

# Sang-Hun Oh

## List of Publications by Year in descending order

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Version: 2024-02-01

34  
papers

481  
citations

758635

12  
h-index

713013

21  
g-index

34  
all docs

34  
docs citations

34  
times ranked

507  
citing authors

#	ARTICLE	IF	CITATIONS
1	Molecular phylogenetic systematics and biogeography of tribe Neillieae (Rosaceae) using DNA sequences of cpDNA, rDNA, and LEAFY. <i>American Journal of Botany</i> , 2005, 92, 179-192.	0.8	72
2	Phylogenetic utility of the second intron of LEAFY in <i>Neillia</i> and <i>Stephanandra</i> (Rosaceae) and implications for the origin of <i>Stephanandra</i> . <i>Molecular Phylogenetics and Evolution</i> , 2003, 29, 203-215.	1.2	70
3	Cytokinin-dependent secondary growth determines root biomass in radish ( <i>Raphanus sativus</i> L.). <i>Journal of Experimental Botany</i> , 2015, 66, 4607-4619.	2.4	47
4	Peaches and almonds: phylogeny of <i>Prunus</i> subg. <i>Amygdalus</i> (Rosaceae) based on DNA sequences and morphology. <i>Plant Systematics and Evolution</i> , 2013, 299, 1403-1418.	0.3	33
5	The complete chloroplast genome sequence of <i>Viburnum erosum</i> (Adoxaceae). <i>Mitochondrial DNA Part B: Resources</i> , 2019, 4, 3278-3279.	0.2	18
6	The complete chloroplast genome sequence of a morphotype of <i>Goodyera schlechtendaliana</i> (Orchidaceae) with the column appendages. <i>Mitochondrial DNA Part B: Resources</i> , 2019, 4, 626-627.	0.2	18
7	Comparative chloroplast genomics and phylogenetic analysis of the <i>Viburnum dilatatum</i> complex (Adoxaceae) in Korea. <i>Korean Journal of Plant Taxonomy</i> , 2020, 50, 8-16.	0.3	18
8	The complete chloroplast genome sequence of <i>Goodyera schlechtendaliana</i> in Korea (Orchidaceae). <i>Mitochondrial DNA Part B: Resources</i> , 2019, 4, 2692-2693.	0.2	17
9	A second complete chloroplast genome sequence of <i>Fagus multinervis</i> Nakai (Fagaceae): intraspecific variations on chloroplast genome. <i>Mitochondrial DNA Part B: Resources</i> , 2020, 5, 1868-1869.	0.2	17
10	The second complete chloroplast genome sequence of the <i>Viburnum erosum</i> (Adoxaceae) showed a low level of intra-species variations. <i>Mitochondrial DNA Part B: Resources</i> , 2020, 5, 271-272.	0.2	16
11	Molecular evidence for hybrid origin of <i>Aster chusanensis</i> , an endemic species of Ulleungdo, Korea. <i>Journal of Plant Biology</i> , 2014, 57, 174-185.	0.9	15
12	Description and Phylogenetic Position of a New Angiosperm Family, Guamatelaceae, Inferred from Chloroplast <i>rbcL</i> , <i>atpB</i> , and <i>matK</i> Sequences. <i>Systematic Botany</i> , 2006, 31, 730-738.	0.2	14
13	<i>Neillia</i> Includes <i>Stephanandra</i> (Rosaceae). <i>Novon</i> , 2006, 16, 91-95.	0.3	12
14	Phylogenetic position of <i>Daphne genkwa</i> (Thymelaeaceae) inferred from complete chloroplast data. <i>Korean Journal of Plant Taxonomy</i> , 2021, 51, 171-175.	0.3	11
15	Phylogeny and Evolution of Endemic Species on Ulleungdo Island, Korea: The Case of <i>Fagus multinervis</i> (Fagaceae). <i>Systematic Botany</i> , 2016, 41, 617-625.	0.2	10
16	Sea, wind, or bird: Origin of <i>Fagus multinervis</i> (Fagaceae) inferred from chloroplast DNA sequences. <i>Korean Journal of Plant Taxonomy</i> , 2015, 45, 213-220.	0.3	10
17	The complete chloroplast genome of the traditional medicinal plant <i>Stellera chamaejasme</i> L. (Thymelaeaceae). <i>Mitochondrial DNA Part B: Resources</i> , 2019, 4, 1796-1797.	0.2	8
18	The complete chloroplast genome of <i>Campanula takesimana</i> Nakai from Dokdo Island in Korea (Campanulaceae). <i>Mitochondrial DNA Part B: Resources</i> , 2021, 6, 135-137.	0.2	8

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19	The complete chloroplast genome of an endangered species in Korea, <i>Halenia corniculata</i> (L.) Cornaz (Gentianaceae). Mitochondrial DNA Part B: Resources, 2019, 4, 1539-1540.	0.2	7
20	The complete chloroplast genome of <i>Euscaphis japonica</i> (Thunb.) Kanitz (Staphyleaceae) isolated in Korea. Mitochondrial DNA Part B: Resources, 2020, 5, 3751-3753.	0.2	7
21	A comparative morphological study of Thymelaeaceae in Korea. Korean Journal of Plant Taxonomy, 2017, 47, 207-221.	0.3	7
22	Phylogenetic analysis of <i>Viburnum</i> (Adoxaceae) in Korea using DNA sequences. Korean Journal of Plant Taxonomy, 2018, 48, 206-217.	0.3	7
23	The complete chloroplast genome of <i>Diarthron linifolium</i> (Thymelaeaceae), a species found on a limestone outcrop in eastern Asia. Korean Journal of Plant Taxonomy, 2021, 51, 345-352.	0.3	7
24	The complete chloroplast genome of <i>Aruncus dioicus</i> var. <i>kamtschaticus</i> (Rosaceae). Mitochondrial DNA Part B: Resources, 2021, 6, 1256-1258.	0.2	6
25	A comparative morphological study of <i>Viburnum</i> (Adoxaceae) in Korea. Korean Journal of Plant Taxonomy, 2019, 49, 107-117.	0.3	6
26	New taxa of <i>Rhododendron tschonoskii</i> alliance (Ericaceae) from East Asia. PhytoKeys, 2019, 134, 97-114.	0.4	6
27	Phylogenetic analysis of PISTILLATA sequences in <i>Neillia</i> (Rosaceae). Journal of Plant Biology, 2013, 56, 145-151.	0.9	3
28	Taxonomy of tribe Neillieae (Rosaceae): <i>Neillia</i> . Korean Journal of Plant Taxonomy, 2016, 46, 13-32.	0.3	2
29	(127-135) Proposals to add new Provisions and Recommendations to Division III 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
30	Report of the Special Purpose Committee on Virtual Participation in the Nomenclature Section. Taxon, 2021, 70, 1399-1401.	0.4	2
31	The complete chloroplast genome of <i>Aruncus aethusifolius</i> (Rosaceae), a species endemic to Korea. Korean Journal of Plant Taxonomy, 2022, 52, 118-122.	0.3	2
32	(1619) Proposal to conserve the name <i>Physocarpus opulifolius</i> (L.) Maxim. against <i>Physocarpus opulifolius</i> Raf. (Rosaceae). Taxon, 2004, 53, 212-213.	0.4	1
33	Genetic variation of the endangered species <i>Halenia coreana</i> (Gentianaceae). Korean Journal of Plant Taxonomy, 2022, 52, 45-53.	0.3	1
34	A New Species of <i>Goodyera</i> (Orchidaceae: Orchidoideae) from Korea and Japan. Journal of Plant Biology, 0, .	0.9	1