

# Shengguan Cai

## List of Publications by Year in descending order

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30  
papers

983  
citations

471509

17  
h-index

477307

29  
g-index

30  
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30  
docs citations

30  
times ranked

1275  
citing authors

#	ARTICLE	IF	CITATIONS
1	Vacuolar H <sup>+</sup> -pyrophosphatase HVP10 enhances salt tolerance via promoting Na <sup>+</sup> translocation into root vacuoles. <i>Plant Physiology</i> , 2022, 188, 1248-1263.	4.8	15
2	Time-Course Comparative Metabolome Analysis of Different Barley Varieties during Malting. <i>Journal of Agricultural and Food Chemistry</i> , 2022, 70, 2051-2059.	5.2	6
3	Evolution of rapid blue-light response linked to explosive diversification of ferns in angiosperm forests. <i>New Phytologist</i> , 2021, 230, 1201-1213.	7.3	33
4	Comprehensive dissection of primary metabolites in response to diverse abiotic stress in barley at seedling stage. <i>Plant Physiology and Biochemistry</i> , 2021, 161, 54-64.	5.8	20
5	Multi-Omics Analysis Reveals the Mechanism Underlying the Edaphic Adaptation in Wild Barley at Evolution Slope (Tabigha). <i>Advanced Science</i> , 2021, 8, e2101374.	11.2	14
6	The zinc finger transcription factor ATF1 regulates aluminum tolerance in barley. <i>Journal of Experimental Botany</i> , 2020, 71, 6512-6523.	4.8	13
7	Transcriptome-Based Analysis of Phosphite-Induced Resistance against Pathogens in Rice. <i>Plants</i> , 2020, 9, 1334.	3.5	4
8	Genetic variation of HvXYN1 associated with endoxylanase activity and TAX content in barley ( <i>Hordeum vulgare</i> L.). <i>BMC Plant Biology</i> , 2019, 19, 170.	3.6	5
9	Molecular Evolution and Interaction of Membrane Transport and Photoreception in Plants. <i>Frontiers in Genetics</i> , 2019, 10, 956.	2.3	21
10	Root and leaf metabolite profiles analysis reveals the adaptive strategies to low potassium stress in barley. <i>BMC Plant Biology</i> , 2018, 18, 187.	3.6	47
11	Association mapping for total polyphenol content, total flavonoid content and antioxidant activity in barley. <i>BMC Genomics</i> , 2018, 19, 81.	2.8	45
12	Development of predictive models for total phenolics and free p-coumaric acid contents in barley grain by near-infrared spectroscopy. <i>Food Chemistry</i> , 2017, 227, 342-348.	8.2	31
13	Isobaric Tags for Relative and Absolute Quantitation Proteomic Analysis of Germinating Barley under Gibberellin and Abscisic Acid Treatments. <i>Journal of Agricultural and Food Chemistry</i> , 2017, 65, 2248-2257.	5.2	6
14	Evolutionary Conservation of ABA Signaling for Stomatal Closure. <i>Plant Physiology</i> , 2017, 174, 732-747.	4.8	158
15	Speedy Grass Stomata: Emerging Molecular and Evolutionary Features. <i>Molecular Plant</i> , 2017, 10, 912-914.	8.3	36
16	The combined treatment of Mn and Al alleviates the toxicity of Al or Mn stress alone in barley. <i>Acta Physiologiae Plantarum</i> , 2016, 38, 1.	2.1	11
17	Identification of Quantitative Trait Loci for the Phenolic Acid Contents and Their Association with Agronomic Traits in Tibetan Wild Barley. <i>Journal of Agricultural and Food Chemistry</i> , 2016, 64, 980-987.	5.2	10
18	The effects of GA and ABA treatments on metabolite profile of germinating barley. <i>Food Chemistry</i> , 2016, 192, 928-933.	8.2	29

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19	Identification of the proteins associated with low potassium tolerance in cultivated and Tibetan wild barley. <i>Journal of Proteomics</i> , 2015, 126, 1-11.	2.4	44
20	Genetic Diversity and QTL Mapping of Thermostability of Limit Dextrinase in Barley. <i>Journal of Agricultural and Food Chemistry</i> , 2015, 63, 3778-3783.	5.2	19
21	Genetic Diversity of Individual Phenolic Acids in Barley and Their Correlation with Barley Malt Quality. <i>Journal of Agricultural and Food Chemistry</i> , 2015, 63, 7051-7057.	5.2	29
22	Comparative Transcriptome Profiling of Two Tibetan Wild Barley Genotypes in Responses to Low Potassium. <i>PLoS ONE</i> , 2014, 9, e100567.	2.5	76
23	Transcriptome profiling reveals mosaic genomic origins of modern cultivated barley. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 13403-13408.	7.1	74
24	The influence of salinity on cell ultrastructures and photosynthetic apparatus of barley genotypes differing in salt stress tolerance. <i>Acta Physiologiae Plantarum</i> , 2014, 36, 1261-1269.	2.1	30
25	The changes in physiological and biochemical traits of Tibetan wild and cultivated barley in response to low phosphorus stress. <i>Soil Science and Plant Nutrition</i> , 2014, 60, 832-842.	1.9	15
26	Grain protein content variation and its association analysis in barley. <i>BMC Plant Biology</i> , 2013, 13, 35.	3.6	95
27	Genome-Wide Association Analysis of Aluminum Tolerance in Cultivated and Tibetan Wild Barley. <i>PLoS ONE</i> , 2013, 8, e69776.	2.5	55
28	EFFECT OF SALINITY AND HEXAVALENT CHROMIUM STRESSES ON UPTAKE AND ACCUMULATION OF MINERAL ELEMENTS IN BARLEY GENOTYPES DIFFERING IN SALT TOLERANCE. <i>Journal of Plant Nutrition</i> , 2012, 35, 827-839.	1.9	24
29	Interactive effects of aluminum and chromium stresses on the uptake of nutrients and the metals in barley. <i>Soil Science and Plant Nutrition</i> , 2011, 57, 68-79.	1.9	17
30	Evolution of phosphate metabolism in Tibetan wild barley to adapt to aluminum stress. <i>Plant and Soil</i> , 0, , .	3.7	1