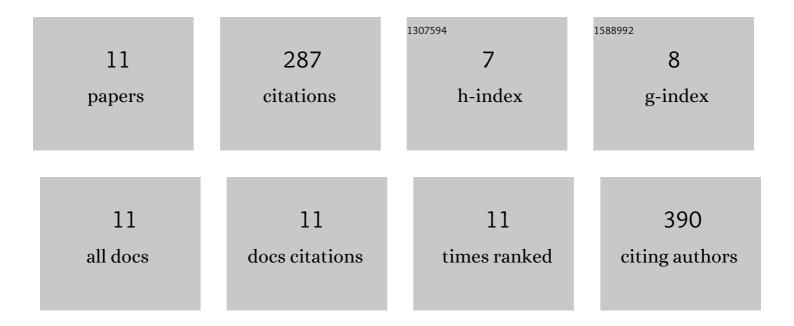
Jihong Li

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/9992671/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Integration of small <scp>RNA</scp> s, degradome and transcriptome sequencing in hyperaccumulator <i>Sedum alfredii</i> uncovers a complex regulatory network and provides insights into cadmium phytoremediation. Plant Biotechnology Journal, 2016, 14, 1470-1483.	8.3	96
2	Construction of a high-density genetic map using specific length amplified fragment markers and identification of a quantitative trait locus for anthracnose resistance in walnut (Juglans regia L.). BMC Genomics, 2015, 16, 614.	2.8	72
3	PzTAC and PzLAZY from a narrow-crown poplar contribute to regulation of branch angles. Plant Physiology and Biochemistry, 2017, 118, 571-578.	5.8	38
4	Identification and Characterization of MicroRNAs in Ginkgo biloba var. epiphylla Mak. PLoS ONE, 2015, 10, e0127184.	2.5	37
5	Transcriptome Profile Analysis from Different Sex Types of Ginkgo biloba L Frontiers in Plant Science, 2016, 7, 871.	3.6	21
6	PagGRF12a interacts with PagGIF1b to regulate secondary xylem development through modulating <i>PagXND1a</i> expression in <i>Populus alba</i> A— <i>P. glandulosa</i> . Journal of Integrative Plant Biology, 2021, 63, 1683-1694.	8.5	13
7	Transcriptome sequencing of active buds from Populus deltoides CL. and Populus × zhaiguanheibaiyang reveals phytohormones involved in branching. Genomics, 2019, 111, 700-709.	2.9	8
8	Transcriptome Analysis of Active Axillary Buds from Narrow-crown and Broad-crown Poplars Provides Insight into the Phytohormone Regulatory Network for Branching Angle. Plant Molecular Biology Reporter, 2021, 39, 595-606.	1.8	2
9	Natural forest conservation hierarchical program with neural network. Frontiers of Forestry in China: Selected Publications From Chinese Universities, 2006, 1, 318-323.	0.2	0
10	\$\$sqrt 2 \$\$ Rule for Controlling the Tree Pattern in Forest Cut. Frontiers of Biology in China: Selected Publications From Chinese Universities, 2006, 1, 71-75.	0.2	0
11	Construction of a high-density genetic map using specific-length amplified fragment markers and identification of QTLs for branching angle in poplar. Molecular Genetics and Genomics, 2022, 297,	2.1	0