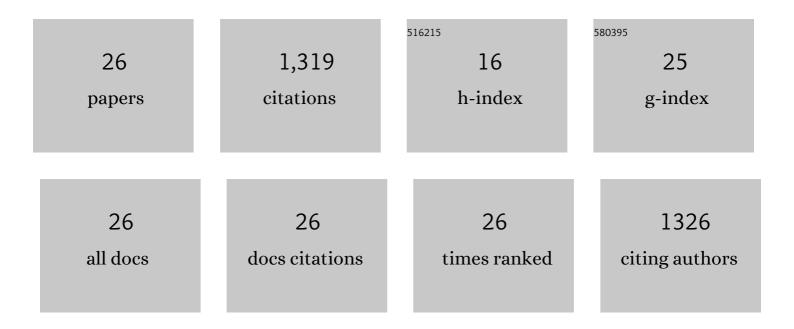
Khaled M A Amiri

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Patterns of Volatile Diversity Yield Insights Into the Genetics and Biochemistry of the Date Palm Fruit Volatilome. Frontiers in Plant Science, 2022, 13, 853651.	1.7	6
2	Transcriptome Profiling and Functional Validation of RINC-Type E3 Ligases in Halophyte Sesuvium verrucosum under Salinity Stress. International Journal of Molecular Sciences, 2022, 23, 2821.	1.8	3
3	Large-scale genome sequencing reveals the driving forces of viruses in microalgal evolution. Cell Host and Microbe, 2021, 29, 250-266.e8.	5.1	48
4	Complete Genome Sequences of Pseudomonas atacamensis Strain SM1 and Pseudomonas toyotomiensis Strain SM2, Isolated from the Date Palm Rhizosphere. Microbiology Resource Announcements, 2021, 10,	0.3	4
5	Genomic Landscape of the Mitochondrial Genome in the United Arab Emirates Native Population. Genes, 2020, 11, 876.	1.0	7
6	Complete Genome Sequence of the Plant Growth-Promoting Bacterium Pantoea agglomerans Strain UAEU18, Isolated from Date Palm Rhizosphere Soil in the United Arab Emirates. Microbiology Resource Announcements, 2020, 9, .	0.3	6
7	Transcriptome Profiling of Haloxylon persicum (Bunge ex Boiss and Buhse) an Endangered Plant Species under PEG-Induced Drought Stress. Genes, 2020, 11, 640.	1.0	8
8	Prospects for the Study and Improvement of Abiotic Stress Tolerance in Date Palms in the Post-genomics Era. Frontiers in Plant Science, 2020, 11, 293.	1.7	34
9	Complete Genome Sequence of Phytobacter diazotrophicus Strain UAEU22, a Plant Growth-Promoting Bacterium Isolated from the Date Palm Rhizosphere. Microbiology Resource Announcements, 2020, 9, .	0.3	4
10	The genome of pest Rhynchophorus ferrugineus reveals gene families important at the plant-beetle interface. Communications Biology, 2020, 3, 323.	2.0	44
11	Genome-wide association mapping of date palm fruit traits. Nature Communications, 2019, 10, 4680.	5.8	75
12	Schisandrin A protects intestinal epithelial cells from deoxynivalenol-induced cytotoxicity, oxidative damage and inflammation. Scientific Reports, 2019, 9, 19173.	1.6	35
13	Mapping of HKT1;5 Gene in Barley Using GWAS Approach and Its Implication in Salt Tolerance Mechanism. Frontiers in Plant Science, 2018, 9, 156.	1.7	95
14	Y-chromosome polymorphisms in the United Arab Emirates population. Forensic Science International: Genetics Supplement Series, 2017, 6, e397-e398.	0.1	0
15	Distribution of AGG interruption patterns within nine world populations. Intractable and Rare Diseases Research, 2014, 3, 153-161.	0.3	20
16	Screening for the Presence of FMR1 Premutation Alleles in Women With Parkinsonism. Archives of Neurology, 2009, 66, 244-9.	4.9	27
17	A Rapid Polymerase Chain Reaction-Based Screening Method for Identification of All Expanded Alleles of the Fragile X (FMR1) Gene in Newborn and High-Risk Populations. Journal of Molecular Diagnostics, 2008, 10, 43-49.	1.2	323
18	Fragile X–Associated Tremor/Ataxia Syndrome. Archives of Neurology, 2008, 65, 19-25.	4.9	67

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#	Article	IF	CITATIONS
19	Seroprevalence of antibodies to repetitive domains of Plasmodium vivax circumsporozoite protein in United Arab Emirates children. Transactions of the Royal Society of Tropical Medicine and Hygiene, 2002, 96, 560-564.	0.7	5
20	The Global Conformation of an Active Hammerhead RNA during the Process of Self-cleavage. Journal of Molecular Biology, 1996, 261, 125-134.	2.0	44
21	Hammering away at RNA global structure. Current Opinion in Structural Biology, 1996, 6, 317-321.	2.6	11
22	Global Conformation of a Self-Cleaving Hammerhead RNA. Biochemistry, 1994, 33, 13172-13177.	1.2	65
23	Interhelix Geometry of Stems I and II of a Self-Cleaving Hammerhead RNA. Biochemistry, 1994, 33, 1788-1796.	1.2	34
24	Heterozygous fragile X female: Historical, physical, cognitive, and cytogenetic features. American Journal of Medical Genetics Part A, 1991, 38, 269-274.	2.4	178
25	Fragile X checklist. American Journal of Medical Genetics Part A, 1991, 38, 283-287.	2.4	142
26	Mental impairment in cytogenetically positive fragile X females. American Journal of Medical Genetics Part A, 1991, 38, 503-504.	2.4	34