

# Joo-Hwan Kim

## List of Publications by Year in descending order

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Version: 2024-02-01

59

papers

1,078

citations

430874

18

h-index

454955

30

g-index

59

all docs

59

docs citations

59

times ranked

1208

citing authors

#	ARTICLE	IF	CITATIONS
1	Phylogenetic relationship, biogeography, and conservation genetics of endangered <i>Fraxinus chiisanensis</i> (Oleaceae), endemic to South Korea. <i>Plant Diversity</i> , 2022, 44, 170-180.	3.7	2
2	Ethanol Extract of <i>Elaeagnus glabra</i> f. <i>oxyphylla</i> Branches Alleviates the Inflammatory Response Through Suppression of Cyclin D3/Cyclin-Dependent Kinase 11p58 Coupled to Lipopolysaccharide-Activated BV-2 Microglia. <i>Natural Product Communications</i> , 2022, 17, 1934578X2210750.	0.5	0
3	Disjunction and Vicariance Between East and West Asia: A Case Study on <i>Euonymus</i> sect. <i>Uniloculares</i> Based on Plastid Genome Analysis. <i>Frontiers in Plant Science</i> , 2022, 13, 825209.	3.6	0
4	Molecular Phylogeny and Historical Biogeography of <i>Goodyera</i> R. Br. (Orchidaceae): A Case of the Vicariance Between East Asia and North America. <i>Frontiers in Plant Science</i> , 2022, 13, 850170.	3.6	3
5	Complete chloroplast genomes shed light on phylogenetic relationships, divergence time, and biogeography of Allioideae (Amaryllidaceae). <i>Scientific Reports</i> , 2021, 11, 3262.	3.3	21
6	Simultaneous Quantification of Four Marker Compounds in <i>Bauhinia coccinea</i> Extract and Their Potential Inhibitory Effects on Alzheimer's Disease Biomarkers. <i>Plants</i> , 2021, 10, 702.	3.5	2
7	<i>Ficus erecta</i> Thunb Leaves Alleviate Memory Loss Induced by Scopolamine in Mice via Regulation of Oxidative Stress and Cholinergic System. <i>Molecular Neurobiology</i> , 2021, 58, 3665-3676.	4.0	7
8	<i>Ficus erecta</i> Thunb. Leaves Ameliorate Cognitive Deficit and Neuronal Damage in a Mouse Model of Amyloid- $\beta$ -Induced Alzheimer's Disease. <i>Frontiers in Pharmacology</i> , 2021, 12, 607403.	3.5	6
9	Insights into phylogenetic relationships and genome evolution of subfamily Commelinoidae (Commelinaceae Mirb.) inferred from complete chloroplast genomes. <i>BMC Genomics</i> , 2021, 22, 231.	2.8	18
10	Comparative analysis and implications of the chloroplast genomes of three thistles ( <i>Carduus</i> ) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5.0	2.0	5
11	POCU1b, the n-Butanol Soluble Fraction of <i>Polygoni Cuspidati Rhizoma et Radix</i> , Attenuates Obesity, Non-Alcoholic Fatty Liver, and Insulin Resistance via Inhibitions of Pancreatic Lipase, cAMP-Dependent PDE Activity, AMPK Activation, and SOCS-3 Suppression. <i>Nutrients</i> , 2020, 12, 3612.	4.1	8
12	Characterization of the complete chloroplast genome of Korean endemic, <i>Habenaria cruciformis</i> (Orchidaceae). <i>Mitochondrial DNA Part B: Resources</i> , 2020, 5, 3269-3271.	0.4	1
13	Rearâ€¢edge, lowâ€¢diversity, and haplotypic uniformity in coldâ€¢adapted <i>Bupleurum euphorbioides</i> interglacial refugia populations. <i>Ecology and Evolution</i> , 2020, 10, 10449-10462.	1.9	5
14	Catechol-Type Flavonoids from the Branches of <i>Elaeagnus glabra</i> f. <i>oxyphylla</i> Exert Antioxidant Activity and an Inhibitory Effect on Amyloid- $\beta$ Aggregation. <i>Molecules</i> , 2020, 25, 4917.	3.8	14
15	Implications of plastome evolution in the true lilies (monocot order Liliales). <i>Molecular Phylogenetics and Evolution</i> , 2020, 148, 106818.	2.7	23
16	Osteomeles schwerinae Extract and Its Major Compounds Inhibit Methylglyoxal-Induced Apoptosis in Human Retinal Pigment Epithelial Cells. <i>Molecules</i> , 2020, 25, 2605.	3.8	3
17	Epitypification of <i>Prunus</i> — <i>nudiflora</i> (Rosaceae), a natural hybrid species in Jeju Island, Korea. <i>Journal of Asia-Pacific Biodiversity</i> , 2019, 12, 718-720.	0.4	0
18	Autophagy Activation by <i>Crepidiastrum Denticulatum</i> Extract Attenuates Environmental Pollutant-Induced Damage in Dermal Fibroblasts. <i>International Journal of Molecular Sciences</i> , 2019, 20, 517.	4.1	10

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19	The implication of plastid transcriptome analysis in petaloid monocotyledons: A case study of <i>Lilium lancifolium</i> (Liliaceae, Liliales). <i>Scientific Reports</i> , 2019, 9, 6662.	3.3	3
20	<i>Elaeagnus glabra</i> f. <i>oxyphylla</i> Attenuates Scopolamine-Induced Learning and Memory Impairments in Mice by Improving Cholinergic Transmission via Activation of CREB/NGF Signaling. <i>Nutrients</i> , 2019, 11, 1205.	4.1	11
21	The newly developed single nucleotide polymorphism (SNP) markers for a potentially medicinal plant, <i>Crepidiastrum denticulatum</i> (Asteraceae), inferred from complete chloroplast genome data. <i>Molecular Biology Reports</i> , 2019, 46, 3287-3297.	2.3	11
22	Historical Biogeography of Melanthiaceae: A Case of Out-of-North America Through the Bering Land Bridge. <i>Frontiers in Plant Science</i> , 2019, 10, 396.	3.6	24
23	Improvement in Diabetic Retinopathy through Protection against Retinal Apoptosis in Spontaneously Diabetic Torii Rats Mediated by Ethanol Extract of <i>Osteomeles schwerinae</i> C.K. Schneid. <i>Nutrients</i> , 2019, 11, 546.	4.1	9
24	Molecular markers for phylogenetic applications derived from comparative plastome analysis of <i>Prunus</i> species. <i>Journal of Systematics and Evolution</i> , 2019, 57, 15-22.	3.1	7
25	Characterization of the complete chloroplast genome of <i>Fraxinus chiisanensis</i> (Oleaceae), an endemic to Korea. <i>Conservation Genetics Resources</i> , 2019, 11, 63-66.	0.8	3
26	Sequencing of the plastome in the leafless green mycoheterotroph <i>Cymbidium macrorhizon</i> helps us to understand an early stage of fully mycoheterotrophic plastome structure. <i>Plant Systematics and Evolution</i> , 2018, 304, 245-258.	0.9	21
27	Draft genome sequence of wild <i>Prunus yedoensis</i> reveals massive inter-specific hybridization between sympatric flowering cherries. <i>Genome Biology</i> , 2018, 19, 127.	8.8	89
28	Updated molecular phylogenetic analysis, dating and biogeographical history of the lily family (Liliaceae: Liliales). <i>Botanical Journal of the Linnean Society</i> , 2018, 187, 579-593.	1.6	33
29	Molecular Phylogeny and Dating of Forsythieae (Oleaceae) Provide Insight into the Miocene History of Eurasian Temperate Shrubs. <i>Frontiers in Plant Science</i> , 2018, 9, 99.	3.6	32
30	Development of 26 microsatellite markers in <i>Bupleurum latissimum</i> (Apiaceae), an endangered plant endemic to Ulleung Island, Korea. <i>Applications in Plant Sciences</i> , 2018, 6, e1144.	2.1	2
31	Molecular systematics and historical biogeography of <i>Maianthemum</i> s.s.. <i>American Journal of Botany</i> , 2017, 104, 939-952.	1.7	9
32	Genomic clues to the parental origin of the wild flowering cherry <i>Prunus yedoensis</i> var. <i>nudiflora</i> (Rosaceae). <i>Plant Biotechnology Reports</i> , 2017, 11, 449-459.	1.5	14
33	A Dynamic Tandem Repeat in Monocotyledons Inferred from a Comparative Analysis of Chloroplast Genomes in Melanthiaceae. <i>Frontiers in Plant Science</i> , 2017, 8, 693.	3.6	13
34	New method for an evaluation of the esthetical improvements resulting from a mandibular angle reduction. <i>Journal of the Korean Association of Oral and Maxillofacial Surgeons</i> , 2017, 43, 239.	0.8	3
35	The complete plastid genome sequence of <i>Bomarea edulis</i> (Alstroemeriaceae: Liliales). <i>Mitochondrial DNA</i> , 2016, 27, 1-2.	0.6	1
36	Insight into infrageneric circumscription through complete chloroplast genome sequences of two Trillium species. <i>AoB PLANTS</i> , 2016, 8, plw015.	2.3	17

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37	Molecular phylogenetic relationships of Melanthiaceae (Liliales) based on plastid DNA sequences. Botanical Journal of the Linnean Society, 2016, 181, 567-584.		1.6	19
38	The complete chloroplast genome sequence of <i>Lilium tsingtauense</i> Gilg (sect. Martagon,) Tj ETQq0 0 0 rgBT <sub>0.4</sub> Overlock <sub>10</sub> Tf 50 T <sub>7</sub>			
39	Development of Chloroplast Microsatellite Markers for the Endangered <i>Maianthemum bicolor</i> (Asparagaceae s.l.). Applications in Plant Sciences, 2016, 4, 1600032.		2.1	18
40	Complete plastid genome of <i>Astragalus mongolicus</i> var. <i>nakaianus</i> (Fabaceae). Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis, 2016, 27, 2838-2839.		0.7	6
41	Phylogenetic Inferences and the Evolution of Plastid DNA in Campynemataceae and the Mycoheterotrophic <i>Corsia dispar</i> D.L Jones & B. Gray (Corsiaceae). Plant Molecular Biology Reporter, 2016, 34, 192-210.		1.8	7
42	The complete plastid genome sequence of <i>Eustrephus latifolius</i> (Asparagaceae: Lomandroideae). Mitochondrial DNA, 2016, 27, 1549-1551.		0.6	5
43	Seven New Complete Plastome Sequences Reveal Rampant Independent Loss of the <i>ndh</i> Gene Family across Orchids and Associated Instability of the Inverted Repeat/Small Single-Copy Region Boundaries. PLoS ONE, 2015, 10, e0142215.		2.5	131
44	The complete chloroplast genome of colchicine plants ( <i>Colchicum autumnale</i> L. and <i>Gloriosa superba</i> ) Tj ETQq0 0 0 rgBT <sub>3.2</sub> Overlock <sub>10</sub> Tf 42			
45	The Largest Plastid Genome of Monocots: a Novel Genome Type Containing AT Residue Repeats in the Slipper Orchid <i>Cypripedium japonicum</i> . Plant Molecular Biology Reporter, 2015, 33, 1210-1220.		1.8	23
46	Molecular Identification of <i>Reynoutria japonica</i> Houtt. and <i>R. sachalinensis</i> (F. Schmidt) Nakai Using SNP Sites. Korean Journal of Plant Resources, 2015, 28, 743-751.		0.2	2
47	A <i>trnl_CAU</i> Triplication Event in the Complete Chloroplast Genome of <i>Paris verticillata</i> M.Bieb. (Melanthiaceae, Liliales). Genome Biology and Evolution, 2014, 6, 1699-1706.		2.5	26
48	Undifferentiated Pleomorphic Sarcoma in Mandible. Maxillofacial Plastic and Reconstructive Surgery, 2014, 36, 303-307.		1.8	9
49	Osteomyelitis in an Osteopathia Striata with Cranial Sclerosis Patient. Maxillofacial Plastic and Reconstructive Surgery, 2014, 36, 285-291.		1.8	1
50	Complete Chloroplast Genome of <i>Chionographis japonica</i> (Willd.) Maxim. (Melanthiaceae): Comparative Genomics and Evaluation of Universal Primers for Liliales. Plant Molecular Biology Reporter, 2013, 31, 1407-1421.		1.8	26
51	Comparative genomics of four Liliales families inferred from the complete chloroplast genome sequence of <i>Veratrum patulum</i> O. Loes. (Melanthiaceae). Gene, 2013, 530, 229-235.		2.2	47
52	Familial relationships of the monocot order Liliales based on a molecular phylogenetic analysis using four plastid loci: <i>matK</i> , <i>rbcL</i> , <i>atpB</i> and <i>atpF</i> - <i>H</i> . Botanical Journal of the Linnean Society, 2013, 172, 5-21.		1.6	50
53	Molecular phylogenetic relationships and implications for the circumscription of Colchicaceae (Liliales). Botanical Journal of the Linnean Society, 2013, 172, 255-269.		1.6	10
54	Networks in a Large-Scale Phylogenetic Analysis: Reconstructing Evolutionary History of Asparagales (Liliinae) Based on Four Plastid Genes. PLoS ONE, 2013, 8, e59472.		2.5	58

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55	Comparative Genome Analysis and Phylogenetic Relationship of Order Liliales Insight from the Complete Plastid Genome Sequences of Two Lilies ( <i>Lilium longiflorum</i> and <i>Alstroemeria aurea</i> ). PLoS ONE, 2013, 8, e68180.	2.5	39
56	Molecular identification of <i>Schisandra chinensis</i> and its allied species using multiplex PCR based on SNPs. Genes and Genomics, 2012, 34, 283-290.	1.4	12
57	The phylogenetic relationships of Asparagales in Korea based on five plastid DNA regions. Journal of Plant Biology, 2012, 55, 325-341.	2.1	14
58	Molecular phylogenetics of Ruscaceae sensu lato and related families (Asparagales) based on plastid and nuclear DNA sequences. Annals of Botany, 2010, 106, 775-790.	2.9	71
59	Single-step separation of bioactive flavonol glucosides from <i>Osteomeles schwerinae</i> by high-speed counter-current chromatography. Journal of Separation Science, 2010, 33, 582-586.	2.5	24