

# Jingye Fu

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/9639957/publications.pdf>

Version: 2024-02-01

10  
papers

393  
citations

1307594

7  
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1372567

10  
g-index

10  
all docs

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docs citations

10  
times ranked

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citing authors

#	ARTICLE	IF	CITATIONS
1	Transcriptional Factors Regulate Plant Stress Responses Through Mediating Secondary Metabolism. <i>Genes</i> , 2020, 11, 346.	2.4	138
2	A Tandem Array of <i>ent</i> -Kaurene Synthases in Maize with Roles in Gibberellin and More Specialized Metabolism. <i>Plant Physiology</i> , 2016, 170, 742-751.	4.8	81
3	ZmWRKY79 positively regulates maize phytoalexin biosynthetic gene expression and is involved in stress response. <i>Journal of Experimental Botany</i> , 2018, 69, 497-510.	4.8	51
4	Maize WRKY Transcription Factor ZmWRKY79 Positively Regulates Drought Tolerance through Elevating ABA Biosynthesis. <i>International Journal of Molecular Sciences</i> , 2021, 22, 10080.	4.1	38
5	Maize transcription factor ZmEREB20 enhanced salt tolerance in transgenic Arabidopsis. <i>Plant Physiology and Biochemistry</i> , 2021, 159, 257-267.	5.8	28
6	ZmMYC2 exhibits diverse functions and enhances JA signaling in transgenic Arabidopsis. <i>Plant Cell Reports</i> , 2020, 39, 273-288.	5.6	23
7	Rice contains a biosynthetic gene cluster associated with production of the casbane-type diterpenoid phytoalexin <i>ent</i> - <i>10</i> - <i>oxodepressin</i> . <i>New Phytologist</i> , 2021, 231, 85-93.	7.3	21
8	A Wheat $\beta$ -Patchoulene Synthase Confers Resistance against Herbivory in Transgenic Arabidopsis. <i>Genes</i> , 2019, 10, 441.	2.4	5
9	Probing Enzymatic Structure and Function in the Dihydroxylating Sesquiterpene Synthase ZmEDS. <i>Biochemistry</i> , 2020, 59, 2660-2666.	2.5	5
10	Promoter Variation Results in Differential Phytoalexin Accumulation in Two Maize Inbred Lines. <i>Plant Molecular Biology Reporter</i> , 2020, 38, 165-174.	1.8	3