## Nirala Ramchiary

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6365948/publications.pdf

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38 879 17 28 papers citations h-index g-index

times ranked

citing authors

docs citations

all docs

#	Article	IF	CITATIONS
1	A review on bioactive phytochemicals and ethnopharmacological potential of purslane (Portulaca) Tj ETQq1 1 0.78	84314 rgB	T <sub>3</sub> /Overlock
2	A comprehensive update on Capsicum proteomics: Advances and future prospects. Journal of Proteomics, 2022, 261, 104578.	2.4	6
3	Genomic Designing for Breeding Biotic Stress Resistant Pepper Crop., 2022,, 65-145.		1
4	A high-throughput RNA-Seq approach to elucidate the transcriptional response of Piriformospora indica to high salt stress. Scientific Reports, 2021, 11, 4129.	3.3	13
5	Role of Traditional Ethnobotanical Knowledge and Indigenous Communities in Achieving Sustainable Development Goals. Sustainability, 2021, 13, 3062.	3.2	24
6	Profiling of miRNAs in Bhut Jolokia (Capsicum chinense) and Kon Jolokia (C. frutescens) of Northeast India. Scientia Horticulturae, 2021, 281, 109952.	3.6	4
7	Integrating Omics and Gene Editing Tools for Rapid Improvement of Traditional Food Plants for Diversified and Sustainable Food Security. International Journal of Molecular Sciences, 2021, 22, 8093.	4.1	33
8	ICP-MS based analysis of mineral elements composition during fruit development in Capsicum germplasm. Journal of Food Composition and Analysis, 2021, 101, 103977.	3.9	5
9	MiR1885 Regulates Disease Tolerance Genes in Brassica rapa during Early Infection with Plasmodiophora brassicae. International Journal of Molecular Sciences, 2021, 22, 9433.	4.1	9
10	Improvement of a Traditional Orphan Food Crop, Portulaca oleracea L. (Purslane) Using Genomics for Sustainable Food Security and Climate-Resilient Agriculture. Frontiers in Sustainable Food Systems, 2021, 5, .	3.9	8
11	Identification and expression analysis of phosphate transporter genes and metabolites in response to phosphate stress in Capsicum annuum. Environmental and Experimental Botany, 2021, 190, 104597.	4.2	9
12	Capsicum chinense MYB Transcription Factor Genes: Identification, Expression Analysis, and Their Conservation and Diversification With Other Solanaceae Genomes. Frontiers in Plant Science, 2021, 12, 721265.	3.6	11
13	Genetics, Genomics and Breeding of Chili Pepper Capsicum frutescens L. and Other Capsicum Species. , 2021, , 59-86.		3
14	Analysis of bioactive components in Ghost chili ( <i>Capsicum chinense</i> ) for antioxidant, genotoxic, and apoptotic effects in mice. Drug and Chemical Toxicology, 2020, 43, 182-191.	2.3	13
15	Development and characterization of non-coding RNA based simple sequence repeat markers in Capsicum species. Genomics, 2020, 112, 1554-1564.	2.9	32
16	Characterizing the Nutrient Composition, Physiological Maturity, and Effect of Cold Storage in Khasi Mandarin ( <i>Citrus reticulata </i> Blanco). International Journal of Fruit Science, 2020, 20, 521-540.	2.4	10
17	Single-base cytosine methylation analysis in fruits of three Capsicum species. Genomics, 2020, 112, 3342-3353.	2.9	9
18	Transcriptome profiling, simple sequence repeat markers development and genetic diversity analysis of potential industrial crops Capsicum chinense and C. frutescens of Northeast India. Industrial Crops and Products, 2020, 154, 112687.	5.2	13

#	Article	IF	Citations
19	Comparative analysis of developmental changes of fruit metabolites, antioxidant activities and mineral elements content in Bhut jolokia and other Capsicum species. LWT - Food Science and Technology, 2019, 105, 363-370.	5.2	23
20	Identification of genes involved in fruit development/ripening in Capsicum and development of functional markers. Genomics, 2019, 111, 1913-1922.	2.9	31
21	Progress and Prospects in Capsicum Breeding for Biotic and Abiotic Stresses. , 2018, , 279-322.		15
22	Integrated analysis of leaf morphological and color traits in different populations of Chinese cabbage (Brassica rapa ssp. pekinensis). Theoretical and Applied Genetics, 2017, 130, 1617-1634.	3.6	9
23	Comparative Analysis of Fruit Metabolites and Pungency Candidate Genes Expression between Bhut Jolokia and Other Capsicum Species. PLoS ONE, 2016, 11, e0167791.	2.5	50
24	Molecular modeling and simulation studies of recombinant laccase from Yersinia enterocolitica suggests significant role in the biotransformation of non-steroidal anti-inflammatory drugs. Biochemical and Biophysical Research Communications, 2016, 469, 306-312.	2.1	29
25	Anatomic Characteristics Associated with Head Splitting in Cabbage (Brassica oleracea var. capitata) Tj ETQq $1\ 1$	0.784314 2.5	rgBT /Overlo
26	Quantitative trait loci mapping of partial resistance to Diamondback moth in cabbage (Brassica) Tj ETQq0 0 0 rg	BT <u>JO</u> verlo	ck <sub>10</sub> 0 Tf 50 4
27	Mapping QTLs of resistance to head splitting in cabbage (Brassica oleracea L.var. capitata L.). Molecular Breeding, 2015, 35, 1.	2.1	11
28	Biotechnological advances on in vitro capsaicinoids biosynthesis in capsicum: a review. Phytochemistry Reviews, 2015, 14, 189-201.	6.5	20
29	Application of genetics and genomics towards Capsicum translational research. Plant Biotechnology Reports, 2014, 8, 101-123.	1.5	49
30	Quantitative Trait Loci Mapping in Brassica rapa Revealed the Structural and Functional Conservation of Genetic Loci Governing Morphological and Yield Component Traits in the A, B, and C Subgenomes of Brassica Species. DNA Research, 2013, 20, 1-16.	3.4	59
31	Development of EST database and transcriptome analysis in the leaves of Brassica rapa using a newly developed pipeline. Genes and Genomics, 2012, 34, 671-679.	1.4	4
32	Identification of Potential microRNAs and Their Targets in Brassica rapa L Molecules and Cells, 2011, 32, 21-38.	2.6	43
33	Mapping quantitative trait loci for leaf and heading-related traits in Chinese cabbage (Brassica rapa L.) Tj ETQq1	1 0,78431 2.1	4 rgBT /Over
34	Genic Microsatellite Markers in Brassica rapa: Development, Characterization, Mapping, and Their Utility in Other Cultivated and Wild Brassica Relatives. DNA Research, 2011, 18, 305-320.	3.4	77
35	Development and linkage mapping of unigene-derived microsatellite markers in Brassica rapa L Breeding Science, 2011, 61, 160-167.	1.9	28
36	Development and characterization of new microsatellite markers in Panax ginseng (C.A. Meyer) from BAC end sequences. Conservation Genetics, 2010, 11, 1223-1225.	1.5	26

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37	Genetics of Clubroot Resistance in Brassica Species. Journal of Plant Growth Regulation, 2009, 28, 252-264.	5.1	120
38	Reaping the Benefits of Next-generation Sequencing Technologies for Crop Improvement — Solanaceae. , 0, , .		1