Abdullah

List of Publications by Citations

36 papers	482 citations	14 h-index	21 g-index
43 ext. papers	802 ext. citations	3.5 avg, IF	4.55 L-index

#	Paper	IF	Citations
36	Chloroplast genome of Hibiscus rosa-sinensis (Malvaceae): Comparative analyses and identification of mutational hotspots. *Genomics*, **2020**, 112, 581-591	4.3	53
35	Characterization of Withania somnifera chloroplast genome and its comparison with other selected species of Solanaceae. *Genomics*, **2020**, 112, 1522-1530	4.3	42
34	Comparative analyses of chloroplast genomes among three Firmiana species: Identification of mutational hotspots and phylogenetic relationship with other species of Malvaceae. *Plant Gene*, **2019**, 19, 100199	3.1	36
33	Chloroplast genome sequences of Artemisia maritima and Artemisia absinthium: Comparative analyses, mutational hotspots in genus Artemisia and phylogeny in family Asteraceae. *Genomics*, **2020**, 112, 1454-1463	4.3	36
32	Molecular evolution of chloroplast genomes in Monsteroideae (Araceae). *Planta*, **2020**, 251, 72	4.7	31
31	Evolutionary dynamics of chloroplast genomes in subfamily Aroideae (Araceae). *Genomics*, **2020**, 112, 2349-2360	4.3	28
30	Chloroplast Genome Sequence of : Comparative Analyses and Screening of Mutational Hotspots. *Plants*, **2019**, 8,	4.5	24
29	Correlations among oligonucleotide repeats, nucleotide substitutions, and insertion□deletion mutations in chloroplast genomes of plant family Malvaceae. *Journal of Systematics and Evolution*, **2021**, 59, 388-402	2.9	24
28	Comparative Plastomics of Ashwagandha (, Solanaceae) and Identification of Mutational Hotspots for Barcoding Medicinal Plants. *Plants*, **2020**, 9,	4.5	21
27	Plastid genomics of (Solanaceae): insights into molecular evolution, positive selection and the origin of the maternal genome of Aztec tobacco (). *PeerJ*, **2020**, 8, e9552	3.1	19
26	The GASA Gene Family in Cacao (Theobroma cacao, Malvaceae): Genome Wide Identification and Expression Analysis. *Agronomy*, **2021**, 11, 1425	3.6	18
25	A genome-wide approach to the comprehensive analysis of GASA gene family in Glycine max. *Plant Molecular Biology*, **2019**, 100, 607-620	4.6	17
24	Comparative analyses of chloroplast genomes of Theobroma cacao and Theobroma grandiflorum. *Biologia (Poland)*, **2020**, 75, 761-771	1.5	17
23	Complete Chloroplast Genomes of Anthurium huixtlense and Pothos scandens (Pothoideae, Araceae): Unique Inverted Repeat Expansion and Contraction Affect Rate of Evolution. *Journal of Molecular Evolution*, **2020**, 88, 562-574	3.1	16
22	Comparative plastome analysis of , with implications for genome evolution and phylogeny of Asteroideae. *Ecology and Evolution*, **2021**, 11, 7810-7826	2.8	14
21	A novel homozygous sequence variant in GLI1 underlies first case of autosomal recessive pre-axial polydactyly. *Clinical Genetics*, **2019**, 95, 540-541	4	13
20	Comparison of Chloroplast Genomes among Species of Unisexual and Bisexual Clades of the Monocot Family Araceae. *Plants*, **2020**, 9,	4.5	12

19	Chloroplast genome evolution in the Dracunculus clade (Aroideae, Araceae). *Genomics*, **2021**, 113, 183-192	4.3	10
18	Mutational Dynamics of Aroid Chloroplast Genomes II. *Frontiers in Genetics*, **2020**, 11, 610838	4.5	9
17	Magnesium transporter Gene Family: Genome-Wide Identification and Characterization in Theobroma cacao, Corchorus capsularis, and Gossypium hirsutum of Family Malvaceae. *Agronomy*, **2021**, 11, 1651	3.6	8
16	Variants in Cause Greig Cephalopolysyndactyly Syndrome. *Genetic Testing and Molecular Biomarkers* , **2019**, 23, 744-750	1.6	6
15	Whole-exome sequencing revealed a nonsense mutation in STKLD1 causing non-syndromic pre-axial polydactyly type A affecting only upper limb. *Clinical Genetics*, **2019**, 96, 134-139	4	4
14	Plastid genomics of Nicotiana (Solanaceae): insights into molecular evolution, positive selection and the origin of the maternal genome of Aztec tobacco (Nicotiana rustica)		4
13	Investigation and Computational Analysis of the Sulfotransferase (SOT) Gene Family in Potato (): Insights into Sulfur Adjustment for Proper Development and Stimuli Responses.. *Plants*, **2021**, 10,	4.5	3
12	Complete chloroplast genomes of Anthurium huixtlense and Pothos scandens (Pothoideae, Araceae): unique inverted repeat expansion and contraction affect rate of evolution		3
11	Sequence variants in the EDAR gene causing hypohidrotic ectodermal dysplasia. *Congenital Anomalies (discontinued)*, **2019**, 59, 145-147	1.1	2
10	Pseudogenization of the chloroplast threonine (trnT-GGU) gene in the sunflower family (Asteraceae). *Scientific Reports*, **2021**, 11, 21122	4.9	2
9	Agro-morphological, yield, and genotyping-by-sequencing data of selected wheat germplasm		2
8	Pseudogenization of trnT-GGU in chloroplast genomes of the plant family Asteraceae		2
7	The BAHD Gene Family in Cacao (Theobroma cacao, Malvaceae): Genome-Wide Identification and Expression Analysis. *Frontiers in Ecology and Evolution*, **2021**, 9,	3.7	1
6	A homozygous nonsense variant in DYM underlies Dyggve-Melchior-Clausen syndrome associated with ectodermal features. *Molecular Biology Reports*, **2020**, 47, 7083-7088	2.8	1
5	Agro-Morphological, Yield, and Genotyping-by-Sequencing Data of Selected Wheat () Germplasm From Pakistan. *Frontiers in Genetics*, **2021**, 12, 617772	4.5	1
4	Comparative Chloroplast Genomics in Phyllanthaceae Species. *Diversity*, **2021**, 13, 403	2.5	1
3	Clinical and genetic characterization of congenital lipoid adrenal hyperplasia. *Clinical Dysmorphology*, **2020**, 29, 173-176	0.9	0
2	A novel nonsense variant in EXOC8 underlies a neurodevelopmental disorder.. *Neurogenetics*, **2022**, 1	3	0

1	Comparative Chloroplast Genome Analyses of the Winter-Blooming Eastern Asian Endemic Genus (Calycanthaceae) With Implications For Its Phylogeny and Diversification.. *Frontiers in Genetics*, **2021**, 12, 709996	4.5