

Liangbo Fu

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/365000/publications.pdf>

Version: 2024-02-01

19
papers

513
citations

687363

13
h-index

794594

19
g-index

19
all docs

19
docs citations

19
times ranked

564
citing authors

#	ARTICLE	IF	CITATIONS
1	Multi-omics analysis reveals molecular mechanisms of shoot adaption to salt stress in Tibetan wild barley. <i>BMC Genomics</i> , 2016, 17, 889.	2.8	68
2	Ionic, metabolomic and proteomic analyses reveal molecular mechanisms of root adaption to salt stress in Tibetan wild barley. <i>Plant Physiology and Biochemistry</i> , 2018, 123, 319-330.	5.8	55
3	Calmodulin HvCaM1 Negatively Regulates Salt Tolerance via Modulation of HvHKT1s and HvCAMTA4. <i>Plant Physiology</i> , 2020, 183, 1650-1662.	4.8	50
4	Metabolite profiling and gene expression of Na/K transporter analyses reveal mechanisms of the difference in salt tolerance between barley and rice. <i>Plant Physiology and Biochemistry</i> , 2018, 130, 248-257.	5.8	44
5	Physiological and molecular mechanisms of cobalt and copper interaction in causing phyto-toxicity to two barley genotypes differing in Co tolerance. <i>Ecotoxicology and Environmental Safety</i> , 2020, 187, 109866.	6.0	42
6	Alleviating effects of calcium on cobalt toxicity in two barley genotypes differing in cobalt tolerance. <i>Ecotoxicology and Environmental Safety</i> , 2017, 139, 488-495.	6.0	37
7	Transcriptome-wide m6A methylation profile reveals regulatory networks in roots of barley under cadmium stress. <i>Journal of Hazardous Materials</i> , 2022, 423, 127140.	12.4	33
8	Time-course of ionic responses and proteomic analysis of a Tibetan wild barley at early stage under salt stress. <i>Plant Growth Regulation</i> , 2017, 81, 11-21.	3.4	26
9	Transcriptomic and alternative splicing analyses reveal mechanisms of the difference in salt tolerance between barley and rice. <i>Environmental and Experimental Botany</i> , 2019, 166, 103810.	4.2	24
10	High accumulation of phenolics and amino acids confers tolerance to the combined stress of cobalt and copper in barley (<i>Hordeum vulgare</i>). <i>Plant Physiology and Biochemistry</i> , 2020, 155, 927-937.	5.8	22
11	Copper alleviates cobalt toxicity in barley by antagonistic interaction of the two metals. <i>Ecotoxicology and Environmental Safety</i> , 2019, 180, 234-241.	6.0	21
12	Genotypic difference of cadmium tolerance and the associated microRNAs in wild and cultivated barley. <i>Plant Growth Regulation</i> , 2019, 87, 389-401.	3.4	15
13	The Influence of Nitrogen Application Level on Eating Quality of the Two Indica-Japonica Hybrid Rice Cultivars. <i>Plants</i> , 2020, 9, 1663.	3.5	15
14	Transcriptome analysis reveals the tolerant mechanisms to cobalt and copper in barley. <i>Ecotoxicology and Environmental Safety</i> , 2021, 209, 111761.	6.0	15
15	Vacuolar H ⁺ -pyrophosphatase HVP10 enhances salt tolerance via promoting Na ⁺ translocation into root vacuoles. <i>Plant Physiology</i> , 2022, 188, 1248-1263.	4.8	15
16	OsC2DP, a Novel C2 Domain-Containing Protein Is Required for Salt Tolerance in Rice. <i>Plant and Cell Physiology</i> , 2019, 60, 2220-2230.	3.1	11
17	Physiological mechanisms for antagonistic interaction of manganese and aluminum in barley. <i>Journal of Plant Nutrition</i> , 2019, 42, 466-476.	1.9	8
18	GWAS and transcriptomic integrating analysis reveals key salt-responding genes controlling Na ⁺ content in barley roots. <i>Plant Physiology and Biochemistry</i> , 2021, 167, 596-606.	5.8	8

#	ARTICLE	IF	CITATIONS
19	Identification of microRNAs Responding to Aluminium, Cadmium and Salt Stresses in Barley Roots. Plants, 2021, 10, 2754.	3.5	4