

# Liang Guo

## List of Publications by Year in descending order

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Version: 2024-02-01

73  
papers

3,510  
citations

136950

32  
h-index

161849

54  
g-index

80  
all docs

80  
docs citations

80  
times ranked

3422  
citing authors

#	ARTICLE	IF	CITATIONS
1	Eight high-quality genomes reveal pan-genome architecture and ecotype differentiation of <i>Brassica napus</i> . <i>Nature Plants</i> , 2020, 6, 34-45.	9.3	449
2	Plant phospholipases D and C and their diverse functions in stress responses. <i>Progress in Lipid Research</i> , 2016, 62, 55-74.	11.6	288
3	Cytosolic Glyceraldehyde-3-Phosphate Dehydrogenases Interact with Phospholipase D $\hat{1}$ to Transduce Hydrogen Peroxide Signals in the <i>Arabidopsis</i> Response to Stress. <i>Plant Cell</i> , 2012, 24, 2200-2212.	6.6	202
4	Comparative transcriptomic analysis uncovers the complex genetic network for resistance to <i>Sclerotinia sclerotiorum</i> in <i>Brassica napus</i> . <i>Scientific Reports</i> , 2016, 6, 19007.	3.3	126
5	Phosphatidic Acid Binds and Stimulates <i>Arabidopsis</i> Sphingosine Kinases. <i>Journal of Biological Chemistry</i> , 2011, 286, 13336-13345.	3.4	109
6	Genome- and transcriptome-wide association studies provide insights into the genetic basis of natural variation of seed oil content in <i>Brassica napus</i> . <i>Molecular Plant</i> , 2021, 14, 470-487.	8.3	107
7	Connections between Sphingosine Kinase and Phospholipase D in the Abscisic Acid Signaling Pathway in <i>Arabidopsis</i> . <i>Journal of Biological Chemistry</i> , 2012, 287, 8286-8296.	3.4	99
8	Patatin-Related Phospholipase pPLAII $\hat{2}$ -Induced Changes in Lipid Metabolism Alter Cellulose Content and Cell Elongation in <i>Arabidopsis</i> . <i>Plant Cell</i> , 2011, 23, 1107-1123.	6.6	94
9	Emerging Roles of Sphingolipid Signaling in Plant Response to Biotic and Abiotic Stresses. <i>Molecular Plant</i> , 2018, 11, 1328-1343.	8.3	87
10	Phosphatidic Acid Interacts with a MYB Transcription Factor and Regulates Its Nuclear Localization and Function in <i>Arabidopsis</i> . <i>Plant Cell</i> , 2014, 25, 5030-5042.	6.6	80
11	Cytosolic Phosphorylating Glyceraldehyde-3-Phosphate Dehydrogenases Affect <i>Arabidopsis</i> Cellular Metabolism and Promote Seed Oil Accumulation. <i>Plant Cell</i> , 2014, 26, 3023-3035.	6.6	80
12	BnTIR: an online transcriptome platform for exploring RNA-seq libraries for oil crop <i>Brassica napus</i> . <i>Plant Biotechnology Journal</i> , 2021, 19, 1895-1897.	8.3	68
13	Spatial analysis of lipid metabolites and expressed genes reveals tissue-specific heterogeneity of lipid metabolism in high- and low-oil <i>Brassica napus</i> L. seeds. <i>Plant Journal</i> , 2018, 94, 915-932.	5.7	66
14	Roles of the <i>Brassica napus</i> DELLA Protein BnaA6.RGA, in Modulating Drought Tolerance by Interacting With the ABA Signaling Component BnaA10.ABF2. <i>Frontiers in Plant Science</i> , 2020, 11, 577.	3.6	66
15	Phosphatidic Acid Binds to Cytosolic Glyceraldehyde-3-phosphate Dehydrogenase and Promotes Its Cleavage in <i>Arabidopsis</i> . <i>Journal of Biological Chemistry</i> , 2013, 288, 11834-11844.	3.4	65
16	Crosstalk between Phospholipase D and Sphingosine Kinase in Plant Stress Signaling. <i>Frontiers in Plant Science</i> , 2012, 3, 51.	3.6	55
17	The genome of jojoba ( <i>Simmondsia chinensis</i> ): A taxonomically isolated species that directs wax ester accumulation in its seeds. <i>Science Advances</i> , 2020, 6, eaay3240.	10.3	53
18	Different effects of phospholipase D $\hat{1}$ 2 and non-specific phospholipase C4 on lipid remodeling and root hair growth in <i>Arabidopsis</i> response to phosphate deficiency. <i>Plant Journal</i> , 2018, 94, 315-326.	5.7	52

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19	Multiple GmWRI1s are redundantly involved in seed filling and nodulation by regulating plastidic glycolysis, lipid biosynthesis and hormone signalling in soybean ( <i>Glycine max</i> ). <i>Plant Biotechnology Journal</i> , 2020, 18, 155-171.	8.3	52
20	The functions of phospholipases and their hydrolysis products in plant growth, development and stress responses. <i>Progress in Lipid Research</i> , 2022, 86, 101158.	11.6	52
21	BnPIR: <i>Brassica napus</i> pan-genome information resource for 1689 accessions. <i>Plant Biotechnology Journal</i> , 2021, 19, 412-414.	8.3	51
22	<i>Arabidopsis</i> GDSL1 overexpression enhances rapeseed <i>Sclerotinia sclerotiorum</i> resistance and the functional identification of its homolog in <i>Brassica napus</i> . <i>Plant Biotechnology Journal</i> , 2020, 18, 1255-1270.	8.3	48
23	Nuclear moonlighting of cytosolic glyceraldehyde-3-phosphate dehydrogenase regulates <i>Arabidopsis</i> response to heat stress. <i>Nature Communications</i> , 2020, 11, 3439.	12.8	48
24	Molecular Basis of Plant Oil Biosynthesis: Insights Gained From Studying the WRINKLED1 Transcription Factor. <i>Frontiers in Plant Science</i> , 2020, 11, 24.	3.6	47
25	Phospholipase D $\gamma$ negatively regulates plant thermotolerance by destabilizing cortical microtubules in <i>Arabidopsis</i> . <i>Plant, Cell and Environment</i> , 2017, 40, 2220-2235.	5.7	45
26	Long-read sequencing reveals widespread intragenic structural variants in a recent allopolyploid crop plant. <i>Plant Biotechnology Journal</i> , 2021, 19, 240-250.	8.3	45
27	An efficient <i>Agrobacterium</i> -mediated transformation method using hypocotyl as explants for <i>Brassica napus</i> . <i>Molecular Breeding</i> , 2020, 40, 1.	2.1	43
28	Genome-wide and transcriptome-wide association studies reveal the genetic basis and the breeding history of seed glucosinolate content in <i>Brassica napus</i> . <i>Plant Biotechnology Journal</i> , 2022, 20, 211-225.	8.3	43
29	Strigolactones promote rhizobia interaction and increase nodulation in soybean ( <i>Glycine max</i> ). <i>Microbial Pathogenesis</i> , 2018, 114, 420-430.	2.9	41
30	<i>De novo</i> transcriptome sequencing and metabolite profiling analyses reveal the complex metabolic genes involved in the terpenoid biosynthesis in Blue Anise Sage ( <i>Salvia guaranitica</i> L.). <i>DNA Research</i> , 2018, 25, 597-617.	3.4	41
31	Characterization of Fatty Acid Exporters involved in fatty acid transport for oil accumulation in the green alga <i>Chlamydomonas reinhardtii</i> . <i>Biotechnology for Biofuels</i> , 2019, 12, 14.	6.2	40
32	Crop height estimation based on UAV images: Methods, errors, and strategies. <i>Computers and Electronics in Agriculture</i> , 2021, 185, 106155.	7.7	40
33	Genome-Wide Analysis of Phospholipase D Gene Family and Profiling of Phospholipids under Abiotic Stresses in <i>Brassica napus</i> . <i>Plant and Cell Physiology</i> , 2019, 60, 1556-1566.	3.1	39
34	A reactive oxygen species burst causes haploid induction in maize. <i>Molecular Plant</i> , 2022, 15, 943-955.	8.3	39
35	Transcriptional regulation of oil biosynthesis in seed plants: Current understanding, applications, and perspectives. <i>Plant Communications</i> , 2022, 3, 100328.	7.7	39
36	Phospholipase D $\mu$ enhances <i>Brassica napus</i> growth and seed production in response to nitrogen availability. <i>Plant Biotechnology Journal</i> , 2016, 14, 926-937.	8.3	35

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37	Nonspecific phospholipase C4 hydrolyzes phosphosphingolipids and sustains plant root growth during phosphate deficiency. <i>Plant Cell</i> , 2021, 33, 766-780.	6.6	31
38	TEOSINTE BRANCHED1/CYCLOIDEA/PROLIFERATING CELL FACTOR4 Interacts with WRINKLED1 to Mediate Seed Oil Biosynthesis. <i>Plant Physiology</i> , 2020, 184, 658-665.	4.8	29
39	ISSR Analysis of the Genetic Diversity of the Endangered Species <i>Sinopodophyllum hexandrum</i> (Royle) Ying from Western Sichuan Province, China. <i>Journal of Integrative Plant Biology</i> , 2006, 48, 1140-1146.	8.5	28
40	Transcriptome analysis reveals genes commonly responding to multiple abiotic stresses in rapeseed. <i>Molecular Breeding</i> , 2019, 39, 1.	2.1	28
41	An efficient and comprehensive plant glycerolipids analysis approach based on high-performance liquid chromatography-quadrupole time-of-flight mass spectrometer. <i>Plant Direct</i> , 2019, 3, e00183.	1.9	26
42	Arabidopsis phospholipase D1 and D1' oppositely modulate EDS1- and SA-independent basal resistance against adapted powdery mildew. <i>Journal of Experimental Botany</i> , 2018, 69, 3675-3688.	4.8	23
43	Multi-omics analysis dissects the genetic architecture of seed coat content in <i>Brassica napus</i> . <i>Genome Biology</i> , 2022, 23, 86.	8.8	23
44	PLD: Phospholipase Ds in Plant Signaling. <i>Signaling and Communication in Plants</i> , 2014, , 3-26.	0.7	22
45	Nonspecific phospholipase C6 increases seed oil production in oilseed Brassicaceae plants. <i>New Phytologist</i> , 2020, 226, 1055-1073.	7.3	22
46	Development and screening of EMS mutants with altered seed oil content or fatty acid composition in <i>Brassica napus</i> . <i>Plant Journal</i> , 2020, 104, 1410-1422.	5.7	21
47	Genome-Wide Association Studies of Salt Tolerance at Seed Germination and Seedling Stages in <i>Brassica napus</i> . <i>Frontiers in Plant Science</i> , 2021, 12, 772708.	3.6	20
48	AFLP Analysis of Genetic Diversity of the Endangered Species <i>Sinopodophyllum hexandrum</i> in the Tibetan Region of Sichuan Province, China. <i>Biochemical Genetics</i> , 2006, 44, 44-57.	1.7	19
49	Two Plastid Fatty Acid Exporters Contribute to Seed Oil Accumulation in <i>Arabidopsis</i> . <i>Plant Physiology</i> , 2020, 182, 1910-1919.	4.8	19
50	CRISPR/Cas9-Targeted Mutagenesis of <i>BnaFAE1</i> Genes Confers Low-Erucic Acid in <i>Brassica napus</i> . <i>Frontiers in Plant Science</i> , 2022, 13, 848723.	3.6	18
51	Genome wide characterization of phospholipase A & C families and pattern of lysolipids and diacylglycerol changes under abiotic stresses in <i>Brassica napus</i> L. <i>Plant Physiology and Biochemistry</i> , 2020, 147, 101-112.	5.8	17
52	Membrane glycerolipidome of soybean root hairs and its response to nitrogen and phosphate availability. <i>Scientific Reports</i> , 2016, 6, 36172.	3.3	16
53	Oil plant genomes: current state of the science. <i>Journal of Experimental Botany</i> , 2022, 73, 2859-2874.	4.8	16
54	Genome-Wide Association Mapping Unravels the Genetic Control of Seed Vigor under Low-Temperature Conditions in Rapeseed ( <i>Brassica napus</i> L.). <i>Plants</i> , 2021, 10, 426.	3.5	15

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55	DELLA proteins BnaA6.RGA and BnaC7.RGA negatively regulate fatty acid biosynthesis by interacting with BnaLEC1s in <i>Brassica napus</i> . <i>Plant Biotechnology Journal</i> , 2021, 19, 2011-2026.	8.3	15
56	The <i>Brassica napus</i> fatty acid exporter FAX1-1 contributes to biological yield, seed oil content, and oil quality. <i>Biotechnology for Biofuels</i> , 2021, 14, 190.	6.2	15
57	Involvement of abscisic acid, ABI5, and PPC2 in plant acclimation to low CO <sub>2</sub> . <i>Journal of Experimental Botany</i> , 2020, 71, 4093-4108.	4.8	13
58	Acylation of non-specific phospholipase C4 determines its function in plant response to phosphate deficiency. <i>Plant Journal</i> , 2021, 106, 1647-1659.	5.7	13
59	BnVIR: bridging the genotype-phenotype gap to accelerate mining of candidate variations underlying agronomic traits in <i>Brassica napus</i> . <i>Molecular Plant</i> , 2022, 15, 779-782.	8.3	13
60	Heterogeneous Distribution of Erucic Acid in <i>Brassica napus</i> Seeds. <i>Frontiers in Plant Science</i> , 2020, 10, 1744.	3.6	12
61	Combining quantitative trait locus and co-expression analysis allowed identification of new candidates for oil accumulation in rapeseed. <i>Journal of Experimental Botany</i> , 2021, 72, 1649-1660.	4.8	12
62	Molecular Cloning, Characterization, and Expression of an $\Delta^6$ -3 Fatty Acid Desaturase Gene from <i>Sapium sebiferum</i> . <i>Journal of Bioscience and Bioengineering</i> , 2008, 106, 375-380.	2.2	11
63	Salt-responsive transcriptome analysis of canola roots reveals candidate genes involved in the key metabolic pathway in response to salt stress. <i>Scientific Reports</i> , 2022, 12, 1666.	3.3	10
64	Sunflower WRINKLED1 Plays a Key Role in Transcriptional Regulation of Oil Biosynthesis. <i>International Journal of Molecular Sciences</i> , 2022, 23, 3054.	4.1	10
65	Proteome-wide identification of S-sulphenylated cysteines in <i>Brassica napus</i> . <i>Plant, Cell and Environment</i> , 2021, 44, 3571-3582.	5.7	9
66	High-throughput unmanned aerial vehicle-based phenotyping provides insights into the dynamic process and genetic basis of rapeseed waterlogging response in the field. <i>Journal of Experimental Botany</i> , 2022, 73, 5264-5278.	4.8	7
67	<i>Brassica napus</i> BnaNTT1 modulates ATP homeostasis in plastids to sustain metabolism and growth. <i>Cell Reports</i> , 2022, 40, 111060.	6.4	7
68	The function of the WR11-TCP4 regulatory module in lipid biosynthesis. <i>Plant Signaling and Behavior</i> , 2020, 15, 1812878.	2.4	6
69	Genome-Wide Association Studies of Salt-Alkali Tolerance at Seedling and Mature Stages in <i>Brassica napus</i> . <i>Frontiers in Plant Science</i> , 2022, 13, 857149.	3.6	5
70	Transcriptome Analysis Reveals Genes of Flooding-Tolerant and Flooding-Sensitive Rapeseeds Differentially Respond to Flooding at the Germination Stage. <i>Plants</i> , 2021, 10, 693.	3.5	4
71	Genetic and Biochemical Investigation of Seed Fatty Acid Accumulation in <i>Arabidopsis</i> . <i>Frontiers in Plant Science</i> , 0, 13, .	3.6	4
72	Critical Roles of Mitochondrial Fatty Acid Synthesis in Tomato Development and Environmental Response. <i>Plant Physiology</i> , 0, , .	4.8	1

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73	Transcriptional regulation of oil biosynthesis in different parts of Wanyou 20 ( <i>Brassica napus</i> ) seeds. <i>Acta Agronomica Sinica</i> (China), 2019, 45, 381.	0.3	0