## Parviz Heidari

## List of Publications by Year in descending order

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430754 454834 1,122 48 18 30 citations h-index g-index papers 55 55 55 946 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	An Overview of Hazardous Impacts of Soil Salinity in Crops, Tolerance Mechanisms, and Amelioration through Selenium Supplementation. International Journal of Molecular Sciences, 2020, 21, 148.	1.8	289
2	The AP2/ERF Gene Family in Triticum durum: Genome-Wide Identification and Expression Analysis under Drought and Salinity Stresses. Genes, 2020, 11, 1464.	1.0	79
3	Genome-Wide Analysis of Potassium Channel Genes in Rice: Expression of the OsAKT and OsKAT Genes under Salt Stress. Genes, 2021, 12, 784.	1.0	49
4	Insights into the SAM Synthetase Gene Family and Its Roles in Tomato Seedlings under Abiotic Stresses and Hormone Treatments. Plants, 2020, 9, 586.	1.6	42
5	The GASA Gene Family in Cacao (Theobroma cacao, Malvaceae): Genome Wide Identification and Expression Analysis. Agronomy, 2021, 11, 1425.	1.3	40
6	Magnesium transporter Gene Family: Genome-Wide Identification and Characterization in Theobroma cacao, Corchorus capsularis, and Gossypium hirsutum of Family Malvaceae. Agronomy, 2021, 11, 1651.	1.3	40
7	In silico study of the CESA and CSL gene family in Arabidopsis thaliana and Oryza sativa: Focus on post-translation modifications. Plant Gene, 2019, 19, 100189.	1.4	35
8	Exogenous EBR Ameliorates Endogenous Hormone Contents in Tomato Species under Low-Temperature Stress. Horticulturae, 2021, 7, 84.	1.2	32
9	New Insights Into Structure and Function of TIFY Genes in Zea mays and Solanum lycopersicum: A Genome-Wide Comprehensive Analysis. Frontiers in Genetics, 2021, 12, 657970.	1.1	31
10	Investigation and Computational Analysis of the Sulfotransferase (SOT) Gene Family in Potato (Solanum tuberosum): Insights into Sulfur Adjustment for Proper Development and Stimuli Responses. Plants, 2021, 10, 2597.	1.6	31
11	Genome-wide characterization, expression profiling, and post-transcriptional study of GASA gene family. Gene Reports, 2020, 20, 100795.	0.4	29
12	Comparative plastome analysis of <i>Blumea</i> , with implications for genome evolution and phylogeny of Asteroideae. Ecology and Evolution, 2021, 11, 7810-7826.	0.8	29
13	Sorption Mechanism and Optimization Study for the Bioremediation of Pb(II) and Cd(II) Contamination by Two Novel Isolated Strains Q3 and Q5 of Bacillus sp International Journal of Environmental Research and Public Health, 2020, 17, 4059.	1.2	28
14	Cadmium and lead removal by new bacterial isolates from coal and aluminum mines. International Journal of Environmental Science and Technology, 2019, 16, 8297-8304.	1.8	27
15	Optimization Study of Nickel and Copper Bioremediation by Microbacterium oxydans Strain CM3 and CM7. Soil and Sediment Contamination, 2020, 29, 438-451.	1.1	27
16	Insights into the genes involved in the ethylene biosynthesis pathway in Arabidopsis thaliana and Oryza sativa. Journal of Genetic Engineering and Biotechnology, 2020, 18, 62.	1.5	25
17	Genome-wide characterization and expression analysis of fatty acid desaturase gene family in Camelina sativa. Gene Reports, 2020, 21, 100894.	0.4	23
18	Phenotypic variability of Pyrus boissieriana Buhse: Implications for conservation and breeding. Scientia Horticulturae, 2019, 247, 1-8.	1.7	22

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19	Genome-wide comparative analysis of Mg transporter gene family between Triticum turgidum and Camelina sativa. BioMetals, 2021, 34, 639-660.	1.8	20
20	Hormone Profiles and Antioxidant Activity of Cultivated and Wild Tomato Seedlings under Low-Temperature Stress. Agronomy, 2021, 11, 1146.	1.3	20
21	Influence of Morphine and Dopamine Receptor Sensitization on Locomotor Activity in Mice. Pharmacology, 2006, 78, 185-192.	0.9	17
22	A genome-wide association study to identify candidate genes for erectile dysfunction. Briefings in Bioinformatics, 2020, 22, .	3.2	16
23	Morphological and pomological variability of a grape (Vitis vinifera L.) germplasm collection. Scientia Horticulturae, 2020, 266, 109285.	1.7	16
24	Haplotype- and SNP-Based GWAS for Growth and Wood Quality Traits in Eucalyptus cladocalyx Trees under Arid Conditions. Plants, 2021, 10, 148.	1.6	15
25	Effects of Clozapine and Sulpiride on Morphine State-Dependent Memory in the Step-Down Passive Avoidance Test. Pharmacology, 2007, 79, 149-153.	0.9	12
26	Identification of superior apricot (Prunus armeniaca L.) genotypes among seedling origin trees. Scientia Horticulturae, 2020, 262, 109062.	1.7	9
27	Comparative Analysis of C-repeat Binding Factors (CBFs) in Tomato and Arabidopsis. Brazilian Archives of Biology and Technology, 0, 62, .	0.5	9
28	Morphological and fruit characterizations of common medlar (Mespilus germanica L.) germplasm. Scientia Horticulturae, 2019, 252, 38-47.	1.7	8
29	The selection of superior pistachio (Pistacia vera L.) genotypes among seedling trees originated from open-pollination. Scientia Horticulturae, 2019, 251, 88-100.	1.7	8
30	Soft computing-based approach on prediction promising pistachio seedling base on leaf characteristics. Scientia Horticulturae, 2020, 274, 109647.	1.7	8
31	Pseudogenization of the chloroplast threonine (trnT-GGU) gene in the sunflower family (Asteraceae). Scientific Reports, 2021, 11, 21122.	1.6	8
32	Using soft computing and leaf dimensions to determine sex in immature Pistacia vera genotypes. Measurement: Journal of the International Measurement Confederation, 2021, 174, 108988.	2.5	7
33	Morphological variabilities of Crataegus monogyna and C. pentagyna in northeastern areas of Iran. Industrial Crops and Products, 2019, 139, 111531.	2.5	6
34	Optimization and Characterization of Lead Bioremediation by Strains of Microbacterium Oxydans. Soil and Sediment Contamination, 2020, 29, 901-913.	1.1	5
35	Identification and DNA Fingerprinting of Some Superior Persian Walnut Genotypes in Iran. Erwerbs-Obstbau, 2021, 63, 393-402.	0.5	5
36	Cell Signaling in Model Plants. International Journal of Molecular Sciences, 2020, 21, 6062.	1.8	4

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37	The BAHD Gene Family in Cacao (Theobroma cacao, Malvaceae): Genome-Wide Identification and Expression Analysis. Frontiers in Ecology and Evolution, 2021, 9, .	1.1	4
38	Does the long-term contamination of lead (PbII) affect the bioremediation mechanisms of <i>Microbacterium oxydans</i> strain CM3 and CM7?. Soil and Sediment Contamination, 2022, 31, 959-973.	1.1	4
39	Effect of Exogenous Brassinosteroid Application on Grain Yield, some Physiological Traits and Expression of Genes Related to This Hormone Signaling Pathway in Wheat under Drought Stress. Plant Genetic Researches, 2020, 6, 157-172.	0.4	3
40	Phenotypical and Pomological Characterization of Non-irrigated Almond (Prunus dulcis Mill.) Trees to Select Superior Genotypes. Erwerbs-Obstbau, 2022, 64, 333-343.	0.5	3
41	Comparative in silico analysis of phosphate transporter gene family, PHT, in Camelina sativa gemome. Gene Reports, 2021, 25, 101351.	0.4	2
42	Morphological and Phenological Characterizations of Some Male and Female Promising Pistachio Genotypes from an Open-pollinated Population. International Journal of Fruit Science, 2021, 21, 456-467.	1.2	2
43	A Neural Network-Based Spectral Approach for the Assignment of Individual Trees to Genetically Differentiated Subpopulations. Remote Sensing, 2022, 14, 2898.	1.8	2
44	Variability of male and female pistachio genotypes with morphological and dominant DNA markers. Nucleus (India), $0$ , $1$ .	0.9	1
45	Validation of some of Housekeeping Genes in Aeluropus littoralis under Salinity Stress. Journal of Crop Breeding, 2018, 10, 110-117.	0.4	1
46	TraitCorr as a workbench for correlating gene expression measurements with phenotypic data. Gene Reports, 2020, 19, 100649.	0.4	0
47	Cell Signaling in Model Plants 2.0. International Journal of Molecular Sciences, 2021, 22, 8007.	1.8	0
48	Morphological variability of indigenous cherry plum (Prunus divaricata Ledeb.) accessions. European Journal of Horticultural Science, 2022, 87, .	0.3	0