Ing-Feng Chang

List of Publications by Year in descending order

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471509 752698 22 1,696 17 20 citations h-index g-index papers 22 22 22 2512 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	CFM6 is an Essential CRM Protein Required for the Splicing of <i>nad5 </i> Transcript in Arabidopsis Mitochondria. Plant and Cell Physiology, 2022, 63, 217-233.	3.1	3
2	Arabidopsis glutamate receptor GLR3.7 is involved in abscisic acid response. Plant Signaling and Behavior, 2021, , 1997513.	2.4	6
3	Regulation of ABI5 expression by ABF3 during salt stress responses in Arabidopsis thaliana. , 2019, 60, 16.		47
4	The Glutamate Receptor-Like Protein GLR3.7 Interacts With 14-3-3ω and Participates in Salt Stress Response in Arabidopsis thaliana. Frontiers in Plant Science, 2019, 10, 1169.	3.6	57
5	Pharmacological Studies with Specific Agonist and Antagonist of Animal iGluR on Root Growth in Arabidopsis thaliana. , 2018 , , .		6
6	The Arabidopsis glutamate receptor-like gene <i>GLR3.6</i> controls root development by repressing the Kip-related protein gene <i>KRP4</i> . Journal of Experimental Botany, 2016, 67, 1853-1869.	4.8	59
7	Engineered xylose utilization enhances bio-products productivity in the cyanobacterium Synechocystis sp. PCC 6803. Metabolic Engineering, 2015, 30, 179-189.	7.0	53
8	A type III ACC synthase, ACS7, is involved in root gravitropism in Arabidopsis thaliana. Journal of Experimental Botany, 2013, 64, 4343-4360.	4.8	63
9	Comparative phosphoproteomic analysis of microsomal fractions of Arabidopsis thaliana and Oryza sativa subjected to high salinity. Plant Science, 2012, 185-186, 131-142.	3.6	33
10	Calcium-Dependent Protein Kinases from Arabidopsis Show Substrate Specificity Differences in an Analysis of 103 Substrates. Frontiers in Plant Science, 2011, 2, 36.	3.6	80
11	Proteomic profiling of proteins associated with the rejuvenation of Sequoia sempervirens (D. Don) Endl. Proteome Science, 2010, 8, 64.	1.7	19
12	Proteomic profiling of tandem affinity purified 14â€3â€3 protein complexes in <i>Arabidopsis thaliana</i> Proteomics, 2009, 9, 2967-2985.	2.2	193
13	Functional phosphoproteomic profiling of phosphorylation sites in membrane fractions of salt-stressed Arabidopsis thaliana. Proteome Science, 2009, 7, 42.	1.7	63
14	A novel yeast two-hybrid approach to identify CDPK substrates: Characterization of the interaction between AtCPK11 and AtDi19, a nuclear zinc finger protein1. FEBS Letters, 2006, 580, 904-911.	2.8	69
15	Mass spectrometry-based proteomic analysis of the epitope-tag affinity purified protein complexes in eukaryotes. Proteomics, 2006, 6, 6158-6166.	2.2	66
16	The Arabidopsis AtDi19 Gene Family Encodes a Novel Type of Cys2/His2 Zinc-finger Protein Implicated in ABA-independent Dehydration, High-salinity Stress and Light Signaling Pathways. Plant Molecular Biology, 2006, 61, 13-30.	3.9	85
17	Proteomic Characterization of Evolutionarily Conserved and Variable Proteins of Arabidopsis Cytosolic Ribosomes. Plant Physiology, 2005, 137, 848-862.	4.8	146
18	Induction of RhoGAP and Pathological Changes Characteristic of Alzheimers Disease by UAHFEMF Discharge in Rat Brain. Current Alzheimer Research, 2005, 2, 559-569.	1.4	13

#	Article	IF	CITATIONS
19	Immunopurification of Polyribosomal Complexes of Arabidopsis for Global Analysis of Gene Expression. Plant Physiology, 2005, 138, 624-635.	4.8	214
20	Humic Substances Affect the Activity of Chlorophyllase. Journal of Chemical Ecology, 2004, 30, 1057-1065.	1.8	34
21	Regulated Phosphorylation of 40S Ribosomal Protein S6 in Root Tips of Maize. Plant Physiology, 2003, 132, 2086-2097.	4.8	115
22	The Organization of Cytoplasmic Ribosomal Protein Genes in the Arabidopsis Genome. Plant Physiology, 2001, 127, 398-415.	4.8	272