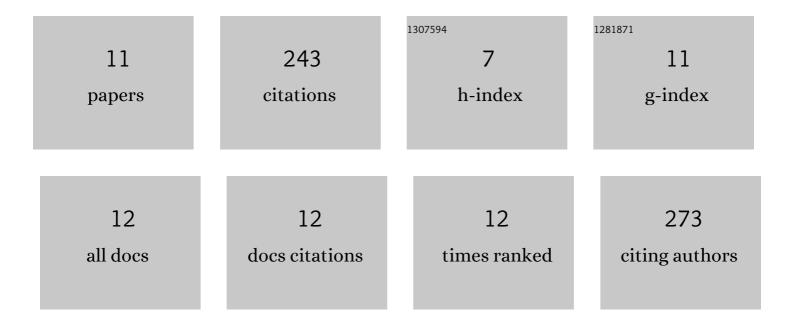
Yong Huang

List of Publications by Year in descending order

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Υονς Ημλις

#	Article	IF	CITATIONS
1	Genome-wide analysis and stress-responsive expression of CCCH zinc finger family genes in Brassica rapa. BMC Plant Biology, 2018, 18, 373.	3.6	73
2	Phylogenetic analysis and classification of the Brassica rapa SET-domain protein family. BMC Plant Biology, 2011, 11, 175.	3.6	42
3	Identification of SET Domain-Containing Proteins in Gossypium raimondii and Their Response to High Temperature Stress. Scientific Reports, 2016, 6, 32729.	3.3	28
4	Evolution and conservation of JmjC domain proteins in the green lineage. Molecular Genetics and Genomics, 2016, 291, 33-49.	2.1	27
5	Conservation and diversification of polycomb repressive complex 2 (PRC2) proteins in the green lineage. Briefings in Functional Genomics, 2017, 16, 106-119.	2.7	24
6	The evolutionary landscape of PRC1 core components in green lineage. Planta, 2016, 243, 825-846.	3.2	20
7	Genome-wide analysis of sulfotransferase genes and their responses to abiotic stresses in Chinese cabbage (Brassica rapa L.). PLoS ONE, 2019, 14, e0221422.	2.5	11
8	Systematic analysis of JmjC gene family and stressÂ-response expression of KDM5 subfamily genes in <i>Brassica napus</i> . PeerJ, 2021, 9, e11137.	2.0	7
9	Systematic analysis of CCCH zinc finger family in Brassica napus showed that BnRR-TZFs are involved in stress resistance. BMC Plant Biology, 2021, 21, 555.	3.6	5
10	Identification and charactering of APX genes provide new insights in abiotic stresses response in <i>Brassica napus</i> . PeerJ, 2022, 10, e13166.	2.0	4
11	An allotetraploid <i>Brassica napus</i> earlyâ€flowering mutant has <i>BnaFLC</i> 2â€regulated flowering. Journal of the Science of Food and Agriculture, 2013, 93, 3763-3768.	3.5	2