Xiaomin Yu

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6535218/publications.pdf

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23	985	16	25
papers	citations	h-index	g-index
26	26	26	1253 citing authors
all docs	docs citations	times ranked	

#	Article	IF	CITATIONS
1	Diversity and abundance of phosphonate biosynthetic genes in nature. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 20759-20764.	7.1	148
2	Interlinked regulatory loops of ABA catabolism and biosynthesis coordinate fruit growth and ripening in woodland strawberry. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, E11542-E11550.	7.1	142
3	Metabolite Profiling of 14 Wuyi Rock Tea Cultivars Using UPLC-QTOF MS and UPLC-QqQ MS Combined with Chemometrics. Molecules, 2018, 23, 104.	3.8	90
4	Metabolite signatures of diverse Camellia sinensis tea populations. Nature Communications, 2020, 11, 5586.	12.8	78
5	Transcriptome dynamics of Camellia sinensis in response to continuous salinity and drought stress. Tree Genetics and Genomes, 2017, 13, 1.	1.6	67
6	Defensive Responses of Tea Plants (Camellia sinensis) Against Tea Green Leafhopper Attack: A Multi-Omics Study. Frontiers in Plant Science, 2019, 10, 1705.	3.6	63
7	Non-targeted metabolomics analysis reveals dynamic changes of volatile and non-volatile metabolites during oolong tea manufacture. Food Research International, 2020, 128, 108778.	6.2	62
8	Endophytic Actinomycetes from Tea Plants (<i>Camellia sinensis</i>): Isolation, Abundance, Antimicrobial, and Plant-Growth-Promoting Activities. BioMed Research International, 2018, 2018, 1-12.	1.9	53
9	Metabolomics Reveals Distinct Carbon and Nitrogen Metabolic Responses to Magnesium Deficiency in Leaves and Roots of Soybean [Glycine max (Linn.) Merr.]. Frontiers in Plant Science, 2017, 8, 2091.	3.6	46
10	Use of a Phosphonate Methyltransferase in the Identification of the Fosfazinomycin Biosynthetic Gene Cluster. Angewandte Chemie - International Edition, 2014, 53, 1334-1337.	13.8	40
11	Purification and Characterization of Phosphonoglycans from Glycomyces sp. Strain NRRL B-16210 and Stackebrandtia nassauensis NRRL B-16338. Journal of Bacteriology, 2014, 196, 1768-1779.	2.2	27
12	Cyanohydrin Phosphonate Natural Product from <i>Streptomyces regensis</i> . Journal of Natural Products, 2014, 77, 243-249.	3.0	24
13	Insights into Tissue-specific Specialized Metabolism in Tieguanyin Tea Cultivar by Untargeted Metabolomics. Molecules, 2018, 23, 1817.	3 . 8	24
14	New Insights into Stress-Induced \hat{l}^2 -Ocimene Biosynthesis in Tea (<i>Camellia sinensis</i>) Leaves during Oolong Tea Processing. Journal of Agricultural and Food Chemistry, 2021, 69, 11656-11664.	5.2	21
15	Conserved biosynthetic pathways for phosalacine, bialaphos and newly discovered phosphonic acid natural products. Journal of Antibiotics, 2016, 69, 15-25.	2.0	20
16	Neuroprotective and Anti-Amyloid \hat{l}^2 Effect and Main Chemical Profiles of White Tea: Comparison Against Green, Oolong and Black Tea. Molecules, 2019, 24, 1926.	3.8	19
17	Production of purple Ma bamboo (Dendrocalamus latiflorus Munro) with enhanced drought and cold stress tolerance by engineering anthocyanin biosynthesis. Planta, 2021, 254, 50.	3. 2	15
18	H and HL synergistically regulate jasmonate-triggered trichome formation in tomato. Horticulture Research, 2022, 9, .	6.3	14

#	Article	IF	CITATIONS
19	Access to chiral \hat{l} ±-substituted- \hat{l}^2 -hydroxy arylphosphonates enabled by biocatalytic dynamic reductive kinetic resolution. Organic and Biomolecular Chemistry, 2020, 18, 2672-2677.	2.8	9
20	Draft Genome Sequence of <i>Streptomyces</i> sp. XY006, an Endophyte Isolated from Tea (<i>Camellia) Tj ET</i>	⁻ Qq0,00 r	gBT _t /Overlock
21	Expression and characterization of inhA gene from Bacillus thuringiensis 8010. World Journal of Microbiology and Biotechnology, 2007, 23, 1621-1625.	3.6	4
22	RAP-PCR fingerprinting reveals time-dependent expression of development-related genes following differentiation process of Bacillus thuringiensis. Canadian Journal of Microbiology, 2015, 61, 683-690.	1.7	2
23	Draft Genome Sequence of Paenibacillus sp. XY044, a Potential Plant Growth Promoter Isolated from a Tea Plant. Genome Announcements, 2017, 5, .	0.8	1