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List of Publications by Year in descending order

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Version: 2024-02-01

		567281	552781
36	770	15	26
papers	citations	h-index	g-index
36	36	36	930
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	Molecular basis of a shattering resistance boosting global dissemination of soybean. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 17797-17802.	7.1	166
2	Somatic embryogenesis induced by the simple application of abscisic acid to carrot (Daucus carota L.) seedlings in culture. Planta, 2000, 211, 756-759.	3.2	94
3	Expression, Cloning, and Immunological Analysis of Buckwheat (<i>Fagopyrum esculentum</i>) Tj ETQq1 1 0.78	4314 rgBT 5.2	/Overlock 10
4	Fine mapping and development of DNA markers for the qPDH1 locus associated with pod dehiscence in soybean. Molecular Breeding, 2010, 25, 407-418.	2.1	37
5	A putative MYB35 ortholog is a candidate for the sex-determining genes in Asparagus officinalis. Scientific Reports, 2017, 7, 41497.	3.3	37
6	Temperature controls nuclear import of Tam3 transposase in <i>Antirrhinum</i> . Plant Journal, 2011, 65, 146-155.	5.7	35
7	A Single-Nucleotide Polymorphism in an Endo-1,4- \hat{l}^2 -Glucanase Gene Controls Seed Coat Permeability in Soybean. PLoS ONE, 2015, 10, e0128527.	2.5	35
8	A Major Soybean QTL, <i>qPDH1</i> , Controls Pod Dehiscence without Marked Morphological Change. Plant Production Science, 2009, 12, 217-223.	2.0	32
9	Assembly and disassembly of the peripheral architecture of the plant cell nucleus during mitosis. Planta, 1999, 210, 165-167.	3.2	27
10	Confirmation of the location and the effects of a major QTL controlling pod dehiscence, qPDH1, in soybean. Breeding Science, 2008, 58, 63-69.	1.9	23
11	A major QTL, qPDH1, is commonly involved in shattering resistance of soybean cultivars. Breeding Science, 2009, 59, 435-440.	1.9	23
12	Protein phosphatase 2A regulates the nuclear accumulation of the Arabidopsis bZIP protein VIP1 under hypo-osmotic stress. Journal of Experimental Botany, 2019, 70, 6101-6112.	4.8	21
13	A GDSL-type esterase/lipase gene, GELP77, is necessary for pollen dissociation and fertility in Arabidopsis. Biochemical and Biophysical Research Communications, 2020, 526, 1036-1041.	2.1	20
14	Mapping and use of QTLs controlling pod dehiscence in soybean. Breeding Science, 2012, 61, 554-558.	1.9	19
15	Localization of Daucus carota NMCP1 to the nuclear periphery: the role of the N-terminal region and an NLS-linked sequence motif, RYNLRR, in the tail domain. Frontiers in Plant Science, 2014, 5, 62.	3.6	17
16	Calcium signalling regulates the functions of the bZIP protein VIP1 in touch responses in <i>Arabidopsis thaliana</i> . Annals of Botany, 2018, 122, 1219-1229.	2.9	17
17	Molecular characterization of buckwheat major immunoglobulin E-reactive proteins in allergic patients. Allergology International, 2000, 49, 117-124.	3.3	15
18	Development of a DNA marker for variety discrimination specific to â€ ⁻ Manten-Kirariâ€ ^{-M} based on an NGS-RNA sequence in Tartary buckwheat (Fagopyrum tataricum). Food Chemistry, 2019, 295, 51-57.	8.2	11

#	Article	IF	Citations
19	The plant nuclear lamina proteins NMCP1 and NMCP2 form a filamentous network with lateral filament associations. Journal of Experimental Botany, 2021, 72, 6190-6204.	4.8	11
20	Detection of immunologically related Kunitz and Bowman-Birk proteinase inhibitors expressed during potato tuber development. Plant Molecular Biology, 1994, 26, 961-969.	3.9	10
21	Determination of the Absolute Configuration of a Monoglyceride Antibolting Compound and Isolation of Related Compounds from Radish Leaves (Raphanus sativus). Journal of Natural Products, 2017, 80, 872-878.	3.0	10
22	Multiple regulatory mechanisms influence the activity of the transposon, <i>Tam3</i> , of <i>Antirrhinum</i> . New Phytologist, 2008, 179, 343-355.	7.3	9
23	Identification of candidates for interacting partners of the tail domain of DcNMCP1, a major component of the Daucus carota nuclear lamina-like structure. Nucleus, 2017, 8, 312-322.	2.2	9
24	A putative AGAMOUS ortholog is a candidate for the gene determining ease of dehulling in Tartary buckwheat (Fagopyrum tataricum). Planta, 2020, 251, 85.	3.2	6
25	Detainment of Tam3 Transposase at Plasma Membrane by Its BED-Zinc Finger Domain. Plant Physiology, 2017, 173, 1492-1501.	4.8	5
26	CRISPR/Cas9-Mediated Editing of Genes Encoding rgs-CaM-like Proteins in Transgenic Potato Plants. Methods in Molecular Biology, 2019, 2028, 153-165.	0.9	5
27	Death of female flower microsporocytes progresses independently of meiosis-like process and can be accelerated by specific transcripts in Asparagus officinalis. Scientific Reports, 2019, 9, 2703.	3.3	5
28	Revision of the relationship between anther morphology and pollen sterility by cold stress at the booting stage in rice. Annals of Botany, 2021, 128, 559-575.	2.9	5
29	B-family subunits of protein phosphatase 2A are necessary for pollen development but not for female gametophyte development in Arabidopsis. Biochemical and Biophysical Research Communications, 2018, 505, 176-180.	2.1	4
30	Possible inhibition of Arabidopsis VIP1-mediated mechanosensory signaling by streptomycin. Plant Signaling and Behavior, 2018, 13, e1521236.	2.4	4
31	Functional characterization and vacuolar localization of fructan exohydrolase derived from onion (<i>Allium cepa</i>). Journal of Experimental Botany, 2022, 73, 4908-4922.	4.8	4
32	VIP1, a bZIP protein, interacts with the catalytic subunit of protein phosphatase 2A in Arabidopsis thaliana. Plant Signaling and Behavior, 2020, 15, 1706026.	2.4	3
33	Isolation of a major genetic interaction associated with an extreme phenotype using assorted F2 populations in rice. Molecular Breeding, 2014, 33, 997-1003.	2.1	2
34	The B″-family subunits of protein phosphatase 2A are necessary for in-vitro dephosphorylation of the Arabidopsis mechanosensory transcription factor VIP1. Biochemical and Biophysical Research Communications, 2021, 534, 353-358.	2.1	2
35	NDR/LATSâ€family protein kinase genes are indispensable for embryogenesis in Arabidopsis. FEBS Open Bio, 2021, 11, 2600-2606.	2.3	2
36	Data of whole genome sequencing of five garden asparagus (Asparagus officinalis) individuals with the MinION nanopore sequencer. Data in Brief, 2020, 28, 104838.	1.0	1