

# Liang Du

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/9528661/publications.pdf>

Version: 2024-02-01

10  
papers

546  
citations

1163117

8  
h-index

1372567

10  
g-index

11  
all docs

11  
docs citations

11  
times ranked

762  
citing authors

#	ARTICLE	IF	CITATIONS
1	Origin and diversification of leucine-rich repeat receptor-like protein kinase (LRR-RLK) genes in plants. <i>BMC Evolutionary Biology</i> , 2017, 17, 47.	3.2	211
2	The Ubiquitin Receptor DA1 Regulates Seed and Organ Size by Modulating the Stability of the Ubiquitin-Specific Protease UBP15/SOD2 in <i>Arabidopsis</i> . <i>Plant Cell</i> , 2014, 26, 665-677.	6.6	149
3	Transcription Factors SOD7/NGAL2 and DPA4/NGAL3 Act Redundantly to Regulate Seed Size by Directly Repressing <i>KLU</i> Expression in <i>Arabidopsis thaliana</i> . <i>Plant Cell</i> , 2015, 27, 620-632.	6.6	77
4	The Pentatricopeptide Repeat Protein Pigment-Defective Mutant2 is Involved in the Regulation of Chloroplast Development and Chloroplast Gene Expression in <i>Arabidopsis</i> . <i>Plant and Cell Physiology</i> , 2017, 58, 747-759.	3.1	38
5	Development and chemical characterization of Casparian strips in the roots of Chinese fir ( <i>Cunninghamia lanceolata</i> ). <i>Trees - Structure and Function</i> , 2019, 33, 827-836.	1.9	22
6	Genome-Wide Identification and Characterization of the UBP Gene Family in Moso Bamboo ( <i>Phyllostachys edulis</i> ). <i>International Journal of Molecular Sciences</i> , 2019, 20, 4309.	4.1	14
7	Protein partners of plant ubiquitin-specific proteases (UBPs). <i>Plant Physiology and Biochemistry</i> , 2019, 145, 227-236.	5.8	13
8	Genome-Wide Identification and Characterization of Hexokinase Genes in Moso Bamboo ( <i>Phyllostachys edulis</i> ). <i>Frontiers in Plant Science</i> , 2020, 11, 600.	3.6	11
9	Transcriptome-wide identification of miRNA targets and a TAS3-homologous gene in <i>Populus</i> by degradome sequencing. <i>Genes and Genomics</i> , 2019, 41, 849-861.	1.4	7
10	The DUB family in <i>Populus</i> : identification, characterization, evolution and expression patterns. <i>BMC Genomics</i> , 2021, 22, 541.	2.8	4