

Abbas Saidi

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/11107911/publications.pdf>

Version: 2024-02-01

27
papers

347
citations

1040056

9
h-index

888059

17
g-index

28
all docs

28
docs citations

28
times ranked

359
citing authors

#	ARTICLE	IF	CITATIONS
1	Characterization of genetic diversity in chickpea using SSR markers, Start Codon Targeted Polymorphism (SCoT) and Conserved DNA-Derived Polymorphism (CDDP). <i>Physiology and Molecular Biology of Plants</i> , 2015, 21, 365-373.	3.1	70
2	Molecular Mapping and Characterization of Genes Governing Time to Flowering, Seed Weight, and Plant Height in an Intraspecific Genetic Linkage Map of Chickpea (<i>Cicer arietinum</i>). <i>Biochemical Genetics</i> , 2013, 51, 387-397.	1.7	38
3	Phytohormones: plant switchers in developmental and growth stages in potato. <i>Journal of Genetic Engineering and Biotechnology</i> , 2021, 19, 89.	3.3	28
4	Genetic variation for resistance to septoria tritici blotch in Iranian tetraploid wheat landraces. <i>European Journal of Plant Pathology</i> , 2012, 132, 191-202.	1.7	23
5	Green synthesis of stable silver nanoparticles by the main reduction component of green tea () Tj ETQq1 1 0.784314 rgBT /Oyerglock 10	3.8	18
6	Study of genetic diversity in local rose varieties (<i>Rosa</i> spp.) using molecular markers. <i>Banat's Journal of Biotechnology</i> , 2017, VIII, 148-157.	0.4	16
7	Application of Next Generation Sequencing, GWAS, RNA seq, WGRS, for genetic improvement of potato (<i>Solanum tuberosum</i> L.) under drought stress. <i>Biocatalysis and Agricultural Biotechnology</i> , 2020, 29, 101801.	3.1	15
8	A linkage map of chickpea (<i>Cicer arietinum</i> L.) based on population from ILC3279Å—ILC588 crosses: Location of genes for time to flowering, seed size and plant height. <i>Genetika</i> , 2015, 47, 253-263.	0.4	14
9	Phylogeny, gene structure and GATA genes expression in different tissues of solanaceae species. <i>Biocatalysis and Agricultural Biotechnology</i> , 2021, 35, 102015.	3.1	12
10	Effects of <i>Foeniculum vulgare</i> ethanol extract on osteogenesis in human mecenchymal stem cells. <i>Avicenna Journal of Phytomedicine</i> , 2013, 3, 135-42.	0.2	12
11	Evaluation of genetic diversity of carnation cultivars using CDDP and DAMD markers and morphological traits. <i>Nucleus (India)</i> , 2018, 61, 129-135.	2.2	10
12	Genome-wide identification of StU-box gene family and assessment of their expression in developmental stages of <i>Solanum tuberosum</i> . <i>Journal of Genetic Engineering and Biotechnology</i> , 2022, 20, 25.	3.3	9
13	Applicability of SCoT and SSR Molecular Markers for Genetic Diversity Analysis in <i>Chrysanthemum morifolium</i> Genotypes. <i>Proceedings of the National Academy of Sciences India Section B - Biological Sciences</i> , 2019, 89, 1067-1077.	1.0	8
14	In-silico analysis of eukaryotic translation initiation factors (eIFs) in response to environmental stresses in rice (<i>Oryza sativa</i>). <i>Biologia (Poland)</i> , 2020, 75, 1731-1738.	1.5	8
15	Characterization of cis-elements in hormonal stress-responsive genes in <i>Oryza sativa</i> . <i>Asia-Pacific Journal of Molecular Biology and Biotechnology</i> , 0, , 95-102.	0.1	8
16	Comparison of Genetic Variation of Anthurium (<i>Anthurium andraeanum</i>) Cultivars Using SCoT, CDDP and RAPD Markers. <i>Plant Tissue Culture and Biotechnology</i> , 2019, 28, 171-182.	0.2	7
17	Genome wide identification of StKNOX gene family and characterization of their expression in <i>Solanum tuberosum</i> . <i>Biocatalysis and Agricultural Biotechnology</i> , 2021, 37, 102160.	3.1	7
18	Suppression mechanism of the calcium sensitivity in <i>Saccharomyces cerevisiae</i> ptp2 ^Δ msg5 ^Δ double disruptant involves a novel HOG-independent function of Åsk2, transcription factor Msn2 and the protein kinase A component Bcy1. <i>Journal of Bioscience and Bioengineering</i> , 2014, 117, 135-141.	2.2	6

#	ARTICLE	IF	CITATIONS
19	Molecular approach to determine taxonomic status of Septoria sp. causing leaf blotch of Castanea sativa in Hyrcanian forests. Journal of Forestry Research, 2017, 28, 661-670.	3.6	6
20	Approaches for developing molecular markers associated with virus resistances in potato (Solanum) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5	2.9	6
21	Identification of responsive genes and analysis of genes with bacterial-inducible cis-regulatory elements in the promoter regions in Oryza sativa L.. Acta Agriculturae Slovenica, 2020, 116, .	0.3	6
22	First report of Cucumber mosaic virus on Canna indica in Iran. Australasian Plant Disease Notes, 2012, 7, 119-121.	0.7	5
23	Senescence-associated proteins and nitrogen remobilization in grain filling under drought stress condition. Journal of Genetic Engineering and Biotechnology, 2022, 20, 101.	3.3	5
24	Transcriptome analysis of Phytophthora infestans and Colletotrichum coccodes in tomato to reveal resistance mechanisms. Asia-Pacific Journal of Molecular Biology and Biotechnology, 0, , 39-51.	0.1	4
25	Monitoring Response of a Few bZip Transcription Factors in Response to Osmotic Stress in Sunflower. Iranian Journal of Biotechnology, 2018, 16, 120-131.	0.3	2
26	Genetic diversity study in lentil (Lens culinaris Medik.) Germplasm: a comparison of CAAT Box Derived Polymorphism (CBDP) and simple sequence repeat (SSR) markers. , 2022, 77, 2793-2803.		2
27	In vitro evaluation of ferutinin on proliferation and osteogenesis differentiation in human unrestricted Somatic stem cells. Steroids, 2021, 172, 108862.	1.8	1