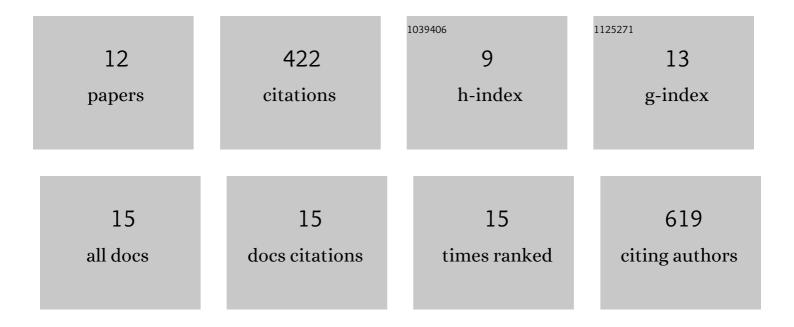
## Kai Hou

## List of Publications by Year in descending order

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KAI HOU

#	Article	IF	CITATIONS
1	Characterizing glycosyltransferases by a combination of sequencing platforms applied to the leaf tissues of Stevia rebaudiana. BMC Genomics, 2020, 21, 794.	1.2	14
2	Sequence variations and expression analysis of FAD2 gene among safflower materials with different linoleic acid content in seed oil. Acta Physiologiae Plantarum, 2019, 41, 1.	1.0	3
3	Bolting, an Important Process in Plant Development, Two Types in Plants. Journal of Plant Biology, 2019, 62, 161-169.	0.9	18
4	Exopolysaccharides from the fungal endophytic Fusarium sp. A14 isolated from Fritillaria unibracteata Hsiao et KC Hsia and their antioxidant and antiproliferation effects. Journal of Bioscience and Bioengineering, 2019, 127, 231-240.	1.1	13
5	Extraction, purification and antioxidation of a polysaccharide from Fritillaria unibracteata var. wabuensis. International Journal of Biological Macromolecules, 2018, 112, 1073-1083.	3.6	26
6	Endophytic fungus strain 28 isolated from Houttuynia cordata possesses wide-spectrum antifungal activity. Brazilian Journal of Microbiology, 2016, 47, 480-488.	0.8	31
7	Deep sequencing reveals transcriptome re-programming of Polygonum multiflorum thunb. roots to the elicitation with methyl jasmonate. Molecular Genetics and Genomics, 2016, 291, 337-348.	1.0	7
8	Transcriptome changes in Polygonum multiflorum Thunb. roots induced by methyl jasmonate. Journal of Zhejiang University: Science B, 2015, 16, 1027-1041.	1.3	6
9	Peimisine and peiminine production by endophytic fungus Fusarium sp. isolated from Fritillaria unibracteata var. wabensis. Phytomedicine, 2014, 21, 1104-1109.	2.3	32
10	RNA-Seq for gene identification and transcript profiling of three Stevia rebaudiana genotypes. BMC Genomics, 2014, 15, 571.	1.2	55
11	<i>SAUR36,</i> a SMALL AUXIN UP RNA Gene, Is Involved in the Promotion of Leaf Senescence in Arabidopsis   Â. Plant Physiology, 2013, 161, 1002-1009.	2.3	173
12	Molecular Cloning and Expression Analysis of Genes Encoding Two Microsomal Oleate Desaturases (FAD2) from Safflower (Carthamus tinctorius L.). Plant Molecular Biology Reporter, 2012, 30, 139-148.	1.0	34