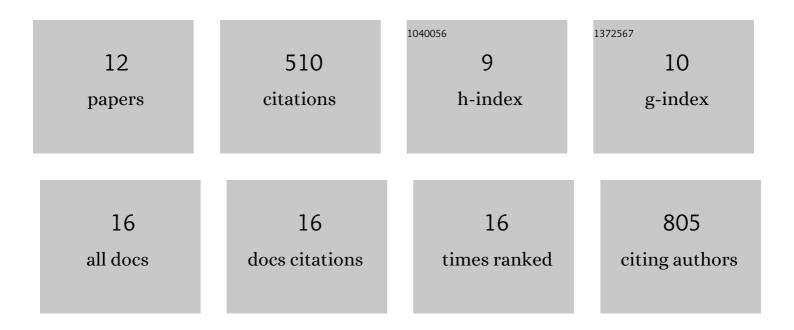
## Li Lei

## List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/338787/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Barcoding the kingdom Plantae: new <scp>PCR</scp> primers for <i><scp>ITS</scp></i> regions of plants with improved universality and specificity. Molecular Ecology Resources, 2016, 16, 138-149.	4.8	280
2	Plant Pan-Genomics Comes of Age. Annual Review of Plant Biology, 2021, 72, 411-435.	18.7	56
3	Expansion and diversification of the SET domain gene family following whole-genome duplications in Populus trichocarpa. BMC Evolutionary Biology, 2012, 12, 51.	3.2	49
4	Comparative Genomics Approaches Accurately Predict Deleterious Variants in Plants. G3: Genes, Genomes, Genetics, 2018, 8, 3321-3329.	1.8	36
5	Development of a Multiparent Population for Genetic Mapping and Allele Discovery in Six-Row Barley. Genetics, 2019, 213, 595-613.	2.9	23
6	Environmental Association Identifies Candidates for Tolerance to Low Temperature and Drought. G3: Genes, Genomes, Genetics, 2019, 9, 3423-3438.	1.8	18
7	Reevaluation of Two Quantitative Trait Loci for Type II Resistance to Fusarium Head Blight in Wheat Germplasm PI 672538. Phytopathology, 2017, 107, 92-99.	2.2	14
8	Collinearity Analysis and High-Density Genetic Mapping of the Wheat Powdery Mildew Resistance Gene Pm40 in Pl 672538. PLoS ONE, 2016, 11, e0164815.	2.5	12
9	Plant organ evolution revealed by phylotranscriptomics in Arabidopsis thaliana. Scientific Reports, 2017, 7, 7567.	3.3	11
10	Temporal fitness fluctuations in experimental Arabidopsis thaliana populations. PLoS ONE, 2017, 12, e0178990.	2.5	9
11	Sequence-related amplified polymorphisms (SRAPs) reveal genetic diversity and variation regions in upland cotton (Gossypium hirsutum L.) in China. Crop Breeding and Applied Biotechnology, 2018, 18, 155-160.	0.4	0
12	Identification of three new resources of resistance to Fusarium head blight in wheat. Czech Journal of Genetics and Plant Breeding, 2019, 55, 15-19.	0.8	0