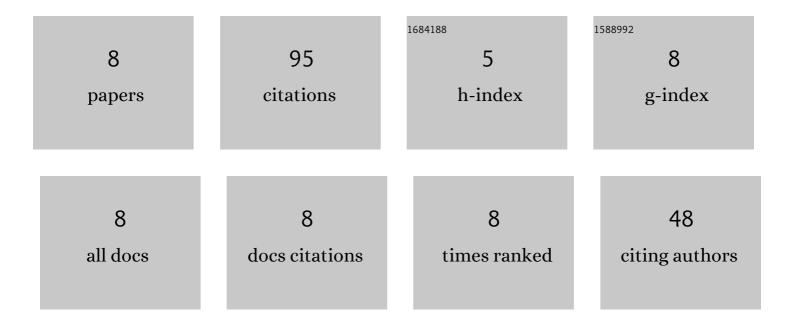
## Haichao Hu

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/2679040/publications.pdf Version: 2024-02-01



Нліснао Нії

#	Article	IF	CITATIONS
1	Transcriptome and Metabolome Dynamics Explain Aroma Differences between Green and Red Prickly Ash Fruit. Foods, 2021, 10, 391.	4.3	28
2	miRNAs and their target genes regulate the antioxidant system of Zanthoxylum bungeanum under drought stress. Plant Physiology and Biochemistry, 2020, 150, 196-203.	5.8	23
3	ZbAGL11, a class D MADS-box transcription factor of Zanthoxylum bungeanum, is involved in sporophytic apomixis. Horticulture Research, 2021, 8, 23.	6.3	14
4	Integrated Analysis of Metabolome and Transcriptome Data for Uncovering Flavonoid Components of Zanthoxylum bungeanum Maxim. Leaves Under Drought Stress. Frontiers in Nutrition, 2021, 8, 801244.	3.7	14
5	Genome-Wide Identification of the NAC Gene Family in Zanthoxylum bungeanum and Their Transcriptional Responses to Drought Stress. International Journal of Molecular Sciences, 2022, 23, 4769.	4.1	8
6	Small RNA sequencing provides candidate miRNA-target pairs for revealing the mechanism of apomixis in Zanthoxylum bungeanum. BMC Plant Biology, 2021, 21, 178.	3.6	5
7	Integrated LC-MS/MS and Transcriptome Sequencing Analysis Reveals the Mechanism of Color Formation During Prickly Ash Fruit Ripening. Frontiers in Nutrition, 2022, 9, 847823.	3.7	2
8	Pollination promotes ABA synthesis but not sexual reproduction in the apomictic species <i>Zanthoxylum bungeanum</i> Maxim Tree Physiology, 2021, 41, 1497-1509.	3.1	1