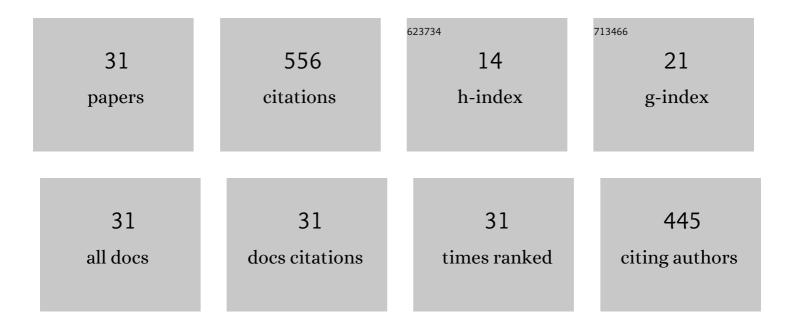
Sagheer Ahmad

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Genome-Wide Analysis of the NAC Transcription Factor Gene Family Reveals Differential Expression Patterns and Cold-Stress Responses in the Woody Plant Prunus mume. Genes, 2018, 9, 494.	2.4	47
2	The genome of <i>Cymbidium sinense</i> revealed the evolution of orchid traits. Plant Biotechnology Journal, 2021, 19, 2501-2516.	8.3	46
3	Comprehensive Cloning of Prunus mume Dormancy Associated MADS-Box Genes and Their Response in Flower Bud Development and Dormancy. Frontiers in Plant Science, 2018, 9, 17.	3.6	40
4	PmCBFs synthetically affect PmDAM6 by alternative promoter binding and protein complexes towards the dormancy of bud for Prunus mume. Scientific Reports, 2018, 8, 4527.	3.3	39
5	Crosstalk of PmCBFs and PmDAMs Based on the Changes of Phytohormones under Seasonal Cold Stress in the Stem of Prunus mume. International Journal of Molecular Sciences, 2018, 19, 15.	4.1	38
6	Highly Efficient Leaf Base Protoplast Isolation and Transient Expression Systems for Orchids and Other Important Monocot Crops. Frontiers in Plant Science, 2021, 12, 626015.	3.6	34
7	Overexpression of LiDXS and LiDXR From Lily (Lilium â€̃Siberia') Enhances the Terpenoid Content in Tobacco Flowers. Frontiers in Plant Science, 2018, 9, 909.	3.6	32
8	Comparative Transcriptome Reveals Benzenoid Biosynthesis Regulation as Inducer of Floral Scent in the Woody Plant Prunus mume. Frontiers in Plant Science, 2017, 8, 319.	3.6	29
9	SEP-class genes in Prunus mume and their likely role in floral organ development. BMC Plant Biology, 2017, 17, 10.	3.6	26
10	Flavonols and Carotenoids in Yellow Petals of Rose Cultivar (<i>Rosa</i> â€~Sun City'): A Possible Rich Source of Bioactive Compounds. Journal of Agricultural and Food Chemistry, 2018, 66, 4171-4181.	5.2	25
11	Dry Storage Effects on Postharvest Performance of Selected Cut Flowers. HortTechnology, 2012, 22, 463-469.	0.9	22
12	Comparative Metabolomic Analysis Reveals Distinct Flavonoid Biosynthesis Regulation for Leaf Color Development of Cymbidium sinense â€~Red Sun'. International Journal of Molecular Sciences, 2020, 21, 1869.	4.1	21
13	Identification and Characterization of NPR1 and PR1 Homologs in Cymbidium orchids in Response to Multiple Hormones, Salinity and Viral Stresses. International Journal of Molecular Sciences, 2020, 21, 1977.	4.1	20
14	Red to Far-Red Light Ratio Modulates Hormonal and Genetic Control of Axillary bud Outgrowth in Chrysanthemum (Dendranthema grandiflorum â€Jinba'). International Journal of Molecular Sciences, 2018, 19, 1590.	4.1	17
15	Transcriptional Cascade in the Regulation of Flowering in the Bamboo Orchid Arundina graminifolia. Biomolecules, 2021, 11, 771.	4.0	12
16	The Genetic and Hormonal Inducers of Continuous Flowering in Orchids: An Emerging View. Cells, 2022, 11, 657.	4.1	12
17	Morpho-physiological integrators, transcriptome and coexpression network analyses signify the novel molecular signatures associated with axillary bud in chrysanthemum. BMC Plant Biology, 2020, 20, 145.	3.6	11
18	Genetic insights into the regulatory pathways for continuous flowering in a unique orchid Arundina graminifolia. BMC Plant Biology, 2021, 21, 587.	3.6	11

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#	Article	IF	CITATIONS
19	Identification of the PmWEEP locus controlling weeping traits in Prunus mume through an integrated genome-wide association study and quantitative trait locus mapping. Horticulture Research, 2021, 8, 131.	6.3	10
20	Organ-Specific Gene Expression Reveals the Role of the Cymbidium ensifolium-miR396/Growth-Regulating Factors Module in Flower Development of the Orchid Plant Cymbidium ensifolium. Frontiers in Plant Science, 2021, 12, 799778.	3.6	9
21	Isolation, functional characterization and evolutionary study of LFY1 gene in Prunus mume. Plant Cell, Tissue and Organ Culture, 2019, 136, 523-536.	2.3	8
22	Genome-wide identification, characterisation, and evolution of <i>ABF/AREB</i> subfamily in nine Rosaceae species and expression analysis in mei (<i>Prunus mume</i>). PeerJ, 2021, 9, e10785.	2.0	8
23	Transcriptional Proposition for Uniquely Developed Protocorm Flowering in Three Orchid Species: Resources for Innovative Breeding. Frontiers in Plant Science, 0, 13, .	3.6	8
24	Comprehensive Analysis for GRF Transcription Factors in Sacred Lotus (Nelumbo nucifera). International Journal of Molecular Sciences, 2022, 23, 6673.	4.1	7
25	The de novo transcriptome identifies important zinc finger signatures associated with flowering in the orchid Arundina graminifolia. Scientia Horticulturae, 2022, 291, 110572.	3.6	6
26	Why Black Flowers? An Extreme Environment and Molecular Perspective of Black Color Accumulation in the Ornamental and Food Crops. Frontiers in Plant Science, 2022, 13, 885176.	3.6	4
27	The Transcriptome Profiling of Flavonoids and Bibenzyls Reveals Medicinal Importance of Rare Orchid Arundina graminifolia. Frontiers in Plant Science, 0, 13, .	3.6	4
28	Selection of optimal reference genes for qRT-PCR analysis of shoot development and graviresponse in prostrate and erect chrysanthemums. PLoS ONE, 2019, 14, e0225241.	2.5	3
29	PmSOC1s and PmDAMs participate in flower bud dormancy of Prunus mume by forming protein complexes and responding to ABA. European Journal of Horticultural Science, 2021, 86, 480-490.	0.7	3
30	Stage Specificity, the Dynamic Regulators and the Unique Orchid Arundina graminifolia. International Journal of Molecular Sciences, 2021, 22, 10935.	4.1	3
31	Decapitation Experiments Combined with the Transcriptome Analysis Reveal the Mechanism of High Temperature on Chrysanthemum Axillary Bud Formation. International Journal of Molecular Sciences, 2021, 22, 9704.	4.1	1