

# Jongsun Park

## List of Publications by Citations

**Source:** <https://exaly.com/author-pdf/7557240/jongsun-park-publications-by-citations.pdf>

**Version:** 2024-04-20

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

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

139  
papers

5,893  
citations

30  
h-index

73  
g-index

206  
ext. papers

7,210  
ext. citations

2.4  
avg, IF

5.93  
L-index

#	Paper	IF	Citations
139	Comparative genomics reveals mobile pathogenicity chromosomes in <i>Fusarium</i> . <i>Nature</i> , <b>2010</b> , 464, 367-73	50.4	1085
138	The Amborella genome and the evolution of flowering plants. <i>Science</i> , <b>2013</b> , 342, 1241089	33.3	546
137	The plant cell wall-decomposing machinery underlies the functional diversity of forest fungi. <i>Science</i> , <b>2011</b> , 333, 762-5	33.3	417
136	Comparative genomics yields insights into niche adaptation of plant vascular wilt pathogens. <i>PLoS Pathogens</i> , <b>2011</b> , 7, e1002137	7.6	335
135	Internet-accessible DNA sequence database for identifying fusaria from human and animal infections. <i>Journal of Clinical Microbiology</i> , <b>2010</b> , 48, 3708-18	9.7	315
134	Genome-wide functional analysis of pathogenicity genes in the rice blast fungus. <i>Nature Genetics</i> , <b>2007</b> , 39, 561-5	36.3	183
133	Insight into trade-off between wood decay and parasitism from the genome of a fungal forest pathogen. <i>New Phytologist</i> , <b>2012</b> , 194, 1001-1013	9.8	168
132	Homeobox transcription factors are required for conidiation and appressorium development in the rice blast fungus <i>Magnaporthe oryzae</i> . <i>PLoS Genetics</i> , <b>2009</b> , 5, e1000757	6	143
131	Fungal secretome database: integrated platform for annotation of fungal secretomes. <i>BMC Genomics</i> , <b>2010</b> , 11, 105	4.5	128
130	FTFD: an informatics pipeline supporting phylogenomic analysis of fungal transcription factors. <i>Bioinformatics</i> , <b>2008</b> , 24, 1024-5	7.2	119
129	A putative MAP kinase kinase kinase, MCK1, is required for cell wall integrity and pathogenicity of the rice blast fungus, <i>Magnaporthe oryzae</i> . <i>Molecular Plant-Microbe Interactions</i> , <b>2008</b> , 21, 525-34	3.6	109
128	Systematic and searchable classification of cytochrome P450 proteins encoded by fungal and oomycete genomes. <i>BMC Genomics</i> , <b>2012</b> , 13, 525	4.5	99
127	Fungal cytochrome P450 database. <i>BMC Genomics</i> , <b>2008</b> , 9, 402	4.5	96
126	MoCRZ1, a gene encoding a calcineurin-responsive transcription factor, regulates fungal growth and pathogenicity of <i>Magnaporthe oryzae</i> . <i>Fungal Genetics and Biology</i> , <b>2009</b> , 46, 243-54	3.9	95
125	Taxonomic status of the <i>Bemisia tabaci</i> complex (Hemiptera: Aleyrodidae) and reassessment of the number of its constituent species. <i>PLoS ONE</i> , <b>2013</b> , 8, e63817	3.7	92
124	In-depth insight into in vivo apoplastic secretome of rice- <i>Magnaporthe oryzae</i> interaction. <i>Journal of Proteomics</i> , <b>2013</b> , 78, 58-71	3.9	80
123	CFGFP: a web-based, comparative fungal genomics platform. <i>Nucleic Acids Research</i> , <b>2008</b> , 36, D562-71	20.1	72

122	Genome-wide analysis of T-DNA integration into the chromosomes of <i>Magnaporthe oryzae</i> . <i>Molecular Microbiology</i> , <b>2007</b> , 66, 371-82	4.1	69
121	Simple sequence repeats in <i>Neurospora crassa</i> : distribution, polymorphism and evolutionary inference. <i>BMC Genomics</i> , <b>2008</b> , 9, 31	4.5	63
120	De novo transcriptome sequencing of <i>Momordica cochinchinensis</i> to identify genes involved in the carotenoid biosynthesis. <i>Plant Molecular Biology</i> , <b>2012</b> , 79, 413-27	4.6	61
119	Whole transcriptome analyses of six thoroughbred horses before and after exercise using RNA-Seq. <i>BMC Genomics</i> , <b>2012</b> , 13, 473	4.5	56
118	Cyber infrastructure for <i>Fusarium</i> : three integrated platforms supporting strain identification, phylogenetics, comparative genomics and knowledge sharing. <i>Nucleic Acids Research</i> , <b>2011</b> , 39, D640-6	20.1	56
117	Phytophthora Database: A Forensic Database Supporting the Identification and Monitoring of <i>Phytophthora</i> . <i>Plant Disease</i> , <b>2008</b> , 92, 966-972	1.5	51
116	Comprehensive genome- and transcriptome-wide analyses of mutations associated with microsatellite instability in Korean gastric cancers. <i>Genome Research</i> , <b>2013</b> , 23, 1109-17	9.7	49
115	Evolution of the large genome in <i>Capsicum annuum</i> occurred through accumulation of single-type long terminal repeat retrotransposons and their derivatives. <i>Plant Journal</i> , <b>2012</b> , 69, 1018-29	6.9	46
114	Combining ChIP-chip and expression profiling to model the MoCRZ1 mediated circuit for Ca/calciueurin signaling in the rice blast fungus. <i>PLoS Pathogens</i> , <b>2010</b> , 6, e1000909	7.6	46
113	Complete sequencing and comparative analyses of the pepper ( <i>Capsicum annuum</i> L.) plastome revealed high frequency of tandem repeats and large insertion/deletions on pepper plastome. <i>Plant Cell Reports</i> , <b>2011</b> , 30, 217-29	5.1	45
112	Global expression profiling of transcription factor genes provides new insights into pathogenicity and stress responses in the rice blast fungus. <i>PLoS Pathogens</i> , <b>2013</b> , 9, e1003350	7.6	43
111	The PEX7-mediated peroxisomal import system is required for fungal development and pathogenicity in <i>Magnaporthe oryzae</i> . <i>PLoS ONE</i> , <b>2011</b> , 6, e28220	3.7	40
110	Identification and analysis of in planta expressed genes of <i>Magnaporthe oryzae</i> . <i>BMC Genomics</i> , <b>2010</b> , 11, 104	4.5	34
109	Comparative analysis of pepper and tomato reveals euchromatin expansion of pepper genome caused by differential accumulation of Ty3/Gypsy-like elements. <i>BMC Genomics</i> , <b>2011</b> , 12, 85	4.5	30
108	Comparison of Whole Plastome Sequences between Thermogenic Skunk Cabbage and Nonthermogenic ( <i>Orontioideae</i> ; <i>Araceae</i> ) in East Asia. <i>International Journal of Molecular Sciences</i> , <b>2019</b> , 20,	6.3	22
107	The complete mitochondrial genome of ( <i>Fallb</i> , 1826) (Hemiptera: <i>Delphacidae</i> ) collected in a mid-Western part of Korean peninsula. <i>Mitochondrial DNA Part B: Resources</i> , <b>2019</b> , 4, 2229-2230	0.5	18
106	Peeling back the evolutionary layers of molecular mechanisms responsive to exercise-stress in the skeletal muscle of the racing horse. <i>DNA Research</i> , <b>2013</b> , 20, 287-98	4.5	18
105	The complete chloroplast genome of Korean Maxim. (): providing genetic background of two types of. <i>Mitochondrial DNA Part B: Resources</i> , <b>2019</b> , 4, 2424-2425	0.5	17

104	SysPIMP: the web-based systematical platform for identifying human disease-related mutated sequences from mass spectrometry. <i>Nucleic Acids Research</i> , <b>2009</b> , 37, D913-20	20.1	17
103	SNUGB: a versatile genome browser supporting comparative and functional fungal genomics. <i>BMC Genomics</i> , <b>2008</b> , 9, 586	4.5	17
102	The complete mitochondrial genome of the fall armyworm, Smith, 1797 (Lepidoptera; Noctuidae), firstly collected in Korea. <i>Mitochondrial DNA Part B: Resources</i> , <b>2019</b> , 4, 3918-3920	0.5	16
101	Comparative chloroplast genomics and phylogenetic analysis of the <i>Viburnum dilatatum</i> complex (Adoxaceae) in Korea. <i>Korean Journal of Plant Taxonomy</i> , <b>2020</b> , 50, 8-16	0.5	16
100	The Complete Chloroplast Genome of Isolated in Korea (Brassicaceae): An Investigation of Intraspecific Variations of the Chloroplast Genome of Korean. <i>International Journal of Genomics</i> , <b>2020</b> , 2020, 3236461	2.5	16
99	IMGD: an integrated platform supporting comparative genomics and phylogenetics of insect mitochondrial genomes. <i>BMC Genomics</i> , <b>2009</b> , 10, 148	4.5	15
98	The complete mitochondrial genome of (Fallb, 1826) (Hemiptera: Delphacidae) collected in a southern part of Korean peninsula. <i>Mitochondrial DNA Part B: Resources</i> , <b>2019</b> , 4, 2242-2243	0.5	14
97	The complete mitochondrial genome of Glover, 1877 (Hemiptera: Aphididae) collected in Korean peninsula. <i>Mitochondrial DNA Part B: Resources</i> , <b>2019</b> , 4, 3007-3009	0.5	13
96	A complete chloroplast genome sequence of (Orchidaceae) represents high sequence variation in the species. <i>Mitochondrial DNA Part B: Resources</i> , <b>2020</b> , 5, 517-519	0.5	13
95	The complete mitochondrial genome of new species candidate of (Rosaceae). <i>Mitochondrial DNA Part B: Resources</i> , <b>2020</b> , 5, 3435-3437	0.5	13
94	The complete chloroplast genome sequence of in Korea (Orchidaceae). <i>Mitochondrial DNA Part B: Resources</i> , <b>2019</b> , 4, 2692-2693	0.5	12
93	The Comparative Analyses of Six Complete Chloroplast Genomes of Morphologically Diverse L. (Amaranthaceae) Collected in Korea. <i>International Journal of Genomics</i> , <b>2021</b> , 2021, 6643444	2.5	12
92	The complete chloroplast genome of candidate new species from in Korea (Rosaceae). <i>Mitochondrial DNA Part B: Resources</i> , <b>2019</b> , 4, 2433-2435	0.5	11
91	The second complete chloroplast genome sequence of the (Adoxaceae) showed a low level of intra-species variations. <i>Mitochondrial DNA Part B: Resources</i> , <b>2019</b> , 5, 271-272	0.5	11
90	The complete mitochondrial genome of Glover, 1877 (Hemiptera: Aphididae) isolated from in Korea. <i>Mitochondrial DNA Part B: Resources</i> , <b>2020</b> , 5, 2878-2880	0.5	11
89	Genome-wide comparative analyses of GATA transcription factors among 19 Arabidopsis ecotype genomes: Intraspecific characteristics of GATA transcription factors. <i>PLoS ONE</i> , <b>2021</b> , 16, e0252181	3.7	11
88	The complete chloroplast genome of Ledeb. isolated in Korea (Rosaceae): investigation of intraspecific variations on its chloroplast genomes. <i>Mitochondrial DNA Part B: Resources</i> , <b>2020</b> , 5, 2264-2266	0.5	10
87	Comparative transcriptome analysis of three color variants of the sea cucumber <i>Apostichopus japonicus</i> . <i>Marine Genomics</i> , <b>2016</b> , 28, 21-24	1.9	10

86	The complete chloroplast genome sequence of (Adoxaceae). <i>Mitochondrial DNA Part B: Resources</i> , <b>2019</b> , 4, 3278-3279	0.5	9
85	Comprehensive Analyses of the Complete Mitochondrial Genome of <i>Figulus binodulus</i> (Coleoptera: Lucanidae). <i>Journal of Insect Science</i> , <b>2020</b> , 20,	2	9
84	The complete chloroplast and mitochondrial genomes of <i>Hyunsasi tree, x</i> (Salicaceae). <i>Mitochondrial DNA Part B: Resources</i> , <b>2019</b> , 4, 2521-2522	0.5	8
83	The complete chloroplast genome of (H. L.) <i>Rapaics</i> (Ranunculaceae). <i>Mitochondrial DNA Part B: Resources</i> , <b>2019</b> , 4, 3404-3406	0.5	8
82	Complete Genome Sequence of the Endosymbiont of. <i>Microbiology Resource Announcements</i> , <b>2020</b> , 9,	1.3	8
81	The complete mitochondrial genome of <i>L.</i> (Ricciaceae, Marchantiophyta): investigation of intraspecific variations on mitochondrial genomes of. <i>Mitochondrial DNA Part B: Resources</i> , <b>2020</b> , 5, 1220-1222	0.5	7
80	The complete mitochondrial genome of (Smith, 1874) (Hymenoptera: Formicidae). <i>Mitochondrial DNA Part B: Resources</i> , <b>2020</b> , 5, 492-494	0.5	7
79	Complete mitochondrial genome sequence of a xerophilic fungus,. <i>Mitochondrial DNA Part B: Resources</i> , <b>2019</b> , 4, 2422-2423	0.5	7
78	The complete mitochondrial genome of (Townsend, 1909) (Diptera:Tachinidae). <i>Mitochondrial DNA Part B: Resources</i> , <b>2019</b> , 4, 2244-2245	0.5	7
77	The complete chloroplast genome of <i>Nakai</i> from Dokdo Island in Korea (Campanulaceae). <i>Mitochondrial DNA Part B: Resources</i> , <b>2021</b> , 6, 135-137	0.5	7
76	The complete mitochondrial genome of the subterranean termite, <i>Morimoto</i> , 1968 (Isoptera: Rhinotermitidae). <i>Mitochondrial DNA Part B: Resources</i> , <b>2017</b> , 2, 178-179	0.5	6
75	The complete mitochondrial genome of (Walker, 1851) (Hemiptera: Ricaniidae): investigation of intraspecific variations on mitochondrial genome. <i>Mitochondrial DNA Part B: Resources</i> , <b>2020</b> , 5, 3814-3816	0.5	6
74	The complete mitochondrial genome of (Mayr, 1862) (Hymenoptera:Formicidae). <i>Mitochondrial DNA Part B: Resources</i> , <b>2019</b> , 5, 147-149	0.5	6
73	The complete chloroplast genome of (Thunb.) <i>Kanitz</i> (Staphyleaceae) isolated in Korea. <i>Mitochondrial DNA Part B: Resources</i> , <b>2020</b> , 5, 3769-3771	0.5	6
72	Phylogenetic position of <i>Daphne genkwa</i> (Thymelaeaceae) inferred from complete chloroplast data. <i>Korean Journal of Plant Taxonomy</i> , <b>2021</b> , 51, 171-175	0.5	6
71	The complete mitochondrial genome of <i>Arabidopsis thaliana</i> (Brassicaceae) isolated in Korea. <i>Korean Journal of Plant Taxonomy</i> , <b>2021</b> , 51, 176-180	0.5	6
70	A Comparative Analyses of the Complete Mitochondrial Genomes of Fungal Endosymbionts in , White-Backed Planthoppers. <i>International Journal of Genomics</i> , <b>2021</b> , 2021, 6652508	2.5	6
69	The complete mitochondrial genome of <i>Glover</i> , 1877 (Hemiptera: Aphididae) isolated from in Korea. <i>Mitochondrial DNA Part B: Resources</i> , <b>2021</b> , 6, 62-65	0.5	6

68	The complete mitochondrial genome of Panzer, 1797 (Coleoptera: Tenebrionidae). <i>Mitochondrial DNA Part B: Resources</i> , <b>2020</b> , 5, 2291-2293	0.5	5
67	The complete chloroplast genome of Prince Ginseng, (Miq.) Pax (Caryophyllaceae). <i>Mitochondrial DNA Part B: Resources</i> , <b>2019</b> , 4, 2251-2253	0.5	5
66	The complete mitochondrial genome of (Baker, 1919) (Hemiptera: Aphididae). <i>Mitochondrial DNA Part B: Resources</i> , <b>2019</b> , 4, 3037-3039	0.5	5
65	Complete mitochondrial genome sequence of lettuce pathogenic fungus, f. sp. 09-002. <i>Mitochondrial DNA Part B: Resources</i> , <b>2019</b> , 4, 3434-3436	0.5	5
64	The complete mitochondrial genome of the subterranean termite, Takematsu, 1999 (Isoptera: Rhinotermitidae). <i>Mitochondrial DNA Part B: Resources</i> , <b>2017</b> , 2, 508-509	0.5	5
63	Functional genomics in the rice blast fungus to unravel the fungal pathogenicity. <i>Journal of Zhejiang University: Science B</i> , <b>2008</b> , 9, 747-52	4.5	5
62	Eukaryotic DNAJ/K Database: A Comprehensive Phylogenomic Analysis Platform for the DNAJ/K Family. <i>Genomics and Informatics</i> , <b>2013</b> , 11, 52-4	1.9	5
61	New lung cancer panel for high-throughput targeted resequencing. <i>Genomics and Informatics</i> , <b>2014</b> , 12, 50-7	1.9	5
60	The complete chloroplast genome of Steph., 1897 (Scapaniaceae, Jungermanniales). <i>Mitochondrial DNA Part B: Resources</i> , <b>2020</b> , 5, 2890-2892	0.5	5
59	Complete Genome Sequence of <i>Lentilactobacillus parabuchneri</i> Strain KEM. <i>Microbiology Resource Announcements</i> , <b>2021</b> , 10,	1.3	5
58	The complete chloroplast genome of a new candidate cultivar, Dae Ryun, of Nakai (Oleaceae). <i>Mitochondrial DNA Part B: Resources</i> , <b>2019</b> , 4, 3713-3715	0.5	5
57	The complete chloroplast genome of a new candidate cultivar, Sang Jae, of Nakai (Oleaceae): initial step of intraspecies variations atlas. <i>Mitochondrial DNA Part B: Resources</i> , <b>2019</b> , 4, 3716-3718	0.5	5
56	The complete mitochondrial genome of (Sulzer, 1776; Hemiptera: Aphididae) isolated in Korea. <i>Mitochondrial DNA Part B: Resources</i> , <b>2021</b> , 6, 10-12	0.5	5
55	The complete chloroplast genome of (L.) Merr. isolated in Korea (Poaceae): investigation of intraspecific variations on chloroplast genomes. <i>Mitochondrial DNA Part B: Resources</i> , <b>2021</b> , 6, 572-574	0.5	5
54	Complete chloroplast genome sequence of the S. Lee, K. Heo & S. C. Kim (Caryophyllaceae). <i>Mitochondrial DNA Part B: Resources</i> , <b>2018</b> , 3, 1296-1297	0.5	5
53	The complete chloroplast genome of coffee tree, L. Blue Mountain (Rubiaceae). <i>Mitochondrial DNA Part B: Resources</i> , <b>2019</b> , 4, 2436-2437	0.5	4
52	The complete chloroplast genome of (brid.) C. Mü. (Leucobryaceae, Bryophyta). <i>Mitochondrial DNA Part B: Resources</i> , <b>2019</b> , 4, 2962-2963	0.5	4
51	The complete mitochondrial genome of (Smith, 1874) (Hymenoptera: Formicidae). <i>Mitochondrial DNA Part B: Resources</i> , <b>2020</b> , 5, 420-421	0.5	4



50	The complete mitochondrial genome of (Miyata & Kishida, 1990) (Lepidoptera: Bombycidae). <i>Mitochondrial DNA Part B: Resources</i> , <b>2019</b> , 5, 355-357	0.5	4
49	The complete chloroplast genome of Bornm. (Rosaceae). <i>Mitochondrial DNA Part B: Resources</i> , <b>2019</b> , 4, 2420-2421	0.5	4
48	The complete chloroplast genome of tulip tree, L. (Magnoliaceae): investigation of intra-species chloroplast variations. <i>Mitochondrial DNA Part B: Resources</i> , <b>2019</b> , 4, 2523-2524	0.5	4
47	The complete chloroplast genome sequence of rose-gold pussy willow, Miq. (Salicaceae). <i>Mitochondrial DNA Part B: Resources</i> , <b>2019</b> , 4, 2118-2120	0.5	4
46	Complete mitochondrial genome sequence of lettuce pathogenic fungus, f. sp. 16-086. <i>Mitochondrial DNA Part B: Resources</i> , <b>2019</b> , 4, 3227-3228	0.5	4
45	Complete Genome Sequence of a Endosymbiont of. <i>Microbiology Resource Announcements</i> , <b>2021</b> , 10,	1.3	4
44	The complete chloroplast genome of f. Nakai (Oleaceae) from the Chungbuk Province, Korea. <i>Mitochondrial DNA Part B: Resources</i> , <b>2021</b> , 6, 1754-1756	0.5	4
43	Complete Genome Sequence of the Endosymbiotic Bacterium " <i>Riesia pediculicola</i> ". <i>Microbiology Resource Announcements</i> , <b>2021</b> , 10,	1.3	4
42	The complete chloroplast genome of <i>Zoysia japonica</i> Steud. isolated in Korea (Poaceae): investigation of potential molecular markers on <i>Z. japonica</i> chloroplast genomes. <i>Plant Biotechnology Reports</i> , <b>2021</b> , 15, 707-715	2.5	4
41	The complete chloroplast genome of coffee tree, L. <i>Rypica</i> (Rubiaceae). <i>Mitochondrial DNA Part B: Resources</i> , <b>2019</b> , 4, 2240-2241	0.5	3
40	The complete chloroplast genome of Korean endemic species, (H.L. & vaniot) Nakai (Asteraceae). <i>Mitochondrial DNA Part B: Resources</i> , <b>2019</b> , 4, 2351-2352	0.5	3
39	The complete mitochondrial genome of (Marsham, 1802) (Coleoptera: Curculionidae). <i>Mitochondrial DNA Part B: Resources</i> , <b>2019</b> , 4, 3096-3098	0.5	3
38	Genome-wide analysis of T-DNA integration into the chromosomes of <i>Magnaporthe oryzae</i> . <i>Molecular Microbiology</i> , <b>2007</b> , 66, 826-826	4.1	3
37	A New Mitochondrial Genome of (Horv�th) (Hemiptera: Delphacidae) and Mitogenome-Wide Investigation on Polymorphisms.. <i>Insects</i> , <b>2021</b> , 12,	2.8	3
36	The complete mitochondrial genome of (Erxleben, 1777), as a model species of Chronic Wasting Disease (CWD). <i>Mitochondrial DNA Part B: Resources</i> , <b>2020</b> , 5, 2621-2623	0.5	3
35	Complete mitochondrial genome sequence of Afla-Guard, commercially available non-toxicogenic. <i>Mitochondrial DNA Part B: Resources</i> , <b>2020</b> , 5, 3590-3592	0.5	3
34	The complete mitochondrial genome of (Mitt.) Steph. (Wiesnerellaceae, Marchantiophyta): large number of intraspecific variations on mitochondrial genomes of. <i>Mitochondrial DNA Part B: Resources</i> , <b>2020</b> , 5, 3351-3353	0.5	3
33	The second complete mitochondrial genome of Panzer, 1797 (Coleoptera: Tenebrionidae): investigation of intraspecific variations on mitochondrial genome. <i>Mitochondrial DNA Part B: Resources</i> , <b>2020</b> , 5, 2979-2981	0.5	3

32	Complete mitochondrial genome of the H3 haplotype Argentine ant (Mayr, 1868) (Formicidae; Hymenoptera). <i>Mitochondrial DNA Part B: Resources</i> , <b>2021</b> , 6, 786-788	0.5	3
31	The complete chloroplast genome sequence of new species candidate of Willd. in Korea (Plantaginaceae). <i>Mitochondrial DNA Part B: Resources</i> , <b>2021</b> , 6, 1961-1963	0.5	3
30	The complete chloroplast genome of Herbich (Asteraceae) isolated in Korea. <i>Mitochondrial DNA Part B: Resources</i> , <b>2021</b> , 6, 1956-1958	0.5	3
29	Investigation of Nucleotide Diversity Based on 17 Sea Cucumber Mitochondrial Genomes and Assessment of Sea Cucumber Mitochondrial Gene Markers <b>2021</b> , 2,		3
28	Complete mitochondrial genome of the acrobat ant Santschi, 1930 (Formicidae; Hymenoptera). <i>Mitochondrial DNA Part B: Resources</i> , <b>2021</b> , 6, 593-595	0.5	3
27	Complete mitochondrial genome of the jet ant Wheeler, W.M., 1910 (Formicidae; Hymenoptera). <i>Mitochondrial DNA Part B: Resources</i> , <b>2021</b> , 6, 505-507	0.5	3
26	The complete chloroplast genome of (Makino) Hatus (Fagaceae). <i>Mitochondrial DNA Part B: Resources</i> , <b>2021</b> , 6, 2743-2745	0.5	3
25	The first complete mitogenome of (Merriam, 1905). <i>Mitochondrial DNA Part B: Resources</i> , <b>2020</b> , 5, 2294-2296		2
24	The chloroplast genome sequence of DC. (Magnoliaceae). <i>Mitochondrial DNA Part B: Resources</i> , <b>2018</b> , 3, 342-343	0.5	2
23	The complete mitochondrial genome of Kubo, 1938 (Decapoda: Atyidae). <i>Mitochondrial DNA Part B: Resources</i> , <b>2019</b> , 4, 2332-2334	0.5	2
22	The complete chloroplast genome of Griff. (Fissidentaceae, Bryophyta). <i>Mitochondrial DNA Part B: Resources</i> , <b>2019</b> , 4, 2225-2226	0.5	2
21	Taxonomic studies of the tribe Potentilleae (Rosaceae) in Korea. <i>Korean Journal of Plant Taxonomy</i> , <b>2019</b> , 49, 28-69	0.5	2
20	Assaying Mitochondrial COI Sequences and Their Molecular Studies in Hexapoda, PART I: From 2000 to 2009. <i>Korean Journal of Applied Entomology</i> , <b>2013</b> , 52, 395-402		2
19	The complete chloroplast genome of <i>Zoysia macrostachya</i> (Poaceae): Insights into intraspecific variations and species delimitation of the <i>Zoysia</i> species. <i>Korean Journal of Plant Taxonomy</i> , <b>2021</b> , 51, 326-331	0.5	2
18	The complete chloroplast genome of <i>Limonium tetragonum</i> (Plumbaginaceae) isolated in Korea. <i>Korean Journal of Plant Taxonomy</i> , <b>2021</b> , 51, 337-344	0.5	2
17	The complete mitochondrial genome of the millipede Verhoeff, 1937 collected in limestone cave of Korea (Polydesmidae: Polydesmida). <i>Mitochondrial DNA Part B: Resources</i> , <b>2020</b> , 5, 3845-3847	0.5	2
16	The complete chloroplast genome of (Lindb.) Konstant. & Vilnet (Scapaniaceae, Jungermanniales). <i>Mitochondrial DNA Part B: Resources</i> , <b>2020</b> , 5, 3698-3700	0.5	2
15	First Record of the Complete Mitochondrial Genome of a Saprotrophic and Opportunistic Human Pathogenic Fungus,. <i>Mycobiology</i> , <b>2020</b> , 48, 528-531	1.7	2



14	The complete chloroplast genome of (Mitt.) Steph. (Wiesnerellaceae, Marchantiophyta). <i>Mitochondrial DNA Part B: Resources</i> , <b>2020</b> , 5, 3124-3126	0.5	2
13	The complete mitochondrial genome of (Lindb.) Konstant. ´et. Vilnet (Scapaniaceae, Jungermanniales). <i>Mitochondrial DNA Part B: Resources</i> , <b>2021</b> , 6, 789-791	0.5	2
12	The complete mitochondrial genome of Steph., 1897 (Scapaniaceae, Jungermanniales). <i>Mitochondrial DNA Part B: Resources</i> , <b>2021</b> , 6, 686-688	0.5	2
11	The complete mitochondrial genome of the far Eastern myotis: Thomas, 1906 in mainland of Korea (Chiroptera, Vespertilionidae). <i>Mitochondrial DNA Part B: Resources</i> , <b>2021</b> , 6, 615-616	0.5	2
10	Genome-wide comparative analyses of GATA transcription factors among seven <i>Populus</i> genomes. <i>Scientific Reports</i> , <b>2021</b> , 11, 16578	4.9	2
9	Comprehensive Analysis of the Effect of Probiotic Intake by the Mother on Human Breast Milk and Infant Fecal Microbiota. <i>Journal of Korean Medical Science</i> , <b>2021</b> , 36, e58	4.7	2
8	Complete mitochondrial genome of the gate-keeper ant (Wheeler, W.M., 1928) (Formicidae: Hymenoptera). <i>Mitochondrial DNA Part B: Resources</i> , <b>2021</b> , 6, 86-88	0.5	2
7	The complete chloroplast genome of <i>Diarthron linifolium</i> (Thymelaeaceae), a species found on a limestone outcrop in eastern Asia. <i>Korean Journal of Plant Taxonomy</i> , <b>2021</b> , 51, 345-352	0.5	2
6	The investigation of intraspecific characteristics and comparative analyses of the complete mitochondrial genome of (Linnaeus, 1758) (Coleoptera: Ptinidae) assembled from public NGS raw reads of the black truffle, .. <i>Science Progress</i> , <b>2022</b> , 105, 368504211072355	1.1	1
5	Complete mitochondrial genome sequence of SRRC1009: insight of intraspecific variations on mitochondrial genomes. <i>Mitochondrial DNA Part B: Resources</i> , <b>2020</b> , 5, 3585-3587	0.5	1
4	The complete chloroplast genome of var. (Rosaceae). <i>Mitochondrial DNA Part B: Resources</i> , <b>2021</b> , 6, 1256-1258	1.2	1
3	The complete chloroplast genome of <i>Glycyrrhiza uralensis</i> Fisch. isolated in Korea (Fabaceae). <i>Korean Journal of Plant Taxonomy</i> , <b>2021</b> , 51, 353-362	0.5	1
2	The complete mitochondrial genome of (Thomas, 1878) (Hemiptera: Aphididae).. <i>Mitochondrial DNA Part B: Resources</i> , <b>2022</b> , 7, 84-86	0.5	1
1	The complete chloroplast genome of C. A. Mey. ex Bong. (Amaranthaceae).. <i>Mitochondrial DNA Part B: Resources</i> , <b>2022</b> , 7, 541-543	0.5	