## Qiang Wang

List of Publications by Year in descending order

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OLANG WANG

#	Article	IF	CITATIONS
1	Deceptive Complexity in Formation of Cleistantha-8,12-diene. Organic Letters, 2022, 24, 2646-2649.	4.6	2
2	Direct formation of the sesquiterpeonid ether liguloxide by a terpene synthase in Senecio scandens. Plant Molecular Biology, 2021, 105, 55-64.	3.9	2
3	Maize transcription factor ZmEREB20 enhanced salt tolerance in transgenic Arabidopsis. Plant Physiology and Biochemistry, 2021, 159, 257-267.	5.8	28
4	Rice contains a biosynthetic gene cluster associated with production of the casbaneâ€type diterpenoid phytoalexin <i>ent</i> â€10â€oxodepressin. New Phytologist, 2021, 231, 85-93.	7.3	21
5	Maize WRKY Transcription Factor ZmWRKY79 Positively Regulates Drought Tolerance through Elevating ABA Biosynthesis. International Journal of Molecular Sciences, 2021, 22, 10080.	4.1	38
6	ZmMYC2 exhibits diverse functions and enhances JA signaling in transgenic Arabidopsis. Plant Cell Reports, 2020, 39, 273-288.	5.6	23
7	Promoter Variation Results in Differential Phytoalexin Accumulation in Two Maize Inbred Lines. Plant Molecular Biology Reporter, 2020, 38, 165-174.	1.8	3
8	Probing Enzymatic Structure and Function in the Dihydroxylating Sesquiterpene Synthase ZmEDS. Biochemistry, 2020, 59, 2660-2666.	2.5	5
9	Transcriptional Factors Regulate Plant Stress Responses Through Mediating Secondary Metabolism. Genes, 2020, 11, 346.	2.4	138
10	A Wheat β-Patchoulene Synthase Confers Resistance against Herbivory in Transgenic Arabidopsis. Genes, 2019, 10, 441.	2.4	5
11	CYP71Z18 overexpression confers elevated blast resistance in transgenic rice. Plant Molecular Biology, 2019, 100, 579-589.	3.9	16
12	Multiple genes recruited from hormone pathways partition maize diterpenoid defences. Nature Plants, 2019, 5, 1043-1056.	9.3	60
13	Discovery, Biosynthesis and Stress-Related Accumulation of Dolabradiene-Derived Defenses in Maize. Plant Physiology, 2018, 176, 2677-2690.	4.8	94
14	Direct production of dihydroxylated sesquiterpenoids by a maize terpene synthase. Plant Journal, 2018, 94, 847-856.	5.7	27
15	ZmWRKY79 positively regulates maize phytoalexin biosynthetic gene expression and is involved in stress response. Journal of Experimental Botany, 2018, 69, 497-510.	4.8	51
16	Indole primes plant defense against necrotrophic fungal pathogen infection. PLoS ONE, 2018, 13, e0207607.	2.5	30
17	CYP701A26 is characterized as an ent-kaurene oxidase with putative involvement in maize gibberellin biosynthesis. Biotechnology Letters, 2017, 39, 1709-1716.	2.2	10
18	Functional characterization of ZmTPS7 reveals a maize Ï"-cadinol synthase involved in stress response. Planta, 2016, 244, 1065-1074.	3.2	17

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19	A Tandem Array of <i>ent</i> -Kaurene Synthases in Maize with Roles in Gibberellin and More Specialized Metabolism. Plant Physiology, 2016, 170, 742-751.	4.8	81
20	Characterization of CYP71Z18 indicates a role in maize zealexin biosynthesis. Phytochemistry, 2016, 121, 4-10.	2.9	43
21	Formation of the Unusual Semivolatile Diterpene Rhizathalene by the <i>Arabidopsis</i> Class I Terpene Synthase TPS08 in the Root Stele Is Involved in Defense against Belowground Herbivory. Plant Cell, 2013, 25, 1108-1125.	6.6	123
22	Picking sides: distinct roles for CYP76M6 and CYP76M8Âin rice oryzalexin biosynthesis. Biochemical Journal, 2013, 454, 209-216.	3.7	48
23	CYP701A8: A Rice <i>ent</i> -Kaurene Oxidase Paralog Diverted to More Specialized Diterpenoid Metabolism  Â. Plant Physiology, 2012, 158, 1418-1425.	4.8	109
24	Characterization of CYP76M5–8 Indicates Metabolic Plasticity within a Plant Biosynthetic Gene Cluster. Journal of Biological Chemistry, 2012, 287, 6159-6168.	3.4	116
25	CYP99A3: functional identification of a diterpene oxidase from the momilactone biosynthetic gene cluster in rice. Plant Journal, 2011, 65, 87-95.	5.7	102
26	Parsing a multifunctional biosynthetic gene cluster from rice: Biochemical characterization of CYP71Z6 & 7. FEBS Letters, 2011, 585, 3446-3451.	2.8	70
27	CYP76M7 Is an <i>ent</i> -Cassadiene C11α-Hydroxylase Defining a Second Multifunctional Diterpenoid Biosynthetic Gene Cluster in Rice Â. Plant Cell, 2009, 21, 3315-3325.	6.6	199