## Meng Yang

List of Publications by Year in descending order

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19	1,728 citations	759055	794469 19
papers	citations	h-index	g-index
19	19	19	2452
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	Reply to: Evaluating two different models of peanut's origin. Nature Genetics, 2020, 52, 560-563.	9.4	8
2	The genome of cultivated peanut provides insight into legume karyotypes, polyploid evolution and crop domestication. Nature Genetics, 2019, 51, 865-876.	9.4	398
3	Development of a massively parallel sequencing assay for investigating sequence polymorphisms of 15 short tandem repeats in a Chinese Northern Han population. Electrophoresis, 2018, 39, 2725-2731.	1.3	11
4	The calciumâ€dependent protein kinase (CDPK) and CDPKâ€related kinase gene families in <i>Hevea brasiliensis</i> àâ€"comparison with five other plant species in structure, evolution, and expression. FEBS Open Bio, 2017, 7, 4-24.	1.0	40
5	Genetic variability and forensic efficiency of 39 microsatellite loci in the Li ethnic group from Hainan Island in the South China Sea. Annals of Human Biology, 2017, 44, 467-474.	0.4	6
6	Development of a rapid 21â€plex autosomal STR typing system for forensic applications. Electrophoresis, 2016, 37, 2789-2799.	1.3	5
7	De novo Transcriptome Analysis Reveals Distinct Defense Mechanisms by Young and Mature Leaves of Hevea brasiliensis (Para Rubber Tree). Scientific Reports, 2016, 6, 33151.	1.6	40
8	Comparative genomics reveals adaptive evolution of Asian tapeworm in switching to a new intermediate host. Nature Communications, 2016, 7, 12845.	5 <b>.</b> 8	43
9	The rubber tree genome reveals new insights into rubber production and species adaptation. Nature Plants, 2016, 2, 16073.	4.7	324
10	Mechanisms of action for 2-phenylethanol isolated from Kloeckera apiculata in control of Penicillium molds of citrus fruits. BMC Microbiology, 2014, 14, 242.	1.3	98
11	Structure and expression profile of the sucrose synthase gene family in the rubber tree: indicative of roles in stress response and sucrose utilization in the laticifers. FEBS Journal, 2014, 281, 291-305.	2.2	62
12	Sequence and Expression Analyses of Ethylene Response Factors Highly Expressed in Latex Cells from Hevea brasiliensis. PLoS ONE, 2014, 9, e99367.	1.1	21
13	Genome sequence of the date palm Phoenix dactylifera L. Nature Communications, 2013, 4, 2274.	<b>5.</b> 8	248
14	The small RNA profile in latex from Hevea brasiliensis trees is affected by tapping panel dryness. Tree Physiology, 2013, 33, 1084-1098.	1.4	38
15	Complete Genome Sequence of Antarctic Bacterium Psychrobacter sp. Strain G. Genome Announcements, $2013,1,.$	0.8	7
16	A Complete Sequence and Transcriptomic Analyses of Date Palm (Phoenix dactylifera L.) Mitochondrial Genome. PLoS ONE, 2012, 7, e37164.	1,1	106
17	A comprehensive analysis of protein phosphatases in rice and Arabidopsis. Plant Systematics and Evolution, 2010, 289, 111-126.	0.3	7
18	Sequence Signatures of Nucleosome Positioning in Caenorhabditis elegans. Genomics, Proteomics and Bioinformatics, 2010, 8, 92-102.	3.0	11

#	Article	lF	CITATIONS
19	The Complete Chloroplast Genome Sequence of Date Palm (Phoenix dactylifera L.). PLoS ONE, 2010, 5, e12762.	1.1	255