

# Xiurong Zhang

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5929527/publications.pdf>

Version: 2024-02-01

9  
papers

118  
citations

1478505

6  
h-index

1372567

10  
g-index

10  
all docs

10  
docs citations

10  
times ranked

85  
citing authors

#	ARTICLE	IF	CITATIONS
1	Identification of Peanut Aux/IAA Genes and Functional Prediction during Seed Development and Maturation. <i>Plants</i> , 2022, 11, 472.	3.5	4
2	Genome-Wide Identification of Auxin Response Factors in Peanut ( <i>Arachis hypogaea</i> L.) and Functional Analysis in Root Morphology. <i>International Journal of Molecular Sciences</i> , 2022, 23, 5309.	4.1	5
3	Genome-wide Identification and Abiotic Stress Response Pattern Analysis of NF-Y Gene Family in Peanut ( <i>Arachis Hypogaea</i> L.). <i>Tropical Plant Biology</i> , 2021, 14, 329-344.	1.9	4
4	Transcriptome and Metabolome Analysis Unveil Anthocyanin Metabolism in Pink and Red Testa of Peanut ( <i>Arachis hypogaea</i> L.). <i>International Journal of Genomics</i> , 2021, 2021, 1-16.	1.6	13
5	AhABI4s Negatively Regulate Salt-Stress Response in Peanut. <i>Frontiers in Plant Science</i> , 2021, 12, 741641.	3.6	12
6	Characterization of glycerol-3-phosphate acyltransferase 9 (AhGPAT9) genes, their allelic polymorphism and association with oil content in peanut ( <i>Arachis hypogaea</i> L.). <i>Scientific Reports</i> , 2020, 10, 14648.	3.3	9
7	Establishment and evaluation of a peanut association panel and analysis of key nutritional traits. <i>Journal of Integrative Plant Biology</i> , 2018, 60, 195-215.	8.5	11
8	QTL mapping and marker analysis of main stem height and the first lateral branch length in peanut ( <i>Arachis hypogaea</i> L.). <i>Euphytica</i> , 2017, 213, 1.	1.2	39
9	Molecular analysis of the chloroplast Cu/Zn-SOD gene (AhCSD2) in peanut. <i>Crop Journal</i> , 2015, 3, 246-257.	5.2	14