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List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Auxinâ€mediated statolith production for root gravitropism. New Phytologist, 2019, 224, 761-774.	7.3	55
2	The K-segments of wheat dehydrin WZY2 are essential for its protective functions under temperature stress. Frontiers in Plant Science, 2015, 6, 406.	3.6	51
3	Abundant RNA editing sites of chloroplast protein-coding genes in Ginkgo biloba and an evolutionary pattern analysis. BMC Plant Biology, 2016, 16, 257.	3.6	49
4	The PIN gene family in cotton (Gossypium hirsutum): genome-wide identification and gene expression analyses during root development and abiotic stress responses. BMC Genomics, 2017, 18, 507.	2.8	46
5	Genome-wide identification of the GhARF gene family reveals that GhARF2 and GhARF18 are involved in cotton fibre cell initiation. Journal of Experimental Botany, 2018, 69, 4323-4337.	4.8	43
6	Two pivotal <scp>RNA</scp> editing sites in the mitochondrial <i>atp1</i> <scp>mRNA</scp> are required for <scp>ATP</scp> synthase to produce sufficient <scp>ATP</scp> for cotton fiber cell elongation. New Phytologist, 2018, 218, 167-182.	7.3	36
7	Comprehensive analysis of WOX genes uncovers that WOX13 is involved in phytohormone-mediated fiber development in cotton. BMC Plant Biology, 2019, 19, 312.	3.6	28
8	Plant transporters: roles in stress responses and effects on growth and development. Plant Growth Regulation, 2021, 93, 253-266.	3.4	17
9	Comprehensive analyses of ZFP gene family and characterization of expression profiles during plant hormone response in cotton. BMC Plant Biology, 2019, 19, 329.	3.6	12
10	GhYGL1d, a pentatricopeptide repeat protein, is required for chloroplast development in cotton. BMC Plant Biology, 2019, 19, 350.	3.6	10
11	GhPIPLC2D promotes cotton fiber elongation by enhancing ethylene biosynthesis. IScience, 2021, 24, 102199.	4.1	5