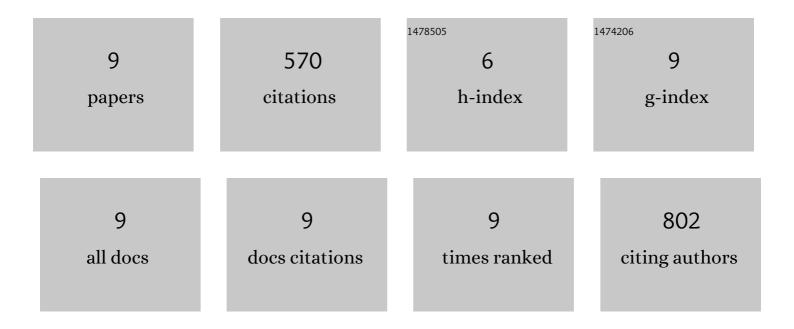
## Wenchun Luo

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/11411684/publications.pdf Version: 2024-02-01



WENCHUNLUO

#	Article	IF	CITATIONS
1	A Comparative Analysis of Transcription Networks Active in Juvenile and Mature Wood in Populus. Frontiers in Plant Science, 2021, 12, 675075.	3.6	7
2	Evolution of RGF/GLV/CLEL Peptide Hormones and Their Roles in Land Plant Growth and Regulation. International Journal of Molecular Sciences, 2021, 22, 13372.	4.1	1
3	Genome-Wide Identification of Long Noncoding RNAs and Their Responses to Salt Stress in Two Closely Related Poplars. Frontiers in Genetics, 2019, 10, 777.	2.3	38
4	A Novel Gene Coding $\hat{I}^3$ -Aminobutyric Acid Transporter May Improve the Tolerance of Populus euphratica to Adverse Environments. Frontiers in Plant Science, 2019, 10, 1083.	3.6	12
5	Transcriptomic Analysis of Seed Germination Under Salt Stress in Two Desert Sister Species (Populus) Tj ETQq1 1	0,784314 2.3	∙rgBT /Over
6	Transcriptome analysis of four poplars exposed to continuous salinity stress. Biochemical Systematics and Ecology, 2017, 70, 311-319.	1.3	4
7	The willow genome and divergent evolution from poplar after the common genome duplication. Cell Research, 2014, 24, 1274-1277.	12.0	148
8	Transcriptome dynamics of a desert poplar (Populus pruinosa) in response to continuous salinity stress. Plant Cell Reports, 2014, 33, 1565-1579.	5.6	42
9	Genomic insights into salt adaptation in a desert poplar. Nature Communications, 2013, 4, 2797.	12.8	286