the Non-photosynthetic Genus Thismia Reveals Ongoing Gene Set Reduction. Frontiers in Plant Science, 2021, 12, 602598.	3.6	13
3	Palynological study of Asian Thismia (Thismiaceae: Dioscoreales) reveals an unusual pollen type. Plant Systematics and Evolution, 2021, 307, 1.	0.9	3
4	Genomic comparison of non-photosynthetic plants from the family Balanophoraceae with their photosynthetic relatives. PeerJ, 2021, 9, e12106.	2.0	7
5	Excessive Parallelism in Protein Evolution of Lake Baikal Amphipod Species Flock. Genome Biology and Evolution, 2020, 12, 1493-1503.	2.5	1
6	Complete genome assembly data of paenibacillus sp. RUD330, a hypothetical symbiont of euglena gracilis. Data in Brief, 2020, 32, 106070.	1.0	0
7	Genome-Wide Transcriptome Profiling Provides Insight on Cholesterol and Lithocholate Degradation Mechanisms in Nocardioides simplex VKM Ac-2033D. Genes, 2020, 11, 1229.	2.4	8
8	Mitochondrial Genome of Fagopyrum esculentum and the Genetic Diversity of Extranuclear Genomes in Buckwheat. Plants, 2020, 9, 618.	3 . 5	16
9	Assembly and Analysis of the Complete Mitochondrial Genome of Capsella bursa-pastoris. Plants, 2020, 9, 469.	3 . 5	14
10	Phylogenetics of the mycoheterotrophic genus <i>Thismia</i> (Thismiaceae: Dioscoreales) with a focus on the Old World taxa: delineation of novel natural groups and insights into the evolution of morphological traits. Botanical Journal of the Linnean Society, 2020, 193, 287-315.	1.6	24
11	Mitochondrial genome of the nonphotosynthetic mycoheterotrophic plant <i>Hypopitys monotropa</i> , its structure, gene expression and RNA editing. PeerJ, 2020, 8, e9309.	2.0	16
12	Genome-wide response on phytosterol in 9-hydroxyandrostenedione-producing strain of Mycobacterium sp. VKM Ac-1817D. BMC Biotechnology, 2019, 19, 39.	3.3	20
13	Knockdown of the neuronal gene Lim3 at the early stages of development affects mitochondrial function and lifespan in Drosophila. Mechanisms of Ageing and Development, 2019, 181, 29-41.	4.6	12
14	<i>Rhopalocnemis phalloides</i> has one of the most reduced and mutated plastid genomes known. PeerJ, 2019, 7, e7500.	2.0	25
15	The complete genome of the oil emulsifying strain Thalassolituus oleivorans K-188 from the Barents Sea. Marine Genomics, 2018, 37, 18-20.	1.1	4
16	RNA-seq highlights parallel and contrasting patterns in the evolution of the nuclear genome of fully mycoheterotrophic plants. BMC Genomics, 2018, 19, 602.	2.8	16
17	Effect of methyl-Î ² -cyclodextrin on gene expression in microbial conversion of phytosterol. Applied Microbiology and Biotechnology, 2017, 101, 4659-4667.	3 . 6	23
18	Genome Sequencing of Steroid-Producing Bacteria with Illumina Technology. Methods in Molecular Biology, 2017, 1645, 29-44.	0.9	1

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19	Comparative analysis of plastid genomes of non-photosynthetic Ericaceae and their photosynthetic relatives. Scientific Reports, 2016, 6, 30042.	3.3	47
20	Genome-wide bioinformatics analysis of steroid metabolism-associated genes in Nocardioides simplex VKM Ac-2033D. Current Genetics, 2016, 62, 643-656.	1.7	40
21	Complete Genome Sequence of Mycobacterium sp. Strain VKM Ac-1817D, Capable of Producing 9α-Hydroxy-androst-4-ene-3,17-dione from Phytosterol. Genome Announcements, 2015, 3, .	0.8	15
22	Complete Genome Sequence of Steroid-Transforming Nocardioides simplex VKM Ac-2033D. Genome Announcements, 2015, 3, .	0.8	13
23	Exploring the Limits for Reduction of Plastid Genomes: A Case Study of the Mycoheterotrophic Orchids Epipogium aphyllum and Epipogium roseum. Genome Biology and Evolution, 2015, 7, 1179-1191.	2.5	116
24	The Plastid Genome of Mycoheterotrophic Monocot Petrosavia stellaris Exhibits Both Gene Losses and Multiple Rearrangements. Genome Biology and Evolution, 2014, 6, 238-246.	2.5	75
25	Complete Genome Sequence of Sterol-Transforming Mycobacterium neoaurum Strain VKM Ac-1815D. Genome Announcements, 2014, 2, .	0.8	23
26	Comparative analysis of genes encoding key steroid core oxidation enzymes in fast-growing Mycobacterium spp. strains. Journal of Steroid Biochemistry and Molecular Biology, 2013, 138, 41-53.	2.5	66
27	Sequencing and Analysis of Plastid Genome in Mycoheterotrophic Orchid Neottia nidus-avis. Genome Biology and Evolution, 2011, 3, 1296-1303.	2.5	111