Sajjad Asaf

List of Publications by Year in descending order

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100	4,493	35	62
papers	citations	h-index	g-index
105	105	105	3661 citing authors
all docs	docs citations	times ranked	

#	Article	IF	CITATIONS
1	Mangrove's rhizospheric engineering with bacterial inoculation improve degradation of diesel contamination. Journal of Hazardous Materials, 2022, 423, 127046.	12.4	8
2	Unraveling the Genome Sequence of Plant Growth Promoting Aspergillus niger (CSR3) Provides Insight into the Synthesis of Secondary Metabolites and Its Comparative Genomics. Journal of Fungi (Basel, Switzerland), 2022, 8, 107.	3.5	7
3	Uncovering the first complete plastome genomics, comparative analyses, and phylogenetic dispositions of endemic medicinal plant Ziziphus hajarensis (Rhamnaceae). BMC Genomics, 2022, 23, 83.	2.8	9
4	Endophytic fungus <i>Bipolaris</i> sp. CSL-1 induces salt tolerance in <i>Glycine max.</i> L via modulating its endogenous hormones, antioxidative system and gene expression. Journal of Plant Interactions, 2022, 17, 319-332.	2.1	16
5	Microbiome Variation Across Populations of Desert Halophyte Zygophyllum qatarensis. Frontiers in Plant Science, 2022, 13, 841217.	3.6	3
6	The Plastome Sequences of Triticum sphaerococcum (ABD) and Triticum turgidum subsp. durum (AB) Exhibit Evolutionary Changes, Structural Characterization, Comparative Analysis, Phylogenomics and Time Divergence. International Journal of Molecular Sciences, 2022, 23, 2783.	4.1	5
7	Drought and UV Radiation Stress Tolerance in Rice Is Improved by Overaccumulation of Non-Enzymatic Antioxidant Flavonoids. Antioxidants, 2022, 11, 917.	5.1	22
8	Silicon- and Boron-Induced Physio-Biochemical Alteration and Organic Acid Regulation Mitigates Aluminum Phytotoxicity in Date Palm Seedlings. Antioxidants, 2022, 11, 1063.	5.1	7
9	Complete mitochondrial genome of endangered Arabian tahr (<i>Arabitragus jayakari)</i> and phylogenetic placement. Mitochondrial DNA Part B: Resources, 2022, 7, 1189-1190.	0.4	O
10	Comparative Analysis of Date Palm (Phoenix dactylifera L.) Mitochondrial Genomics. Compendium of Plant Genomes, 2021, , 211-222.	0.5	3
11	Discovery and Validation of a Novel Step Catalyzed by OsF3H in the Flavonoid Biosynthesis Pathway. Biology, 2021, 10, 32.	2.8	5
12	Mangrove tree (Avicennia marina): insight into chloroplast genome evolutionary divergence and its comparison with related species from family Acanthaceae. Scientific Reports, 2021, 11, 3586.	3.3	11
13	From Traditional Breeding to Genome Editing for Boosting Productivity of the Ancient Grain Tef [Eragrostis tef (Zucc.) Trotter]. Plants, 2021, 10, 628.	3.5	16
14	Molecular epidemiology of COVID-19 in Oman: A molecular and surveillance study for the early transmission of COVID-19 in the country. International Journal of Infectious Diseases, 2021, 104, 139-149.	3.3	16
15	Halotolerant bacteria mitigate the effects of salinity stress on soybean growth by regulating secondary metabolites and molecular responses. BMC Plant Biology, 2021, 21, 176.	3.6	76
16	Decoding first complete chloroplast genome of toothbrush tree (Salvadora persica L.): insight into genome evolution, sequence divergence and phylogenetic relationship within Brassicales. BMC Genomics, 2021, 22, 312.	2.8	9
17	Plant Secondary Metabolite Biosynthesis and Transcriptional Regulation in Response to Biotic and Abiotic Stress Conditions. Agronomy, 2021, 11, 968.	3.0	256
18	Flavonone 3-hydroxylase Relieves Bacterial Leaf Blight Stress in Rice via Overaccumulation of Antioxidant Flavonoids and Induction of Defense Genes and Hormones. International Journal of Molecular Sciences, 2021, 22, 6152.	4.1	26

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19	Rhizospheric Bacillus spp. Rescues Plant Growth Under Salinity Stress via Regulating Gene Expression, Endogenous Hormones, and Antioxidant System of Oryza sativa L. Frontiers in Plant Science, 2021, 12, 665590.	3.6	38
20	The dynamic history of gymnosperm plastomes: Insights from structural characterization, comparative analysis, phylogenomics, and time divergence. Plant Genome, 2021, 14, e20130.	2.8	7
21	Over-Expression of Chorismate Mutase Enhances the Accumulation of Salicylic Acid, Lignin, and Antioxidants in Response to the White-Backed Planthopper in Rice Plants. Antioxidants, 2021, 10, 1680.	5.1	8
22	Transcriptomics of tapping and healing process in frankincense tree during resin production. Genomics, 2021, 113, 4337-4351.	2.9	2
23	Enhanced Flavonoid Accumulation Reduces Combined Salt and Heat Stress Through Regulation of Transcriptional and Hormonal Mechanisms. Frontiers in Plant Science, 2021, 12, 796956.	3.6	35
24	<i>Sphingomonas</i> : from diversity and genomics to functional role in environmental remediation and plant growth. Critical Reviews in Biotechnology, 2020, 40, 138-152.	9.0	264
25	Overexpression of OsCM alleviates BLB stress via phytohormonal accumulation and transcriptional modulation of defense-related genes in Oryza sativa. Scientific Reports, 2020, 10, 19520.	3.3	17
26	Complete Chloroplast Genome Characterization of Oxalis Corniculata and Its Comparison with Related Species from Family Oxalidaceae. Plants, 2020, 9, 928.	3.5	12
27	Transcriptomic analysis of Dubas bug (Ommatissus lybicus Bergevin) infestation to Date Palm. Scientific Reports, 2020, 10, 11505.	3.3	5
28	Modulation of sugar and nitrogen in callus induction media alter PAL pathway, SA and biomass accumulation in rice callus. Plant Cell, Tissue and Organ Culture, 2020, 143, 517-530.	2.3	5
29	Overexpression of OsF3H modulates WBPH stress by alteration of phenylpropanoid pathway at a transcriptomic and metabolomic level in Oryza sativa. Scientific Reports, 2020, 10, 14685.	3.3	35
30	Unraveling the Chloroplast Genomes of Two Prosopis Species to Identify Its Genomic Information, Comparative Analyses and Phylogenetic Relationship. International Journal of Molecular Sciences, 2020, 21, 3280.	4.1	21
31	Thermotolerance effect of plant growth-promoting Bacillus cereus SA1 on soybean during heat stress. BMC Microbiology, 2020, 20, 175.	3.3	147
32	Silicon-induced thermotolerance in Solanum lycopersicum L. via activation of antioxidant system, heat shock proteins, and endogenous phytohormones. BMC Plant Biology, 2020, 20, 248.	3.6	56
33	Expanded inverted repeat region with large scale inversion in the first complete plastid genome sequence of Plantago ovata. Scientific Reports, 2020, 10, 3881.	3.3	34
34	Complete Genome Sequence of Pseudomonas psychrotolerans CS51, a Plant Growth-Promoting Bacterium, Under Heavy Metal Stress Conditions. Microorganisms, 2020, 8, 382.	3.6	45
35	Rhizosphere Microbiome of Arid Land Medicinal Plants and Extra Cellular Enzymes Contribute to Their Abundance. Microorganisms, 2020, 8, 213.	3.6	37
36	Comparative Chloroplast Genomics of Endangered Euphorbia Species: Insights into Hotspot Divergence, Repetitive Sequence Variation, and Phylogeny. Plants, 2020, 9, 199.	3 . 5	12

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37	Extending thermotolerance to tomato seedlings by inoculation with SA1 isolate of Bacillus cereus and comparison with exogenous humic acid application. PLoS ONE, 2020, 15, e0232228.	2.5	59
38	Melatonin: Awakening the Defense Mechanisms during Plant Oxidative Stress. Plants, 2020, 9, 407.	3.5	124
39	Halo-tolerant rhizospheric Arthrobacter woluwensis AK1 mitigates salt stress and induces physio-hormonal changes and expression of GmST1 and GmLAX3 in soybean. Symbiosis, 2019, 77, 9-21.	2.3	47
40	Halotolerant Rhizobacterial Strains Mitigate the Adverse Effects of NaCl Stress in Soybean Seedlings. BioMed Research International, 2019, 2019, 1-15.	1.9	69
41	Integrated phytohormone production by the plant growth-promoting rhizobacterium <i>Bacillus tequilensis</i> SSB07 induced thermotolerance in soybean. Journal of Plant Interactions, 2019, 14, 416-423.	2.1	82
42	Metal Resistant Endophytic Bacteria Reduces Cadmium, Nickel Toxicity, and Enhances Expression of Metal Stress Related Genes with Improved Growth of Oryza Sativa, via Regulating Its Antioxidant Machinery and Endogenous Hormones. Plants, 2019, 8, 363.	3.5	111
43	Metabolic and proteomic alteration in phytohormone-producing endophytic Bacillus amyloliquefaciens RWL-1 during methanol utilization. Metabolomics, 2019, 15, 16.	3.0	28
44	Taxonomy, Distribution and Ecology of Boswellia. , 2019, , 11-34.		4
45	Frankincense: Tapping, Harvesting and Production. , 2019, , 35-51.		0
46	Frankincense Tree Physiology and Its Responses to Wounding Stress. , 2019, , 53-70.		2
47	Boswellia sacra Plastid Genome Sequencing and Comparative Analysis. , 2019, , 103-121.		0
48	Microbial Communities Accompanying Cultivated and Wild Boswellia sacra Trees., 2019, , 123-132.		6
49	Endophytic Microbial Communities of Boswellia. , 2019, , 133-151.		0
50	Biology of Genus Boswellia. , 2019, , .		10
51	Frankincense and Human Civilization: A Historical Review. , 2019, , 1-9.		1
52	Resin Composition and Structural Diversity. , 2019, , 153-162.		0
53	Genetic Diversity and Differentiation Among Species and Populations of Boswellia., 2019,, 85-101.		0
54	Indole-3-acetic-acid and ACC deaminase producing Leclercia adecarboxylata MO1 improves Solanum lycopersicum L. growth and salinity stress tolerance by endogenous secondary metabolites regulation. BMC Microbiology, 2019, 19, 80.	3.3	146

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55	WRKYs, the Jack-of-various-Trades, Modulate Dehydration Stress in Populus davidiana—A Transcriptomic Approach. International Journal of Molecular Sciences, 2019, 20, 414.	4.1	6
56	Alleviation of salt stress response in soybean plants with the endophytic bacterial isolate Curtobacterium sp. SAK1. Annals of Microbiology, 2019, 69, 797-808.	2.6	88
57	Growth-promoting bioactivities of Bipolaris sp. CSL-1 isolated from Cannabis sativa suggest a distinctive role in modifying host plant phenotypic plasticity and functions. Acta Physiologiae Plantarum, 2019, 41, 1.	2.1	14
58	Biotransformation of benzoin by Sphingomonas sp. LK11 and ameliorative effects on growth of Cucumis sativus. Archives of Microbiology, 2019, 201, 591-601.	2.2	14
59	Complete Chloroplast Genomes of Vachellia nilotica and Senegalia senegal: Comparative Genomics and Phylogenomic Placement in a New Generic System. PLoS ONE, 2019, 14, e0225469.	2.5	11
60	First complete chloroplast genomics and comparative phylogenetic analysis of Commiphora gileadensis and C. foliacea: Myrrh producing trees. PLoS ONE, 2019, 14, e0208511.	2.5	31
61	Rhizobacteria AK1 remediates the toxic effects of salinity stress via regulation of endogenous phytohormones and gene expression in soybean. Biochemical Journal, 2019, 476, 2393-2409.	3.7	36
62	Complete chloroplast genomes of medicinally important <i>Teucrium</i> species and comparative analyses with related species from Lamiaceae. PeerJ, 2019, 7, e7260.	2.0	12
63	Exogenous application of nitric oxide donors regulates short-term flooding stress in soybean. PeerJ, 2019, 7, e7741.	2.0	20
64	<i>Aspergillus niger</i> CSR3 regulates plant endogenous hormones and secondary metabolites by producing gibberellins and indoleacetic acid. Journal of Plant Interactions, 2018, 13, 100-111.	2.1	75
65	<i>Preussia</i> sp. BSL-10 producing nitric oxide, gibberellins, and indole acetic acid and improving rice plant growth. Journal of Plant Interactions, 2018, 13, 112-118.	2.1	26
66	Transcriptome wide identification and characterization of NO-responsive WRKY transcription factors in Arabidopsis thaliana L Environmental and Experimental Botany, 2018, 148, 128-143.	4.2	39
67	Plant growth promoting endophytic fungi Asprgillus fumigatus TS1 and Fusarium proliferatum BRL1 produce gibberellins and regulates plant endogenous hormones. Symbiosis, 2018, 76, 117-127.	2.3	165
68	Endophytic fungus Paecilomyces formosus LHL10 produces sester-terpenoid YW3548 and cyclic peptide that inhibit urease and $\hat{1}$ ±-glucosidase enzyme activities. Archives of Microbiology, 2018, 200, 1493-1502.	2.2	35
69	First reported chloroplast genome sequence of Punica granatum (cultivar Helow) from Jabal Al-Akhdar, Oman: phylogenetic comparative assortment with Lagerstroemia. Genetica, 2018, 146, 461-474.	1.1	7
70	Complete genome sequencing and analysis of endophytic Sphingomonas sp. LK11 and its potential in plant growth. 3 Biotech, 2018, 8, 389.	2.2	58
71	The first complete mitochondrial genome of wild soybean (<i>Glycine soja</i>). Mitochondrial DNA Part B: Resources, 2018, 3, 527-528.	0.4	4
72	First chloroplast genomics study of Phoenix dactylifera (var. Naghal and Khanezi): A comparative analysis. PLoS ONE, 2018, 13, e0200104.	2.5	15

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73	What Is There in Seeds? Vertically Transmitted Endophytic Resources for Sustainable Improvement in Plant Growth. Frontiers in Plant Science, 2018, 9, 24.	3.6	208
74	Biochar amendment changes jasmonic acid levels in two rice varieties and alters their resistance to herbivory. PLoS ONE, 2018, 13, e0191296.	2.5	32
75	First complete mitochondrial genome of Phoenix dact ylifera var. Khanezi. Mitochondrial DNA Part B: Resources, 2018, 3, 778-779.	0.4	3
76	Salt tolerance of Glycine max .L induced by endophytic fungus Aspergillus flavus CSH1, via regulating its endogenous hormones and antioxidative system. Plant Physiology and Biochemistry, 2018, 128, 13-23.	5.8	84
77	Gibberellin application ameliorates the adverse impact of short-term flooding on Glycine max L Biochemical Journal, 2018, 475, 2893-2905.	3.7	21
78	Complete chloroplast genome sequence and comparative analysis of loblolly pine (Pinus taeda L.) with related species. PLoS ONE, 2018, 13, e0192966.	2.5	44
79	Bacterial endophytes from arid land plants regulate endogenous hormone content and promote growth in crop plants: an example of <i>Sphingomonas</i> sp. and <i>Serratia marcescens</i> Journal of Plant Interactions, 2017, 12, 31-38.	2.1	90
80	Additive effects due to biochar and endophyte application enable soybean to enhance nutrient uptake and modulate nutritional parameters. Journal of Zhejiang University: Science B, 2017, 18, 109-124.	2.8	29
81	Endophytes from medicinal plants and their potential for producing indole acetic acid, improving seed germination and mitigating oxidative stress. Journal of Zhejiang University: Science B, 2017, 18, 125-137.	2.8	46
82	Complete mitochondrial genome sequence of <i>Aspergillus oryzae</i> RIB 127 and its comparative analysis with related species. Mitochondrial DNA Part B: Resources, 2017, 2, 632-633.	0.4	1
83	Gibberellins and indole-3-acetic acid producing rhizospheric bacterium <i>Leifsonia xyli</i> SE134 mitigates the adverse effects of copper-mediated stress on tomato. Journal of Plant Interactions, 2017, 12, 373-380.	2.1	48
84	Osmoprotective functions conferred to soybean plants via inoculation with Sphingomonas sp. LK11 and exogenous trehalose. Microbiological Research, 2017, 205, 135-145.	5.3	100
85	Chloroplast genomes of Arabidopsis halleri ssp. gemmifera and Arabidopsis lyrata ssp. petraea: Structures and comparative analysis. Scientific Reports, 2017, 7, 7556.	3.3	86
86	Plant growth-promoting endophyte Sphingomonas sp. LK11 alleviates salinity stress in Solanum pimpinellifolium. Environmental and Experimental Botany, 2017, 133, 58-69.	4.2	131
87	Plant growth-promoting endophytic bacteria versus pathogenic infections: an example of < i > Bacillus amylolique faciens < i > RWL-1 and < i > Fusarium oxysporum < i > f. sp. < i > lycopersici < i > in tomato. Peerl, 2017, 5, e3107.	2.0	86
88	The Complete Chloroplast Genome of Wild Rice (Oryza minuta) and Its Comparison to Related Species. Frontiers in Plant Science, 2017, 8, 304.	3.6	115
89	Endophytic Paecilomyces formosus LHL10 Augments Glycine max L. Adaptation to Ni-Contamination through Affecting Endogenous Phytohormones and Oxidative Stress. Frontiers in Plant Science, 2017, 8, 870.	3.6	59
90	The First Chloroplast Genome Sequence of Boswellia sacra, a Resin-Producing Plant in Oman. PLoS ONE, 2017, 12, e0169794.	2.5	19

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91	Complete genome analysis of Serratia marcescens RSC-14: A plant growth-promoting bacterium that alleviates cadmium stress in host plants. PLoS ONE, 2017, 12, e0171534.	2.5	52
92	Comparative analysis of complete plastid genomes from wild soybean (Glycine soja) and nine other Glycine species. PLoS ONE, 2017, 12, e0182281.	2.5	53
93	Rhizospheric microbial communities associated with wild and cultivated frankincense producing Boswellia sacra tree. PLoS ONE, 2017, 12, e0186939.	2.5	13
94	Endophytic Fungi from Frankincense Tree Improves Host Growth and Produces Extracellular Enzymes and Indole Acetic Acid. PLoS ONE, 2016, 11, e0158207.	2.5	124
95	Mitochondrial Genome Analysis of Wild Rice (Oryza minuta) and Its Comparison with Other Related Species. PLoS ONE, 2016, 11, e0152937.	2.5	31
96	Complete Chloroplast Genome of Nicotiana otophora and its Comparison with Related Species. Frontiers in Plant Science, 2016, 7, 843.	3.6	108
97	First draft genome sequencing of indole acetic acid producing and plant growth promoting fungus Preussia sp. BSL10. Journal of Biotechnology, 2016, 225, 44-45.	3.8	18
98	Seed-borne endophytic Bacillus amyloliquefaciens RWL-1 produces gibberellins and regulates endogenous phytohormones of Oryza sativa. Plant Physiology and Biochemistry, 2016, 106, 236-243.	5.8	219
99	Complete genome sequence of plant growth-promoting bacterium Leifsonia xyli SE134, a possible gibberellin and auxin producer. Journal of Biotechnology, 2016, 239, 34-38.	3.8	21
100	Isolation and characterization of a novel silicate-solubilizing bacterial strain <i>Burkholderia eburnea</i> CS4-2 that promotes growth of japonica rice (<i>Oryza sativa</i> L. cv. Dongjin). Soil Science and Plant Nutrition. 0 1-9.	1.9	28