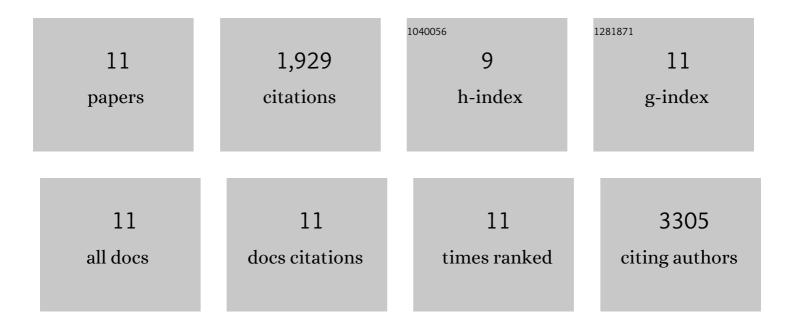
Yupeng Li

List of Publications by Year in descending order

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VUDENCL

#	Article	IF	CITATIONS
1	Draft genome sequence of pigeonpea (Cajanus cajan), an orphan legume crop of resource-poor farmers. Nature Biotechnology, 2012, 30, 83-89.	17.5	788
2	MicroRNAs as master regulators of the plant <i>NB-LRR</i> defense gene family via the production of phased, <i>trans</i> -acting siRNAs. Genes and Development, 2011, 25, 2540-2553.	5.9	668
3	Current state-of-art of sequencing technologies for plant genomics research. Briefings in Functional Genomics, 2012, 11, 3-11.	2.7	123
4	Gene Networks in Plant Biology: Approaches in Reconstruction and Analysis. Trends in Plant Science, 2015, 20, 664-675.	8.8	73
5	Identification and characterization of functional centromeres of the common bean. Plant Journal, 2013, 76, 47-60.	5.7	61
6	Fine mapping of Co-x, an anthracnose resistance gene to a highly virulent strain of Colletotrichum lindemuthianum in common bean. Theoretical and Applied Genetics, 2014, 127, 1653-1666.	3.6	59
7	The soybean (<i>Glycine max</i>) nodulationâ€suppressive <scp>CLE</scp> peptide, Gm <scp>RIC</scp> 1, functions interspecifically in common white bean (<i>Phaseolus vulgaris</i>), but not in a supernodulating line mutated in the receptor Pv <scp>NARK</scp> . Plant Biotechnology Journal, 2014, 12. 1085-1097.	8.3	55
8	Landscape and evolutionary dynamics of terminal repeat retrotransposons in miniature in plant genomes. Genome Biology, 2016, 17, 7.	8.8	50
9	Gene Network Reconstruction by Integration of Prior Biological Knowledge. G3: Genes, Genomes, Genetics, 2015, 5, 1075-1079.	1.8	39
10	A Bayesian gene network reveals insight into the JAK-STAT pathway in systemic lupus erythematosus. PLoS ONE, 2019, 14, e0225651.	2.5	8
11	Crowdsourcing the nodulation gene network discovery environment. BMC Bioinformatics, 2016, 17, 223.	2.6	5