Hye Sun Cho

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Construction of SARS-CoV-2 virus-like particles in plant. Scientific Reports, 2022, 12, 1005.	3.3	26
2	Temporally distinct regulatory pathways coordinate thermo-responsive storage organ formation in potato. Cell Reports, 2022, 38, 110579.	6.4	10
3	The Arabidopsis cyclophilin CYP18-1 facilitates PRP18 dephosphorylation and the splicing of introns retained under heat stress. Plant Cell, 2022, 34, 2383-2403.	6.6	10
4	Complete genome sequence of artemisia virus B, a new polerovirus infecting Artemisia princeps in South Korea. Archives of Virology, 2021, 166, 1495-1499.	2.1	4
5	A More Accessible, Time-Saving, and Efficient Method for In Vitro Plant Regeneration from Potato Protoplasts. Plants, 2021, 10, 781.	3.5	10
6	Nitrogen Signaling Genes and SOC1 Determine the Flowering Time in a Reciprocal Negative Feedback Loop in Chinese Cabbage (Brassica rapa L.) Based on CRISPR/Cas9-Mediated Mutagenesis of Multiple BrSOC1 Homologs. International Journal of Molecular Sciences, 2021, 22, 4631.	4.1	10
7	Complete genome sequence of platycodon closterovirus 1, a novel putative member of the genus Closterovirus. Archives of Virology, 2021, 166, 2051-2054.	2.1	3
8	A single amino acid insertion in LCYB2 deflects carotenoid biosynthesis in red carrot. Plant Cell Reports, 2021, 40, 1793-1795.	5.6	5
9	Efficient plant regeneration from embryogenic cell suspension cultures of Euonymus alatus. Scientific Reports, 2021, 11, 15120.	3.3	6
10	FERONIA Confers Resistance to Photooxidative Stress in Arabidopsis. Frontiers in Plant Science, 2021, 12, 714938.	3.6	7
11	SUMO Modification of OsFKBP20-1b Is Integral to Proper Pre-mRNA Splicing upon Heat Stress in Rice. International Journal of Molecular Sciences, 2021, 22, 9049.	4.1	7
12	Complete genome sequence and genome organization of scorzonera virus A (SCoVA), a novel member of the genus Potyvirus. Archives of Virology, 2021, 166, 2901-2904.	2.1	0
13	<scp>PIN</scp> â€mediated polar auxin transport facilitates rootâ~'obstacle avoidance. New Phytologist, 2020, 225, 1285-1296.	7.3	39
14	Golgi-localized cyclophilin 21 proteins negatively regulate ABA signalling via the peptidyl prolyl isomerase activity during early seedling development. Plant Molecular Biology, 2020, 102, 19-38.	3.9	7
15	Development of Systems for the Production of Plant-Derived Biopharmaceuticals. Plants, 2020, 9, 30.	3.5	67
16	Submergence deactivates wound-induced plant defence against herbivores. Communications Biology, 2020, 3, 651.	4.4	5
17	Complete genome sequence and genome organization of achyranthes virus A, a novel member of the genus Potyvirus. Archives of Virology, 2020, 165, 2695-2698.	2.1	3
18	Gibberellin Promotes Bolting and Flowering via the Floral Integrators RsFT and RsSOC1-1 under Marginal Vernalization in Radish. Plants, 2020, 9, 594.	3.5	13

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19	Comparative proteomic analysis of host responses to Plasmodiophora brassicae infection in susceptible and resistant Brassica oleracea. Plant Biotechnology Reports, 2020, 14, 263-274.	1.5	11
20	OsFKBP20â€1b interacts with the splicing factor OsSR45 and participates in the environmental stress response at the postâ€transcriptional level in rice. Plant Journal, 2020, 102, 992-1007.	5.7	21
21	Genomic detection and molecular characterization of two distinct isolates of cycas necrotic stunt virus from Paeonia suffruticosa and Daphne odora. Virus Genes, 2019, 55, 734-737.	1.6	7
22	Complete genome sequence of a tentative new member of the genus Badnavirus identified in Codonopsis lanceolata. Archives of Virology, 2019, 164, 1733-1737.	2.1	3
23	Label-free quantitative proteomic analysis determines changes in amino acid and carbohydrate metabolism in three cultivars of Jerusalem artichoke tubers. Plant Biotechnology Reports, 2019, 13, 111-122.	1.5	8
24	Expression of Jerusalem artichoke (Helianthus tuberosus L.) fructosyltransferases, and high fructan accumulation in potato tubers. Applied Biological Chemistry, 2019, 62, .	1.9	8
25	Genome-wide identification of flowering time genes associated with vernalization and the regulatory flowering networks in Chinese cabbage. Plant Biotechnology Reports, 2018, 12, 347-363.	1.5	10
26	Transcriptome Profiling and Characterization of Drought-Tolerant Potato Plant (L.). Molecules and Cells, 2018, 41, 979-992.	2.6	24
27	Nicotiana benthamiana Matrix Metalloprotease 1 (NMMP1) gene confers disease resistance to Phytophthora infestans in tobacco and potato plants. Journal of Plant Physiology, 2017, 218, 189-195.	3.5	10
28	Overexpression of Golgi Protein CYP21-4s Improves Crop Productivity in Potato and Rice by Increasing the Abundance of Mannosidic Glycoproteins. Frontiers in Plant Science, 2017, 8, 1250.	3.6	15
29	Genome-wide Analysis of Alternative Splicing in An Inbred Cabbage (Brassica oleracea L.) Line â€~HO' in Response to Heat Stress. Current Genomics, 2017, 19, 12-20.	1.6	10
30	The OsCYP19-4 Gene Is Expressed as Multiple Alternatively Spliced Transcripts Encoding Isoforms with Distinct Cellular Localizations and PPIase Activities under Cold Stress. International Journal of Molecular Sciences, 2016, 17, 1154.	4.1	20
31	The Last Ten Years of Advancements in Plant-Derived Recombinant Vaccines against Hepatitis B. International Journal of Molecular Sciences, 2016, 17, 1715.	4.1	24
32	Identification of Flowering-Related Genes Responsible for Differences in Bolting Time between Two Radish Inbred Lines. Frontiers in Plant Science, 2016, 7, 1844.	3.6	26
33	Comparative transcriptome profiling and SSR marker identification in three Jerusalem artichoke (Helianthus tuberosus L.) cultivars exhibiting phenotypic variation. Plant Biotechnology Reports, 2016, 10, 447-461.	1.5	9
34	The complete sequence and genome organization of ligustrum virus A, a novel carlavirus. Archives of Virology, 2016, 161, 3593-3596.	2.1	3
35	Overexpression of <i>OsCYP19-4</i> increases tolerance to cold stress and enhances grain yield in rice (<i>Oryza sativa</i>). Journal of Experimental Botany, 2016, 67, 69-82.	4.8	51
36	Highly efficient plant regeneration and Agrobacterium-mediated transformation of Helianthus tuberosus L Industrial Crops and Products, 2016, 83, 670-679.	5.2	24

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37	OsCYP21-4, a novel Golgi-resident cyclophilin, increases oxidative stress tolerance in rice. Frontiers in Plant Science, 2015, 6, 797.	3.6	26
38	Rice cyclophilin OsCYP18â€2 is translocated to the nucleus by an interaction with SKIP and enhances drought tolerance in rice and <i>Arabidopsis</i> . Plant, Cell and Environment, 2015, 38, 2071-2087.	5.7	37
39	Physiological and molecular characterization of two inbred radish lines with different bolting times. Journal of Plant Biotechnology, 2015, 42, 215-222.	0.4	1
40	<scp>AtFKBP16</scp> â€1, a chloroplast lumenal immunophilin, mediates response to photosynthetic stress by regulating <scp>PsaL</scp> stability. Physiologia Plantarum, 2014, 150, 620-631.	5.2	25
41	RNA-Seq Analysis and De Novo Transcriptome Assembly of Jerusalem Artichoke (Helianthus tuberosus) Tj ETQq1	1 0.7843 2.5	l4 ggBT /Ove
42	Cucumber Pti1-L is a cytoplasmic protein kinase involved in defense responses and salt tolerance. Journal of Plant Physiology, 2014, 171, 817-822.	3.5	11
43	Use of Heat Stress Responsive Gene Expression Levels for Early Selection of Heat Tolerant Cabbage (Brassica oleracea L.). International Journal of Molecular Sciences, 2013, 14, 11871-11894.	4.1	39
44	A Rice Immunophilin Gene, OsFKBP16-3, Confers Tolerance to Environmental Stress in Arabidopsis and Rice. International Journal of Molecular Sciences, 2013, 14, 5899-5919.	4.1	15
45	Comparison of Major Nutrients in Eels Anguilla japonica Cultured with Different Formula Feeds or at Different Farms. Fisheries and Aquatic Sciences, 2013, 16, 85-92.	0.8	11
46	A novel gibberellin 2-oxidase gene CaGA2ox1 in pepper is specifically induced by incompatible plant pathogens. Plant Biotechnology Reports, 2012, 6, 381-390.	1.5	9
47	The rice thylakoid lumenal cyclophilin OsCYP20-2 confers enhanced environmental stress tolerance in tobacco and Arabidopsis. Plant Cell Reports, 2012, 31, 417-426.	5.6	45
48	Near-UV cyanobacteriochrome signaling system elicits negative phototaxis in the cyanobacterium <i>Synechocystis</i> sp. PCC 6803. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 10780-10785.	7.1	162
49	Evaluation of Major Nutrients of Domestic Farmed Eels Anguilla japonica. Han'guk Susan Hakhoe Chi = Bulletin of the Korean Fisheries Society, 2011, 44, 237-242.	0.1	3
50	A novel WD40 protein, BnSWD1, is involved in salt stress in Brassica napus. Plant Biotechnology Reports, 2010, 4, 165-172.	1.5	23
51	Classification of rice (Oryza satival. japonica nipponbare) immunophilins (FKBPs, CYPs) and expression patterns under water stress. BMC Plant Biology, 2010, 10, 253.	3.6	78
52	CaMsrB2, Pepper Methionine Sulfoxide Reductase B2, Is a Novel Defense Regulator against Oxidative Stress and Pathogen Attack. Plant Physiology, 2010, 154, 245-261.	4.8	86
53	Silencing of a BYPASS1 homolog results in root-independent pleiotrophic developmental defects in Nicotiana benthamiana. Plant Molecular Biology, 2008, 68, 423-437.	3.9	13
54	Pepper EST database: comprehensive in silico tool for analyzing the chili pepper (Capsicum annuum) transcriptome. BMC Plant Biology, 2008, 8, 101.	3.6	54

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55	A chloroplast cyclophilin functions in the assembly and maintenance of photosystem II in Arabidopsis thaliana. Proceedings of the National Academy of Sciences of the United States of America, 2007, 104, 15947-15952.	7.1	162
56	Tomato plants overexpressing CaKR1 enhanced tolerance to salt and oxidative stress. Biochemical and Biophysical Research Communications, 2007, 363, 983-988.	2.1	47
57	Induction of enhanced tolerance to cold stress and disease by overexpression of the pepper CaPIF1 gene in tomato. Physiologia Plantarum, 2007, 129, 555-566.	5.2	19
58	Capsicum annuum CCR4-associated factor CaCAF1 is necessary for plant development and defence response. Plant Journal, 2007, 51, 792-802.	5.7	65
59	Characterization of a Stress-Responsive Ankyrin Repeat-Containing Zinc Finger Protein of Capsicum annuum (CaKR1). BMB Reports, 2007, 40, 952-958.	2.4	41
60	Suppression of pepper SGT1 and SKP1 causes severe retardation of plant growth and compromises basal resistance. Physiologia Plantarum, 2006, 126, 060217072449001-???.	5.2	3
61	Inactivation of Organellar Glutamyl- and Seryl-tRNA Synthetases Leads to Developmental Arrest of Chloroplasts and Mitochondria in Higher Plants. Journal of Biological Chemistry, 2005, 280, 37098-37106.	3.4	35
62	DNA Gyrase Is Involved in Chloroplast Nucleoid Partitioning. Plant Cell, 2004, 16, 2665-2682.	6.6	80
63	CHRK1, a chitinase-related receptor-like kinase, plays a role in plant development and cytokinin homeostasis in tobacco. Plant Molecular Biology, 2003, 53, 877-890.	3.9	29
64	Interaction of NtCDPK1 calcium-dependent protein kinase with NtRpn3 regulatory subunit of the 26S proteasome inNicotiana tabacum. Plant Journal, 2003, 33, 825-840.	5.7	113
65	CHRK1, a chitinase-related receptor-like kinase, interacts with NtPUB4, an armadillo repeat protein, in tobacco. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2003, 1651, 50-59.	2.3	45
66	A novel dual-specificity protein kinase targeted to the chloroplast in tobacco1. FEBS Letters, 2001, 497, 124-130.	2.8	10
67	Expression patterns of diverse genes in response to gamma irradiation inNicotiana tabacum. Journal of Plant Biology, 2000, 43, 82-87.	2.1	23
68	CHRK1, a Chitinase-Related Receptor-Like Kinase in Tobacco. Plant Physiology, 2000, 123, 905-916.	4.8	68
69	Characterization of NtCDPK1, a calcium-dependent protein kinase gene in Nicotiana tabacum, and the activity of its encoded protein. Plant Molecular Biology, 1999, 39, 991-1001.	3.9	134
70	Multiple genes encoding serine/threonine protein phosphatases and their differential expression in Nicotiana tabacum. Plant Molecular Biology, 1998, 36, 315-322.	3.9	20
71	Temporally Distinct Regulatory Pathways Coordinate Thermo-Responsive Storage Organ Formation in Potato. SSRN Electronic Journal, 0, , .	0.4	0