## Lin Fang

## List of Publications by Year in descending order

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840776 642732 24 610 11 23 citations h-index g-index papers 28 28 28 706 all docs docs citations times ranked citing authors

#	Article	IF	Citations
1	Eudicot plant-specific sphingolipids determine host selectivity of microbial NLP cytolysins. Science, 2017, 358, 1431-1434.	12.6	167
2	Loss of Inositol Phosphorylceramide Sphingolipid Mannosylation Induces Plant Immune Responses and Reduces Cellulose Content in Arabidopsis. Plant Cell, 2016, 28, 2991-3004.	6.6	71
3	Pectin methylesterase31 positively regulates salt stress tolerance in Arabidopsis. Biochemical and Biophysical Research Communications, 2018, 496, 497-501.	2.1	57
4	Cell wall $\hat{l}^2$ -1,4-galactan regulated by the BPC1/BPC2-GALS1 module aggravates salt sensitivity in Arabidopsis thaliana. Molecular Plant, 2021, 14, 411-425.	8.3	54
5	Identification and functional characterization of three new terpene synthase genes involved in chemical defense and abiotic stresses in Santalum album. BMC Plant Biology, 2019, 19, 115.	3.6	43
6	Xyloglucan endotransglucosylase-hydrolase30 negatively affects salt tolerance in Arabidopsis. Journal of Experimental Botany, 2019, 70, 5495-5506.	4.8	38
7	GLUCOSAMINE INOSITOLPHOSPHORYLCERAMIDE TRANSFERASE1 (GINT1) Is a GlcNAc-Containing Glycosylinositol Phosphorylceramide Glycosyltransferase. Plant Physiology, 2018, 177, 938-952.	4.8	35
8	Transcriptome analysis provides insights into the non-methylated lignin synthesis in Paphiopedilum armeniacum seed. BMC Genomics, 2020, 21, 524.	2.8	19
9	Abscisic acid positively regulates <scp>l</scp> â€arabinose metabolism to inhibit seed germination through ABSCISIC ACID INSENSITIVE4â€mediated transcriptional promotions of <i>MUR4</i> in <i>Arabidopsis thaliana</i> New Phytologist, 2020, 225, 823-834.	7.3	18
10	Comparative Chloroplast Genomics and Phylogenetic Analysis of Thuniopsis and Closely Related Genera within Coelogyninae (Orchidaceae). Frontiers in Genetics, 2022, 13, 850201.	2.3	14
11	Insights on the aerobic biodegradation of agricultural wastes under simulated rapid composting conditions. Journal of Cleaner Production, 2019, 220, 688-697.	9.3	13
12	Functional characterization of an Indian sandalwood (Santalum album L.) dual-localized bifunctional nerolidol/linalool synthase gene involved in stress response. Phytochemistry, 2021, 183, 112610.	2.9	12
13	Elicitors Modulate Young Sandalwood (Santalum album L.) Growth, Heartwood Formation, and Concrete Oil Synthesis. Plants, 2021, 10, 339.	3 <b>.</b> 5	10
14	BASIC PENTACYSTEINE2 negatively regulates osmotic stress tolerance by modulating LEA4-5 expression in Arabidopsis thaliana. Plant Physiology and Biochemistry, 2021, 168, 373-380.	5.8	9
15	Exogenous GA3 promotes flowering in <i>Paphiopedilum callosum</i> (Orchidaceae) through bolting and lateral flower development regulation. Horticulture Research, 2022, 9, .	<b>6.</b> 3	9
16	The <i>Arabidopsis thaliana</i> nucleotide sugar transporter GONST2 is a functional homolog of GONST1. Plant Direct, 2021, 5, e00309.	1.9	7
17	Characterization of embryo and protocorm development of Paphiopedilum spicerianum. Plant Physiology and Biochemistry, 2021, 167, 1024-1034.	5 <b>.</b> 8	6
18	Ovule Development and in Planta Transformation of Paphiopedilum Maudiae by Agrobacterium-Mediated Ovary-Injection. International Journal of Molecular Sciences, 2021, 22, 84.	4.1	6

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19	Characterization of phytohormone and transcriptome profiles during protocorm-like bodies development of Paphiopedilum. BMC Genomics, 2021, 22, 806.	2.8	6
20	Abscisic Acid Inhibits Asymbiotic Germination of Immature Seeds of Paphiopedilum armeniacum. International Journal of Molecular Sciences, 2020, 21, 9561.	4.1	5
21	Transcriptomic analyses provide insight into adventitious root formation of Euryodendron excelsum H. T. Chang during ex vitro rooting. Plant Cell, Tissue and Organ Culture, 2022, 148, 649-666.	2.3	4
22	Cytological, Biochemical, and Transcriptomic Analyses of a Novel Yellow Leaf Variation in a Paphiopedilum (Orchidaceae) SCBG COP15. Genes, 2022, 13, 71.	2.4	4
23	Characterization of the complete chloroplast genome of Coelogyne fimbriata (Orchidaceae). Mitochondrial DNA Part B: Resources, 2020, 5, 3507-3509.	0.4	1
24	<strong>Reconsideration of the taxonomic status of <em>Bulbophyllum obtusangulum</em> (Orchidaceae) from southern China</strong> . Phytotaxa, 2021, 494, 219-224.	0.3	0