## Qin Qiao

## List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8179501/publications.pdf

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11	389	1040056	1281871
papers	citations	h-index	g-index
11	11	11	561
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	Genomic analysis of field pennycress (Thlaspi arvense) provides insights into mechanisms of adaptation to high elevation. BMC Biology, 2021, 19, 143.	3.8	23
2	Evaluation of genetic diversity and population structure of Fragaria nilgerrensis using EST-SSR markers. Gene, 2021, 796-797, 145791.	2.2	9
3	Evolutionary history and pan-genome dynamics of strawberry ( <i>Fragaria (i) spp.). Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .</i>	7.1	43
4	Dynamic Changes of DNA Methylation During Wild Strawberry (Fragaria nilgerrensis) Tissue Culture. Frontiers in Plant Science, 2021, 12, 765383.	3.6	5
5	Reply to: Revisiting the origin of octoploid strawberry. Nature Genetics, 2020, 52, 5-7.	21.4	44
6	Genome of <i>Crucihimalaya himalaica</i> , a close relative of <i>Arabidopsis</i> , shows ecological adaptation to high altitude. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 7137-7146.	7.1	108
7	Enriched networks â€~nucleoside/nucleotide and ribonucleoside/ribonucleotide metabolic processes' and â€~response to stimulus' potentially conferred to drought adaptation of the epiphytic orchid Dendrobium wangliangii. Physiology and Molecular Biology of Plants, 2019, 25, 31-45.	3.1	13
8	Probing the transcriptome of Aconitum carmichaelii reveals the candidate genes associated with the biosynthesis of the toxic aconitine-type C19-diterpenoid alkaloids. Phytochemistry, 2018, 152, 113-124.	2.9	20
9	Comparative Transcriptomics of Strawberries (Fragaria spp.) Provides Insights into Evolutionary Patterns. Frontiers in Plant Science, 2016, 7, 1839.	3.6	33
10	Transcriptome sequencing of Crucihimalaya himalaica (Brassicaceae) reveals how Arabidopsis close relative adapt to the Qinghai-Tibet Plateau. Scientific Reports, 2016, 6, 21729.	3.3	47
11	Identification of Glutathione S-Transferase (GST) Genes from a Dark Septate Endophytic Fungus (Exophiala pisciphila) and Their Expression Patterns under Varied Metals Stress. PLoS ONE, 2015, 10, e0123418.	2.5	44