

# Fangqian Chen

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/9946132/publications.pdf>

Version: 2024-02-01

12  
papers

170  
citations

1163117

8  
h-index

1199594

12  
g-index

12  
all docs

12  
docs citations

12  
times ranked

166  
citing authors

#	ARTICLE	IF	CITATIONS
1	HBI1 acts downstream of ERECTA and SWR1 in regulating inflorescence architecture through the activation of the brassinosteroid and auxin signaling pathways. <i>New Phytologist</i> , 2021, 229, 414-428.	7.3	17
2	ERECTA signaling regulates plant immune responses via chromatin-mediated promotion of <i>WRKY33</i> binding to target genes. <i>New Phytologist</i> , 2021, 230, 737-756.	7.3	20
3	Spatiotemporal control of miR398 biogenesis, via chromatin remodeling and kinase signaling, ensures proper ovule development. <i>Plant Cell</i> , 2021, 33, 1530-1553.	6.6	16
4	The bZIP transcription factor GmbZIP15 facilitates resistance against <i>Sclerotinia sclerotiorum</i> and <i>Phytophthora sojae</i> infection in soybean. <i>IScience</i> , 2021, 24, 102642.	4.1	10
5	SDG2 regulates <i>Arabidopsis</i> inflorescence architecture through SWR1-ERECTA signaling pathway. <i>IScience</i> , 2021, 24, 103236.	4.1	9
6	Genome-Wide Identification and Evaluation of New Reference Genes in Pineapple ( <i>Ananas comosus</i> L.) during Stamen and Ovule Development. <i>Tropical Plant Biology</i> , 2020, 13, 371-381.	1.9	4
7	Genome-wide investigation of calcium-dependent protein kinase gene family in pineapple: evolution and expression profiles during development and stress. <i>BMC Genomics</i> , 2020, 21, 72.	2.8	22
8	A Soybean bZIP Transcription Factor GmbZIP19 Confers Multiple Biotic and Abiotic Stress Responses in Plant. <i>International Journal of Molecular Sciences</i> , 2020, 21, 4701.	4.1	21
9	Genome-wide identification and expression analysis of the ERF transcription factor family in pineapple ( <i>Ananas comosus</i> (L.) Merr.). <i>PeerJ</i> , 2020, 8, e10014.	2.0	10
10	Identification and expression analysis of the DREB transcription factor family in pineapple ( <i>Ananas</i> ) <i>Tj ETQq0 0 0,rgBT /Overlock 10 T</i>	2.6	21
11	Genome-Wide Identification and Expression Analysis of the NAC Transcription Factor Family in Pineapple. <i>Tropical Plant Biology</i> , 2019, 12, 255-267.	1.9	7
12	Genome-Wide Classification and Evolutionary and Functional Analyses of the VQ Family. <i>Tropical Plant Biology</i> , 2019, 12, 117-131.	1.9	13