

Abdullah

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

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	A novel nonsense variant in EXOC8 underlies a neurodevelopmental disorder.. <i>Neurogenetics</i> , 2022 , 1	3	0
35	The BAHD Gene Family in Cacao (<i>Theobroma cacao</i> , Malvaceae): Genome-Wide Identification and Expression Analysis. <i>Frontiers in Ecology and Evolution</i> , 2021 , 9,	3.7	1
34	Comparative Chloroplast Genome Analyses of the Winter-Blooming Eastern Asian Endemic Genus (<i>Calycanthaceae</i>) With Implications For Its Phylogeny and Diversification.. <i>Frontiers in Genetics</i> , 2021 , 12, 709996	4.5	
33	Investigation and Computational Analysis of the Sulfotransferase (SOT) Gene Family in Potato (): Insights into Sulfur Adjustment for Proper Development and Stimuli Responses.. <i>Plants</i> , 2021 , 10,	4.5	3
32	Pseudogenization of the chloroplast threonine (trnT-GGU) gene in the sunflower family (<i>Asteraceae</i>). <i>Scientific Reports</i> , 2021 , 11, 21122	4.9	2
31	Agro-Morphological, Yield, and Genotyping-by-Sequencing Data of Selected Wheat () Germplasm From Pakistan. <i>Frontiers in Genetics</i> , 2021 , 12, 617772	4.5	1
30	Comparative plastome analysis of , with implications for genome evolution and phylogeny of <i>Asteroidae</i> . <i>Ecology and Evolution</i> , 2021 , 11, 7810-7826	2.8	14
29	The GASA Gene Family in Cacao (<i>Theobroma cacao</i> , Malvaceae): Genome Wide Identification and Expression Analysis. <i>Agronomy</i> , 2021 , 11, 1425	3.6	18
28	Correlations among oligonucleotide repeats, nucleotide substitutions, and insertion/deletion mutations in chloroplast genomes of plant family Malvaceae. <i>Journal of Systematics and Evolution</i> , 2021 , 59, 388-402	2.9	24
27	Chloroplast genome evolution in the <i>Dracunculus</i> clade (<i>Aroideae</i> , <i>Araceae</i>). <i>Genomics</i> , 2021 , 113, 183-192	4.2	10
26	Comparative Chloroplast Genomics in <i>Phyllanthaceae</i> Species. <i>Diversity</i> , 2021 , 13, 403	2.5	1
25	Magnesium transporter Gene Family: Genome-Wide Identification and Characterization in <i>Theobroma cacao</i> , <i>Corchorus capsularis</i> , and <i>Gossypium hirsutum</i> of Family Malvaceae. <i>Agronomy</i> , 2021 , 11, 1651	3.6	8
24	Comparative Plastomics of <i>Ashwagandha</i> (<i>Solanaceae</i>) and Identification of Mutational Hotspots for Barcoding Medicinal Plants. <i>Plants</i> , 2020 , 9,	4.5	21
23	Comparison of Chloroplast Genomes among Species of Unisexual and Bisexual Clades of the Monocot Family <i>Araceae</i> . <i>Plants</i> , 2020 , 9,	4.5	12
22	Complete Chloroplast Genomes of <i>Anthurium huixtlense</i> and <i>Pothos scandens</i> (<i>Pothoideae</i> , <i>Araceae</i>): Unique Inverted Repeat Expansion and Contraction Affect Rate of Evolution. <i>Journal of Molecular Evolution</i> , 2020 , 88, 562-574	3.1	16
21	Molecular evolution of chloroplast genomes in <i>Monsteroideae</i> (<i>Araceae</i>). <i>Planta</i> , 2020 , 251, 72	4.7	31
20	Evolutionary dynamics of chloroplast genomes in subfamily <i>Aroideae</i> (<i>Araceae</i>). <i>Genomics</i> , 2020 , 112, 2349-2360	4.3	28

19	Plastid genomics of (Solanaceae): insights into molecular evolution, positive selection and the origin of the maternal genome of Aztec tobacco (). <i>PeerJ</i> , 2020 , 8, e9552	3.1	19
18	Clinical and genetic characterization of congenital lipoid adrenal hyperplasia. <i>Clinical Dysmorphology</i> , 2020 , 29, 173-176	0.9	0
17	Comparative analyses of chloroplast genomes of Theobroma cacao and Theobroma grandiflorum. <i>Biologia (Poland)</i> , 2020 , 75, 761-771	1.5	17
16	A homozygous nonsense variant in DYM underlies Dyggve-Melchior-Clausen syndrome associated with ectodermal features. <i>Molecular Biology Reports</i> , 2020 , 47, 7083-7088	2.8	1
15	Chloroplast genome of Hibiscus rosa-sinensis (Malvaceae): Comparative analyses and identification of mutational hotspots. <i>Genomics</i> , 2020 , 112, 581-591	4.3	53
14	Characterization of Withania somnifera chloroplast genome and its comparison with other selected species of Solanaceae. <i>Genomics</i> , 2020 , 112, 1522-1530	4.3	42
13	Chloroplast genome sequences of Artemisia maritima and Artemisia absinthium: Comparative analyses, mutational hotspots in genus Artemisia and phylogeny in family Asteraceae. <i>Genomics</i> , 2020 , 112, 1454-1463	4.3	36
12	Mutational Dynamics of Aroid Chloroplast Genomes II. <i>Frontiers in Genetics</i> , 2020 , 11, 610838	4.5	9
11	A genome-wide approach to the comprehensive analysis of GASA gene family in Glycine max. <i>Plant Molecular Biology</i> , 2019 , 100, 607-620	4.6	17
10	Whole-exome sequencing revealed a nonsense mutation in STKLD1 causing non-syndromic pre-axial polydactyly type A affecting only upper limb. <i>Clinical Genetics</i> , 2019 , 96, 134-139	4	4
9	Sequence variants in the EDAR gene causing hypohidrotic ectodermal dysplasia. <i>Congenital Anomalies (discontinued)</i> , 2019 , 59, 145-147	1.1	2
8	Comparative analyses of chloroplast genomes among three Firmiana species: Identification of mutational hotspots and phylogenetic relationship with other species of Malvaceae. <i>Plant Gene</i> , 2019 , 19, 100199	3.1	36
7	Variants in Cause Greig Cephalopolysyndactyly Syndrome. <i>Genetic Testing and Molecular Biomarkers</i> , 2019 , 23, 744-750	1.6	6
6	Chloroplast Genome Sequence of : Comparative Analyses and Screening of Mutational Hotspots. <i>Plants</i> , 2019 , 8,	4.5	24
5	A novel homozygous sequence variant in GLI1 underlies first case of autosomal recessive pre-axial polydactyly. <i>Clinical Genetics</i> , 2019 , 95, 540-541	4	13
4	Plastid genomics of Nicotiana (Solanaceae): insights into molecular evolution, positive selection and the origin of the maternal genome of Aztec tobacco (Nicotiana rustica)		4
3	Complete chloroplast genomes of Anthurium huixtlense and Pothos scandens (Pothoideae, Araceae): unique inverted repeat expansion and contraction affect rate of evolution		3
2	Agro-morphological, yield, and genotyping-by-sequencing data of selected wheat germplasm		2

