Sang-Hun Oh

List of Publications by Year
in descending order

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34	481	758635 12	713013 21
papers	citations	h-index	g-index

34	34	34	507
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Genetic variation of the endangered species Halenia coreana (Gentianaceae). Korean Journal of Plant Taxonomy, 2022, 52, 45-53.	0.3	1
2	The complete chloroplast genome of Aruncus aethusifolius (Rosaceae), a species endemic to Korea. Korean Journal of Plant Taxonomy, 2022, 52, 118-122.	0.3	2
3	The complete chloroplast genome of Aruncus dioicus var. kamtschaticus (Rosaceae). Mitochondrial DNA Part B: Resources, 2021, 6, 1256-1258.	0.2	6
4	Phylogenetic position of Daphne genkwa (Thymelaeaceae) inferred from complete chloroplast data. Korean Journal of Plant Taxonomy, 2021, 51, 171-175.	0.3	11
5	The complete chloroplast genome of Campanula takesimana Nakai from Dokdo Island in Korea (Campanulaceae). Mitochondrial DNA Part B: Resources, 2021, 6, 135-137.	0.2	8
6	The complete chloroplast genome of Diarthron linifolium (Thymelaeaceae), a species found on a limestone outcrop in eastern Asia. Korean Journal of Plant Taxonomy, 2021, 51, 345-352.	0.3	7
7	(127â€“135) Proposals to add new Provisions and Recommendations to Division <scp>III</scp> of the International Code of Nomenclature for algae, fungi, and plants related to virtual participation in the Nomenclature Section. Taxon, 2021, 70, 1397-1398.	0.4	2
8	Report of the Specialâ€purpose Committee on Virtual Participation in the Nomenclature Section. Taxon, 2021, 70, 1399-1401.	0.4	2
9	The complete chloroplast genome of Euscaphis japonica (Thunb.) Kanitz (Staphyleaceae) isolated in Korea. Mitochondrial DNA Part B: Resources, 2020, 5, 3751-3753.	0.2	7
10	A second complete chloroplast genome sequence of Fagus multinervis Nakai (Fagaceae): intraspecific variations on chloroplast genome. Mitochondrial DNA Part B: Resources, 2020, 5, 1868-1869.	0.2	17
11	The second complete chloroplast genome sequence of the Viburnum erosum (Adoxaceae) showed a low level of intra-species variations. Mitochondrial DNA Part B: Resources, 2020, 5, 271-272.	0.2	16
12	Comparative chloroplast genomics and phylogenetic analysis of the <i>Viburnum dilatatum</i> complex (Adoxaceae) in Korea. Korean Journal of Plant Taxonomy, 2020, 50, 8-16.	0.3	18
13	The complete chloroplast genome of an endangered species in Korea, Halenia corniculata (L.) Cornaz (Gentianaceae). Mitochondrial DNA Part B: Resources, 2019, 4, 1539-1540.	0.2	7
14	The complete chloroplast genome sequence of Viburnum erosum (Adoxaceae). Mitochondrial DNA Part B: Resources, 2019, 4, 3278-3279.	0.2	18
15	The complete chloroplast genome of the traditional medicinal plant Stellera chamaejasme L. (Thymelaeaceae). Mitochondrial DNA Part B: Resources, 2019, 4, 1796-1797.	0.2	8
16	The complete chloroplast genome sequence of Goodyera schlechtendaliana in Korea (Orchidaceae). Mitochondrial DNA Part B: Resources, 2019, 4, 2692-2693.	0.2	17
17	The complete chloroplast genome sequence of a morphotype of Goodyera schlechtendaliana (Orchidaceae) with the column appendages. Mitochondrial DNA Part B: Resources, 2019, 4, 626-627.	0.2	18
18	A comparative morphological study of Viburnum (Adoxaceae) in Korea. Korean Journal of Plant Taxonomy, 2019, 49, 107-117.	0.3	6

#	Article	IF	Citations
19	New taxa of Rhododendron tschonoskii alliance (Ericaceae) from East Asia. PhytoKeys, 2019, 134, 97-114.	0.4	6
20	Phylogenetic analysis of Viburnum (Adoxaceae) in Korea using DNA sequences. Korean Journal of Plant Taxonomy, 2018, 48, 206-217.	0.3	7
21	A comparative morphological study of Thymelaeaceae in Korea. Korean Journal of Plant Taxonomy, 2017, 47, 207-221.	0.3	7
22	Phylogeny and Evolution of Endemic Species on Ulleungdo Island, Korea: The Case of <I>Fagus multinervis</I> (Fagaceae). Systematic Botany, 2016, 41, 617-625.	0.2	10
23	Taxonomy of tribe Neillieae (Rosaceae): Neillia. Korean Journal of Plant Taxonomy, 2016, 46, 13-32.	0.3	2
24	Cytokinin-dependent secondary growth determines root biomass in radish (Raphanus sativusL.). Journal of Experimental Botany, 2015, 66, 4607-4619.	2.4	47
25	Sea, wind, or bird: Origin of <i>Fagus multinervis</i> (Fagaceae) inferred from chloroplast DNA sequences. Korean Journal of Plant Taxonomy, 2015, 45, 213-220.	0.3	10
26	Molecular evidence for hybrid origin of Aster chusanensis, an endemic species of Ulleungdo, Korea. Journal of Plant Biology, 2014, 57, 174-185.	0.9	15
27	Phylogenetic analysis of PISTILLATA sequences in Neillia (Rosaceae). Journal of Plant Biology, 2013, 56, 145-151.	0.9	3
28	Peaches and almonds: phylogeny of Prunus subg. Amygdalus (Rosaceae) based on DNA sequences and morphology. Plant Systematics and Evolution, 2013, 299, 1403-1418.	0.3	33
29	Description and Phylogenetic Position of a New Angiosperm Family, Guamatelaceae, Inferred from Chloroplast <I>rbcL, atpB</I>, and <I>matK</I> Sequences. Systematic Botany, 2006, 31, 730-738.	0.2	14
30	Neillia Includes Stephanandra (Rosaceae). Novon, 2006, 16, 91-95.	0.3	12
31	Molecular phylogenetic systematics and biogeography of tribe Neillieae (Rosaceae) using DNA sequences of cpDNA, rDNA, and LEAFY. American Journal of Botany, 2005, 92, 179-192.	0.8	72
32	(1619) Proposal to conserve the name Physocarpus opulifolius (L.) Maxim. against Physocarpus opulifolius Raf. (Rosaceae). Taxon, 2004, 53, 212-213.	0.4	1
33	Phylogenetic utility of the second intron of LEAFY in Neillia and Stephanandra (Rosaceae) and implications for the origin of Stephanandra. Molecular Phylogenetics and Evolution, 2003, 29, 203-215.	1.2	70
34	A New Species of Goodyera (Orchidaceae: Orchidoideae) from Korea and Japan. Journal of Plant Biology, 0, , .	0.9	1