

Erik Alexandersson

List of Publications by Year in descending order

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Version: 2024-02-01

37
papers

2,527
citations

279798

23
h-index

345221

36
g-index

41
all docs

41
docs citations

41
times ranked

3912
citing authors

#	ARTICLE	IF	CITATIONS
1	Whole Gene Family Expression and Drought Stress Regulation of Aquaporins. <i>Plant Molecular Biology</i> , 2005, 59, 469-484.	3.9	429
2	Arabidopsis Plasma Membrane Proteomics Identifies Components of Transport, Signal Transduction and Membrane Trafficking. <i>Plant and Cell Physiology</i> , 2004, 45, 1543-1556.	3.1	236
3	Normalyzer: A Tool for Rapid Evaluation of Normalization Methods for Omics Data Sets. <i>Journal of Proteome Research</i> , 2014, 13, 3114-3120.	3.7	218
4	The short-term growth response to salt of the developing barley leaf. <i>Journal of Experimental Botany</i> , 2006, 57, 1079-1095.	4.8	150
5	Transcriptional regulation of aquaporins in accessions of Arabidopsis in response to drought stress. <i>Plant Journal</i> , 2010, 61, 650-660.	5.7	150
6	High-Throughput Field-Phenotyping Tools for Plant Breeding and Precision Agriculture. <i>Agronomy</i> , 2019, 9, 258.	3.0	144
7	The genes and enzymes of the carotenoid metabolic pathway in <i>Vitis vinifera</i> L.. <i>BMC Genomics</i> , 2012, 13, 243.	2.8	112
8	Grapevine Plasticity in Response to an Altered Microclimate: Sauvignon Blanc Modulates Specific Metabolites in Response to Increased Berry Exposure. <i>Plant Physiology</i> , 2016, 170, 1235-1254.	4.8	91
9	Phosphite-induced changes of the transcriptome and secretome in <i>Solanum tuberosum</i> leading to resistance against <i>Phytophthora infestans</i> . <i>BMC Plant Biology</i> , 2014, 14, 254.	3.6	77
10	Quantitative proteomics and transcriptomics of potato in response to <i>Phytophthora infestans</i> in compatible and incompatible interactions. <i>BMC Genomics</i> , 2014, 15, 497.	2.8	77
11	The effects of the loss of TIP1;1 and TIP1;2 aquaporins in <i>Arabidopsis thaliana</i> . <i>Plant Journal</i> , 2008, 56, 756-767.	5.7	71
12	Potassium phosphite combined with reduced doses of fungicides provides efficient protection against potato late blight in large-scale field trials. <i>Crop Protection</i> , 2016, 86, 42-55.	2.1	70
13	Integrative Genomic Signatures Of Hepatocellular Carcinoma Derived from Nonalcoholic Fatty Liver Disease. <i>PLoS ONE</i> , 2015, 10, e0124544.	2.5	70
14	Plant secretome proteomics. <i>Frontiers in Plant Science</i> , 2013, 4, 9.	3.6	67
15	Proteomics and transcriptomics of the BABA-induced resistance response in potato using a novel functional annotation approach. <i>BMC Genomics</i> , 2014, 15, 315.	2.8	67
16	Plant Resistance Inducers against Pathogens in Solanaceae Species—From Molecular Mechanisms to Field Application. <i>International Journal of Molecular Sciences</i> , 2016, 17, 1673.	4.1	61
17	Field-omics—Understanding large-scale molecular data from field crops. <i>Frontiers in Plant Science</i> , 2014, 5, 286.	3.6	53
18	Automatic late blight lesion recognition and severity quantification based on field imagery of diverse potato genotypes by deep learning. <i>Knowledge-Based Systems</i> , 2021, 214, 106723.	7.1	46

#	ARTICLE	IF	CITATIONS
19	HvPIP1;6, a Barley (<i>Hordeum vulgare</i> L.) Plasma Membrane Water Channel Particularly Expressed in Growing Compared with Non-Growing Leaf Tissues. <i>Plant and Cell Physiology</i> , 2007, 48, 1132-1147.	3.1	44
20	Targeted Proteomics Approach for Precision Plant Breeding. <i>Journal of Proteome Research</i> , 2016, 15, 638-646.	3.7	44
21	Evaluation and integration of functional annotation pipelines for newly sequenced organisms: the potato genome as a test case. <i>BMC Plant Biology</i> , 2014, 14, 329.	3.6	42
22	Intact salicylic acid signalling is required for potato defence against the necrotrophic fungus <i>Alternaria solani</i> . <i>Plant Molecular Biology</i> , 2020, 104, 1-19.	3.9	32
23	Constitutive expression of a grapevine polygalacturonase-inhibiting protein affects gene expression and cell wall properties in uninfected tobacco. <i>BMC Research Notes</i> , 2011, 4, 493.	1.4	30
24	Botanicals and plant strengtheners for potato and tomato cultivation in Africa. <i>Journal of Integrative Agriculture</i> , 2020, 19, 406-427.	3.5	26
25	A novel workflow correlating RNA-seq data to <i>Phytophthora infestans</i> resistance levels in wild <i>Solanum</i> species and potato clones. <i>Frontiers in Plant Science</i> , 2015, 6, 718.	3.6	21
26	Linking crop traits to transcriptome differences in a progeny population of tetraploid potato. <i>BMC Plant Biology</i> , 2020, 20, 120.	3.6	18
27	Inoculation of Transgenic Resistant Potato by <i>Phytophthora infestans</i> Affects Host Plant Choice of a Generalist Moth. <i>PLoS ONE</i> , 2015, 10, e0129815.	2.5	16
28	Phosphite protects against potato and tomato late blight in tropical climates and has varying toxicity depending on the <i>Phytophthora infestans</i> isolate. <i>Crop Protection</i> , 2019, 121, 139-146.	2.1	14
29	Nongenetic Inheritance of Induced Resistance in a Wild Annual Plant. <i>Phytopathology</i> , 2016, 106, 877-883.	2.2	12
30	Functional phenomics for improved climate resilience in Nordic agriculture. <i>Journal of Experimental Botany</i> , 2022, 73, 5111-5127.	4.8	10
31	RNA seq analysis of potato cyst nematode interactions with resistant and susceptible potato roots. <i>European Journal of Plant Pathology</i> , 2018, 152, 531-539.	1.7	9
32	Gene Expression and Metabolite Profiling of Thirteen Nigerian Cassava Landraces to Elucidate Starch and Carotenoid Composition. <i>Agronomy</i> , 2020, 10, 424.	3.0	7
33	Phosphite alters the behavioral response of potato tuber moth (<i>Phthorimaea operculella</i>) to field-grown potato. <i>Pest Management Science</i> , 2019, 75, 616-621.	3.4	5
34	The MORPH web server and software tool for predicting missing genes in biological pathways. <i>Physiologia Plantarum</i> , 2015, 155, 12-20.	5.2	3
35	Phosphite Integrated in Late Blight Treatment Strategies in Starch Potato Does Not Cause Residues in the Starch Product. <i>Plant Disease</i> , 2020, 104, 3026-3032.	1.4	3
36	Leaf Apoplast of Field-Grown Potato Analyzed by Quantitative Proteomics and Activity-Based Protein Profiling. <i>International Journal of Molecular Sciences</i> , 2021, 22, 12033.	4.1	1

#	ARTICLE	IF	CITATIONS
37	Computer Vision and Less Complex Image Analyses to Monitor Potato Traits in Fields. Methods in Molecular Biology, 2021, 2354, 273-299.	0.9	0