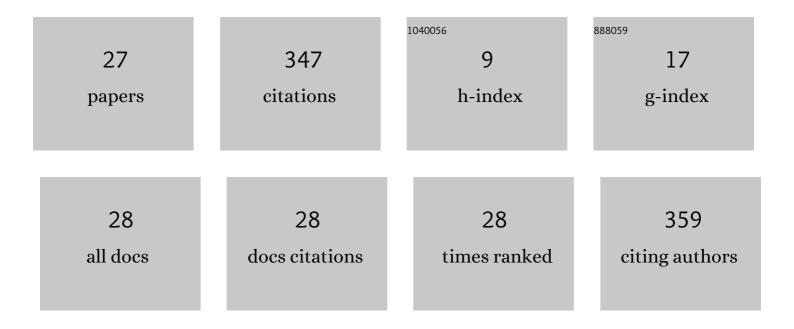
Abbas Saidi

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Genome-wide identification of StU-box gene family and assessment of their expression in developmental stages of Solanum tuberosum. Journal of Genetic Engineering and Biotechnology, 2022, 20, 25.	3.3	9
2	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
3	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
4	Approaches for developing molecular markers associated with virus resistances in potato (Solanum) Tj ETQq0 0	0 rgBT /Ov	erlock 10 Tf 5

5	Phytohormones: plant switchers in developmental and growth stages in potato. Journal of Genetic Engineering and Biotechnology, 2021, 19, 89.	3.3	28
6	In vitro evaluation of ferutinin on proliferation and osteogenesis differentiation in human unrestricted Somatic stem cells. Steroids, 2021, 172, 108862.	1.8	1
7	Phylogeny, gene structure and GATA genes expression in different tissues of solanaceae species. Biocatalysis and Agricultural Biotechnology, 2021, 35, 102015.	3.1	12
8	Genome wide identification of StKNOX gene family and characterization of their expression in Solanum tuberosum. Biocatalysis and Agricultural Biotechnology, 2021, 37, 102160.	3.1	7
9	Application of Next Generation Sequencing, GWAS, RNA seq, WGRS, for genetic improvement of potato (Solanum tuberosum L.) under drought stress. Biocatalysis and Agricultural Biotechnology, 2020, 29, 101801.	3.1	15
10	In-silico analysis of eukaryotic translation initiation factors (eIFs) in response to environmental stresses in rice (Oryza sativa). Biologia (Poland), 2020, 75, 1731-1738.	1.5	8
11	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

Green synthesis of stable silver nanoparticles by the main reduction component of green tea () Tj ETQq000 rgBT / $\frac{10}{3.8}$ rf 50 30.

13	Comparison of Genetic Variation of Anthurium (Anthurium andraeanum) Cultivars Using SCoT, CDDP and RAPD Markers. Plant Tissue Culture and Biotechnology, 2019, 28, 171-182.	0.2	7
14	Applicability of SCoT and SSR Molecular Markers for Genetic Diversity Analysis in Chrysanthemum morifolium Genotypes. Proceedings of the National Academy of Sciences India Section B - Biological Sciences, 2019, 89, 1067-1077.	1.0	8
15	Evaluation of genetic diversity of carnation cultivars using CDDP and DAMD markers and morphological traits. Nucleus (India), 2018, 61, 129-135.	2.2	10
16	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
17	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
18	Study of genetic diversity in local rose varieties (Rosa spp.) using molecular markers. Banat's Journal of Biotechnology, 2017, VIII, 148-157.	0.4	16

Abbas Saidi

#	Article	IF	CITATIONS
19	Characterization of genetic diversity in chickpea using SSR markers, Start Codon Targeted Polymorphism (SCoT) and Conserved DNA-Derived Polymorphism (CDDP). Physiology and Molecular Biology of Plants, 2015, 21, 365-373.	3.1	70
20	A linkage map of chickpea (Cicer arietinum L.) based on population from ILC3279×ILC588 crosses: Location of genes for time to flowering, seed size and plant height. Genetika, 2015, 47, 253-263.	0.4	14
21	Suppression mechanism of the calcium sensitivity in Saccharomyces cerevisiae ptp2î"msg5î" double disruptant involves a novel HOG-independent function ofÂSsk2, transcription factor Msn2 and the protein kinase A component Bcy1. Journal of Bioscience and Bioengineering, 2014, 117, 135-141.	2.2	6
22	Molecular Mapping and Characterization of Genes Governing Time to Flowering, Seed Weight, and Plant Height in an Intraspecific Genetic Linkage Map of Chickpea (Cicer arietinum). Biochemical Genetics, 2013, 51, 387-397.	1.7	38
23	Effects of Foeniculum vulgare ethanol extract on osteogenesis in human mecenchymal stem cells. Avicenna Journal of Phytomedicine, 2013, 3, 135-42.	0.2	12
24	First report of Cucumber mosaic virus on Canna indica in Iran. Australasian Plant Disease Notes, 2012, 7, 119-121.	0.7	5
25	Genetic variation for resistance to septoria tritici blotch in Iranian tetraploid wheat landraces. European Journal of Plant Pathology, 2012, 132, 191-202.	1.7	23
26	Characterization of cis-elements in hormonal stress-responsive genes in Oryza sativa. Asia-Pacific Journal of Molecular Biology and Biotechnology, 0, , 95-102.	0.1	8
27	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