

# Jongsun Park

## List of Publications by Year in descending order

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206  
papers

7,814  
citations

147566

31  
h-index

60497

81  
g-index

206  
all docs

206  
docs citations

206  
times ranked

8968  
citing authors

#	ARTICLE	IF	CITATIONS
1	Comparative genomics reveals mobile pathogenicity chromosomes in <i>Fusarium</i> . <i>Nature</i> , 2010, 464, 367-373.	13.7	1,442
2	The <i>Amborella</i> Genome and the Evolution of Flowering Plants. <i>Science</i> , 2013, 342, 1241089.	6.0	743
3	The Plant Cell Wallâ€™s Decomposing Machinery Underlies the Functional Diversity of Forest Fungi. <i>Science</i> , 2011, 333, 762-765.	6.0	512
4	Comparative Genomics Yields Insights into Niche Adaptation of Plant Vascular Wilt Pathogens. <i>PLoS Pathogens</i> , 2011, 7, e1002137.	2.1	477
5	Internet-Accessible DNA Sequence Database for Identifying <i>Fusaria</i> from Human and Animal Infections. <i>Journal of Clinical Microbiology</i> , 2010, 48, 3708-3718.	1.8	446
6	Insight into trade-off between wood decay and parasitism from the genome of a fungal forest pathogen. <i>New Phytologist</i> , 2012, 194, 1001-1013.	3.5	210
7	Genome-wide functional analysis of pathogenicity genes in the rice blast fungus. <i>Nature Genetics</i> , 2007, 39, 561-565.	9.4	205
8	Homeobox Transcription Factors Are Required for Conidiation and Appressorium Development in the Rice Blast Fungus <i>Magnaporthe oryzae</i> . <i>PLoS Genetics</i> , 2009, 5, e1000757.	1.5	195
9	FTFD: an informatics pipeline supporting phylogenomic analysis of fungal transcription factors. <i>Bioinformatics</i> , 2008, 24, 1024-1025.	1.8	162
10	Fungal Secretome Database: Integrated platform for annotation of fungal secretomes. <i>BMC Genomics</i> , 2010, 11, 105.	1.2	160
11	Systematic and searchable classification of cytochrome P450 proteins encoded by fungal and oomycete genomes. <i>BMC Genomics</i> , 2012, 13, 525.	1.2	150
12	Fungal cytochrome P450 database. <i>BMC Genomics</i> , 2008, 9, 402.	1.2	134
13	A Putative MAP Kinase Kinase Kinase, <i>MCK1</i> , Is Required for Cell Wall Integrity and Pathogenicity of the Rice Blast Fungus, <i>Magnaporthe oryzae</i> . <i>Molecular Plant-Microbe Interactions</i> , 2008, 21, 525-534.	1.4	132
14	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> , 2009, 46, 243-254.	0.9	132
15	Taxonomic Status of the <i>Bemisia tabaci</i> Complex (Hemiptera: Aleyrodidae) and Reassessment of the Number of Its Constituent Species. <i>PLoS ONE</i> , 2013, 8, e63817.	1.1	105
16	In-depth insight into in vivo apoplastic secretome of rice- <i>Magnaporthe oryzae</i> interaction. <i>Journal of Proteomics</i> , 2013, 78, 58-71.	1.2	104
17	Genome-wide analysis of Tâ€œDNA integration into the chromosomes of <i>Magnaporthe oryzae</i> . <i>Molecular Microbiology</i> , 2007, 66, 371-382.	1.2	90
18	CFGP: a web-based, comparative fungal genomics platform. <i>Nucleic Acids Research</i> , 2007, 36, D562-D571.	6.5	76

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19	Simple sequence repeats in <i>Neurospora crassa</i> : distribution, polymorphism and evolutionary inference. <i>BMC Genomics</i> , 2008, 9, 31.	1.2	74
20	Whole transcriptome analyses of six thoroughbred horses before and after exercise using RNA-Seq. <i>BMC Genomics</i> , 2012, 13, 473.	1.2	73
21	Complete sequencing and comparative analyses of the pepper ( <i>Capsicum annum</i> L.) plastome revealed high frequency of tandem repeats and large insertion/deletions on pepper plastome. <i>Plant Cell Reports</i> , 2011, 30, 217-229.	2.8	67
22	De novo transcriptome sequencing of <i>Momordica cochinchinensis</i> to identify genes involved in the carotenoid biosynthesis. <i>Plant Molecular Biology</i> , 2012, 79, 413-427.	2.0	66
23	<i>Phytophthora</i> Database: A Forensic Database Supporting the Identification and Monitoring of <i>Phytophthora</i>. <i>Plant Disease</i> , 2008, 92, 966-972.	0.7	64
24	Cyber infrastructure for <i>Fusarium</i> : three integrated platforms supporting strain identification, phylogenetics, comparative genomics and knowledge sharing. <i>Nucleic Acids Research</i> , 2011, 39, D640-D646.	6.5	63
25	Global Expression Profiling of Transcription Factor Genes Provides New Insights into Pathogenicity and Stress Responses in the Rice Blast Fungus. <i>PLoS Pathogens</i> , 2013, 9, e1003350.	2.1	61
26	Combining ChIP-chip and Expression Profiling to Model the MoCRZ1 Mediated Circuit for Ca <sup>2+</sup> /Calcineurin Signaling in the Rice Blast Fungus. <i>PLoS Pathogens</i> , 2010, 6, e1000909.	2.1	57
27	Comprehensive genome- and transcriptome-wide analyses of mutations associated with microsatellite instability in Korean gastric cancers. <i>Genome Research</i> , 2013, 23, 1109-1117.	2.4	56
28	The PEX7-Mediated Peroxisomal Import System Is Required for Fungal Development and Pathogenicity in <i>Magnaporthe oryzae</i> . <i>PLoS ONE</i> , 2011, 6, e28220.	1.1	52
29	Evolution of the large genome in <i>Capsicum annum</i> occurred through accumulation of single-type long terminal repeat retrotransposons and their derivatives. <i>Plant Journal</i> , 2012, 69, 1018-1029.	2.8	51
30	Identification and analysis of in planta expressed genes of <i>Magnaporthe oryzae</i> . <i>BMC Genomics</i> , 2010, 11, 104.	1.2	37
31	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> , 2011, 12, 85.	1.2	34
32	Comparison of Whole Plastome Sequences between Thermogenic Skunk Cabbage <i>Symplocarpus renifolius</i> and Nonthermogenic <i>S. nipponicus</i> (Orontioideae; Araceae) in East Asia. <i>International Journal of Molecular Sciences</i> , 2019, 20, 4678.	1.8	33
33	The complete chloroplast genome of common camellia tree, <i>Camellia japonica</i> L. (Theaceae), adapted to cold environment in Korea. <i>Mitochondrial DNA Part B: Resources</i> , 2019, 4, 1038-1040.	0.2	25
34	Genome-wide comparative analyses of GATA transcription factors among 19 <i>Arabidopsis</i> ecotype genomes: Intraspecific characteristics of GATA transcription factors. <i>PLoS ONE</i> , 2021, 16, e0252181.	1.1	25
35	The Complete Chloroplast Genome of <i>Arabidopsis thaliana</i> Isolated in Korea (Brassicaceae): An Investigation of Intraspecific Variations of the Chloroplast Genome of Korean <i>A. thaliana</i>. <i>International Journal of Genomics</i> , 2020, 2020, 1-18.	0.8	24
36	The second complete chloroplast genome sequence of <i>Pseudostellaria palibiniana</i> (Takeda) Ohwi (Caryophyllaceae): intraspecies variations based on geographical distribution. <i>Mitochondrial DNA Part B: Resources</i> , 2019, 4, 1310-1311.	0.2	22

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37	The complete mitochondrial genome of the fall armyworm, <i>Spodoptera frugiperda</i> Smith, 1797 (Lepidoptera; Noctuidae), firstly collected in Korea. Mitochondrial DNA Part B: Resources, 2019, 4, 3918-3920.	0.2	22
38	A complete chloroplast genome sequence of <i>Gastrodia elata</i> (Orchidaceae) represents high sequence variation in the species. Mitochondrial DNA Part B: Resources, 2020, 5, 517-519.	0.2	21
39	Peeling Back the Evolutionary Layers of Molecular Mechanisms Responsive to Exercise-Stress in the Skeletal Muscle of the Racing Horse. DNA Research, 2013, 20, 287-298.	1.5	20
40	The complete chloroplast genome of Korean <i>Pyrus ussuriensis</i> Maxim. (Rosaceae): providing genetic background of two types of <i>P. ussuriensis</i> . Mitochondrial DNA Part B: Resources, 2019, 4, 2424-2425.	0.2	20
41	The complete mitochondrial genome of <i>Laodelphax striatellus</i> (Fallén, 1826) (Hemiptera: Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 Resources, 2019, 4, 2229-2230.	0.2	20
42	The complete chloroplast genome of Korean <i>Marchantia polymorpha</i> subsp. <i>ruderalis</i> Bischl. & Boisselier: low genetic diversity between Korea and Japan. Mitochondrial DNA Part B: Resources, 2019, 4, 959-960.	0.2	20
43	SysPIMP: the web-based systematical platform for identifying human disease-related mutated sequences from mass spectrometry. Nucleic Acids Research, 2009, 37, D913-D920.	6.5	19
44	The complete chloroplast genome of Korean mock strawberry, <i>Duchesnea chrysantha</i> (Zoll.) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50	0.2	19
45	The Comparative Analyses of Six Complete Chloroplast Genomes of Morphologically Diverse <i>Chenopodium album</i> L. (Amaranthaceae) Collected in Korea. International Journal of Genomics, 2021, 2021, 1-15.	0.8	19
46	The complete chloroplast genome sequence of <i>Viburnum erosum</i> (Adoxaceae). Mitochondrial DNA Part B: Resources, 2019, 4, 3278-3279.	0.2	18
47	The complete chloroplast genome sequence of a morphotype of <i>Goodyera schlechtendaliana</i> (Orchidaceae) with the column appendages. Mitochondrial DNA Part B: Resources, 2019, 4, 626-627.	0.2	18
48	The complete chloroplast genome of ornamental coffee tree, <i>Coffea arabica</i> L. (Rubiaceae). Mitochondrial DNA Part B: Resources, 2019, 4, 1059-1060.	0.2	18
49	The complete mitochondrial genome of new species candidate of <i>Rosa rugosa</i> (Rosaceae). Mitochondrial DNA Part B: Resources, 2020, 5, 3435-3437.	0.2	18
50	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
51	SNUGB: a versatile genome browser supporting comparative and functional fungal genomics. BMC Genomics, 2008, 9, 586.	1.2	17
52	IMGD: an integrated platform supporting comparative genomics and phylogenetics of insect mitochondrial genomes. BMC Genomics, 2009, 10, 148.	1.2	17
53	The complete mitochondrial genome of <i>Nilaparvata lugens</i> (Stål, 1854) captured in Korea (Hemiptera: Delphacidae). Mitochondrial DNA Part B: Resources, 2019, 4, 1674-1676.	0.2	17
54	The complete mitochondrial genome of <i>Laodelphax striatellus</i> (Fallén, 1826) (Hemiptera: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 2019, 4, 2242-2243.	0.2	17

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55	The complete chloroplast genome sequence of <i>Goodyera schlechtendaliana</i> in Korea (Orchidaceae). Mitochondrial DNA Part B: Resources, 2019, 4, 2692-2693.	0.2	17
56	The complete mitochondrial genome sequence of Korean <i>Chilo suppressalis</i> (Walker, 1863) (Lepidoptera: Crambidae). Mitochondrial DNA Part B: Resources, 2019, 4, 850-851.	0.2	17
57	The complete mitochondrial genome of Korean <i>Marchantia polymorpha</i> subsp. <i>ruderalis</i> Bischl. & Boisselier: inverted repeats on mitochondrial genome between Korean and Japanese isolates. Mitochondrial DNA Part B: Resources, 2019, 4, 769-770.	0.2	17
58	A second complete chloroplast genome sequence of <i>Fagus multinervis</i> Nakai (Fagaceae): intraspecific variations on chloroplast genome. Mitochondrial DNA Part B: Resources, 2020, 5, 1868-1869.	0.2	17
59	The complete mitochondrial genome of <i>Nilaparvata lugens</i> (Stål, 1854) captured in China (Hemiptera: Delphacidae): investigation of intraspecies variations between countries. Mitochondrial DNA Part B: Resources, 2019, 4, 1677-1678.	0.2	16
60	The complete chloroplast genome of common camellia tree in Jeju island, Korea, <i>Camellia japonica</i> L. (Theaceae): intraspecies variations on common camellia chloroplast genomes. Mitochondrial DNA Part B: Resources, 2019, 4, 1292-1293.	0.2	16
61	The second complete chloroplast genome sequence of the <i>Viburnum erosum</i> (Adoxaceae) showed a low level of intra-species variations. Mitochondrial DNA Part B: Resources, 2020, 5, 271-272.	0.2	16
62	The complete chloroplast genome sequence of male individual of Korean endemic willow, <i>Salix koriyanagi</i> Kimura ex Goerz (Salicaceae). Mitochondrial DNA Part B: Resources, 2019, 4, 1619-1621.	0.2	15
63	The complete chloroplast genome of <i>Aconitum coreanum</i> (H. Lév.) Rapaics (Ranunculaceae). Mitochondrial DNA Part B: Resources, 2019, 4, 3404-3406.	0.2	15
64	The complete chloroplast genome of <i>Abeliophyllum distichum</i> Nakai (Oleaceae), cultivar Ok Hwang 1ho: insights of cultivar specific variations of <i>A. distichum</i> . Mitochondrial DNA Part B: Resources, 2019, 4, 1640-1642.	0.2	15
65	The complete mitochondrial genome of <i>Aphis gossypii</i> Glover, 1877 (Hemiptera: Aphididae) collected in Korean peninsula. Mitochondrial DNA Part B: Resources, 2019, 4, 3007-3009.	0.2	15
66	The second complete chloroplast genome of <i>Dysphania pumilio</i> (R.Br.) mosyakin & clemants (Amaranthaceae): intraspecies variation of invasive weeds. Mitochondrial DNA Part B: Resources, 2019, 4, 1428-1429.	0.2	14
67	The complete mitochondrial genome of tulip tree, <i>Liriodendron tulipifera</i> L. (Magnoliaceae): intra-species variations on mitochondrial genome. Mitochondrial DNA Part B: Resources, 2019, 4, 1308-1309.	0.2	14
68	Comprehensive Analyses of the Complete Mitochondrial Genome of <i>Figulus binodulus</i> (Coleoptera: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5	0.6	14
69	Comparative transcriptome analysis of three color variants of the sea cucumber <i>Apostichopus japonicus</i> . Marine Genomics, 2016, 28, 21-24.	0.4	13
70	The complete chloroplast genome of <i>Artemisia fukudo</i> Makino (Asteraceae): providing insight of intraspecies variations. Mitochondrial DNA Part B: Resources, 2019, 4, 1510-1512.	0.2	13
71	The complete chloroplast and mitochondrial genomes of Hyunsasi tree, <i>Populus alba</i> x <i>Populus glandulosa</i> (Salicaceae). Mitochondrial DNA Part B: Resources, 2019, 4, 2521-2522.	0.2	13
72	The complete chloroplast genome of candidate new species from <i>Rosa rugosa</i> in Korea (Rosaceae). Mitochondrial DNA Part B: Resources, 2019, 4, 2433-2435.	0.2	13

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73	The complete chloroplast genome of aniseed tree, <i>Illicium anisatum</i> L. (Schisandraceae). Mitochondrial DNA Part B: Resources, 2019, 4, 1023-1024.	0.2	13
74	The complete chloroplast genome of <i>Agrimonia pilosa</i> Ledeb. isolated in Korea (Rosaceae): investigation of intraspecific variations on its chloroplast genomes. Mitochondrial DNA Part B: Resources, 2020, 5, 2264-2266.	0.2	13
75	The complete chloroplast genome of high production individual tree of <i>Coffea arabica</i> L. (Rubiaceae). Mitochondrial DNA Part B: Resources, 2019, 4, 1541-1542.	0.2	12
76	The complete chloroplast genome sequence of <i>Dumortiera hirsuta</i> (Sw.) Nees (Marchantiophyta, Dumortieraceae). Mitochondrial DNA Part B: Resources, 2019, 4, 318-319.	0.2	12
77	The complete mitochondrial genome of <i>Aphaenogaster famelica</i> (Smith, 1874) (Hymenoptera: Tj ETQq1 1 0.784314 rgBT / Overlock 10 Tf 50 222 Td (	0.2	12
78	The complete chloroplast genome of horned holly, <i>Ilex cornuta</i> Lindl. & Paxton (Aquifoliaceae). Mitochondrial DNA Part B: Resources, 2019, 4, 1275-1276.	0.2	11
79	The complete chloroplast genome of cold hardiness individual of <i>Coffea arabica</i> L. (Rubiaceae). Mitochondrial DNA Part B: Resources, 2019, 4, 1083-1084.	0.2	11
80	The complete mitochondrial genome of <i>Aphis gossypii</i> Glover, 1877 (Hemiptera: Aphididae) isolated from <i>Plantago asiatica</i> in Korea. Mitochondrial DNA Part B: Resources, 2020, 5, 2878-2880.	0.2	11
81	Phylogenetic position of <i>Daphne genkwa</i> (Thymelaeaceae) inferred from complete chloroplast data. Korean Journal of Plant Taxonomy, 2021, 51, 171-175.	0.3	11
82	Genome-wide comparative analyses of GATA transcription factors among seven <i>Populus</i> genomes. Scientific Reports, 2021, 11, 16578.	1.6	11
83	Complete mitochondrial genome sequence of a xerophilic fungus, <i>Aspergillus pseudoglaucus</i> . Mitochondrial DNA Part B: Resources, 2019, 4, 2422-2423.	0.2	10
84	The complete mitochondrial genome of <i>Exorista japonica</i> (Townsend, 1909) (Diptera:Tachinidae). Mitochondrial DNA Part B: Resources, 2019, 4, 2244-2245.	0.2	10
85	The complete chloroplast genome of coffee tree, <i>Coffea arabica</i> L. "Blue Mountain"™ (Rubiaceae). Mitochondrial DNA Part B: Resources, 2019, 4, 2436-2437.	0.2	10
86	The complete mitochondrial genome of <i>Dumortiera hirsuta</i> (Sw.) Nees (Dumortieraceae,) Tj ETQq0 0 0 rgBT / Overlock 10 Tf 50 222 Td (	0.2	10
87	The complete chloroplast genome sequence of <i>Hibiscus syriacus</i> L. "Mamonde"™ (Malvaceae). Mitochondrial DNA Part B: Resources, 2019, 4, 558-559.	0.2	10
88	The complete chloroplast genome of a new candidate cultivar, Sang Jae, of <i>Abeliophyllum distichum</i> Nakai (Oleaceae): initial step of <i>A. distichum</i> intraspecies variations atlas. Mitochondrial DNA Part B: Resources, 2019, 4, 3716-3718.	0.2	10
89	The complete mitochondrial genome of <i>Nilaparvata lugens</i> (Stål, 1854) captured in Guangxi province, China (Hemiptera: Delphacidae): identification of the origin of <i>N. lugens</i> migrated to Korea. Mitochondrial DNA Part B: Resources, 2020, 5, 1960-1961.	0.2	10
90	The comparison of the complete chloroplast genome of <i>Suaeda japonica</i> Makino presenting different external morphology (Amaranthaceae). Mitochondrial DNA Part B: Resources, 2020, 5, 1616-1618.	0.2	10

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91	A Comparative Analyses of the Complete Mitochondrial Genomes of Fungal Endosymbionts in <i>Sogatella furcifera</i> , White-Backed Planthoppers. <i>International Journal of Genomics</i> , 2021, 2021, 1-20.	0.8	10
92	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> , 2021, 15, 707-715.	0.9	10
93	Complete chloroplast genome sequence of the <i>Salix koriyanagi</i> Kimura ex Goerz (Salicaceae). <i>Mitochondrial DNA Part B: Resources</i> , 2019, 4, 549-550.	0.2	10
94	Eukaryotic DNA/K Database: A Comprehensive Phylogenomic Analysis Platform for the DNAJ/K Family. <i>Genomics and Informatics</i> , 2013, 11, 52.	0.4	10
95	Complete chloroplast genome sequence of the <i>Pseudostellaria longipedicellata</i> S. Lee, K. Heo & S. C. Kim (Caryophyllaceae). <i>Mitochondrial DNA Part B: Resources</i> , 2018, 3, 1296-1297.	0.2	9
96	The complete chloroplast genome of tulip tree, <i>Liriodendron tulifipera</i> L. (Magnoliaceae): investigation of intra-species chloroplast variations. <i>Mitochondrial DNA Part B: Resources</i> , 2019, 4, 2523-2524.	0.2	9
97	The complete chloroplast genome of mock strawberry, <i>Duchesnea indica</i> (Andrews) Th. Wolf (Rosaceae). <i>Mitochondrial DNA Part B: Resources</i> , 2019, 4, 560-562.	0.2	9
98	The second complete chloroplast genome sequence of <i>Nymphaea alba</i> L. (Nymphaeaceae) to investigate inner-species variations. <i>Mitochondrial DNA Part B: Resources</i> , 2019, 4, 1014-1015.	0.2	9
99	The complete chloroplast genome sequence of traditional medical herb, <i>Plantago depressa</i> Willd. (Plantaginaceae). <i>Mitochondrial DNA Part B: Resources</i> , 2019, 4, 437-438.	0.2	9
100	The complete mitochondrial genome of <i>Riccia fluitans</i> L. (Ricciaceae, Marchantiophyta): investigation of intraspecific variations on mitochondrial genomes of <i>R. fluitans</i> . <i>Mitochondrial DNA Part B: Resources</i> , 2020, 5, 1220-1222.	0.2	9
101	The complete chloroplast genome of <i>Selaginella tamariscina</i> (Beauv.) Spring (Selaginellaceae) isolated in Korea. <i>Mitochondrial DNA Part B: Resources</i> , 2020, 5, 1654-1656.	0.2	9
102	The complete mitochondrial genome of <i>Aiolocaria hexaspilota</i> (Hope, 1831) (Coleoptera:Coccinellidae). <i>Mitochondrial DNA Part B: Resources</i> , 2019, 4, 1472-1474.	0.2	8
103	The complete chloroplast genome of Prince Ginseng, <i>Pseudostellaria heterophylla</i> (Miq.) Pax (Caryophyllaceae). <i>Mitochondrial DNA Part B: Resources</i> , 2019, 4, 2251-2253.	0.2	8
104	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
105	The complete chloroplast genome of coffee tree, <i>Coffea arabica</i> L. "Typica" (Rubiaceae). <i>Mitochondrial DNA Part B: Resources</i> , 2019, 4, 2240-2241.	0.2	8
106	Complete mitochondrial genome sequence of an aflatoxin B and G producing fungus, <i>Aspergillus parasiticus</i> . <i>Mitochondrial DNA Part B: Resources</i> , 2019, 4, 947-948.	0.2	8
107	The complete chloroplast genome sequence of <i>Dysphania pumilio</i> (R.Br.) Mosyakin & Clemants (Amaranthaceae). <i>Mitochondrial DNA Part B: Resources</i> , 2019, 4, 403-404.	0.2	8
108	The complete mitochondrial genome of <i>Cryptopone sauteri</i> Wheeler, W.M., 1906 (Hymenoptera: Tj ETQq0 0,0,rgBT /Qverlock 10	0.2	8

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109	Complete Genome Sequence of the <i>Blochmannia</i> Endosymbiont of <i>Camponotus nipponensis</i> . <i>Microbiology Resource Announcements</i> , 2020, 9, .	0.3	8
110	The complete mitochondrial genome of <i>Ricania speculum</i> (Walker, 1851) (Hemiptera: Ricaniidae): investigation of intraspecific variations on mitochondrial genome. <i>Mitochondrial DNA Part B: Resources</i> , 2020, 5, 3796-3798.	0.2	8
111	The complete mitochondrial genome of <i>Ochetellus glaber</i> (Mayr, 1862) (Hymenoptera:Formicidae). <i>Mitochondrial DNA Part B: Resources</i> , 2020, 5, 147-149.	0.2	8
112	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
113	The complete mitochondrial genome of the subterranean termite, <i>Reticulitermes speratus kyushuensis</i> Morimoto, 1968 (Isoptera: Rhinotermitidae). <i>Mitochondrial DNA Part B: Resources</i> , 2017, 2, 178-179.	0.2	7
114	The complete chloroplast genome of <i>Chenopodium ficifolium</i> Sm. (Amaranthaceae). <i>Mitochondrial DNA Part B: Resources</i> , 2019, 4, 872-873.	0.2	7
115	Complete mitochondrial genome sequence of the food fermentation fungus, <i>Aspergillus luchuensis</i> . <i>Mitochondrial DNA Part B: Resources</i> , 2019, 4, 945-946.	0.2	7
116	The complete chloroplast genome of <i>Reboulia hemisphaerica</i> (L.) Raddi (Aytoniaceae,) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 462	0.2	7
117	The complete chloroplast genome of an endangered species in Korea, <i>Halenia corniculata</i> (L.) Cornaz (Gentianaceae). <i>Mitochondrial DNA Part B: Resources</i> , 2019, 4, 1539-1540.	0.2	7
118	The complete chloroplast genome of new variety candidate in Korea, <i>Potentilla freyniana</i> var. <i>chejuensis</i> (Rosoidae). <i>Mitochondrial DNA Part B: Resources</i> , 2019, 4, 1354-1356.	0.2	7
119	The complete chloroplast genome of <i>Riccia fluitans</i> L. (Ricciaceae, Marchantiophyta). <i>Mitochondrial DNA Part B: Resources</i> , 2019, 4, 1895-1896.	0.2	7
120	The complete mitochondrial genome of <i>Paracolopha morrisoni</i> (Baker, 1919) (Hemiptera: Aphididae). <i>Mitochondrial DNA Part B: Resources</i> , 2019, 4, 3037-3039.	0.2	7
121	The complete mitochondrial genome of Siberian odorous ant, <i>Dolichoderus sibiricus</i> Emery, 1889 (Hymenoptera: Formicidae). <i>Mitochondrial DNA Part B: Resources</i> , 2019, 4, 525-526.	0.2	7
122	The complete mitochondrial genome of <i>Camponotus concavus</i> Kim & Kim, 1994 (Hymenoptera:) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 2	0.2	7
123	The complete chloroplast genome of a new candidate cultivar, Dae Ryun, of <i>Abeliophyllum distichum</i> Nakai (Oleaceae). <i>Mitochondrial DNA Part B: Resources</i> , 2019, 4, 3713-3715.	0.2	7
124	The complete chloroplast genome of <i>Euscaphis japonica</i> (Thunb.) Kanitz (Staphyleaceae) isolated in Korea. <i>Mitochondrial DNA Part B: Resources</i> , 2020, 5, 3751-3753.	0.2	7
125	The complete chloroplast genome of <i>Scapania ampliata</i> Steph., 1897 (Scapaniaceae,) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 7	0.2	7
126	The complete mitochondrial genome of fungal endosymbiont, <i>Ophiocordycipitaceae</i> sp., isolated from <i>Ricania speculum</i> (Hemiptera: Ricaniidae). <i>Mitochondrial DNA Part B: Resources</i> , 2020, 5, 1888-1889.	0.2	7



#	ARTICLE	IF	CITATIONS
127	The complete mitochondrial genome of <i>Hipparchia autonoe</i> (Esper, 1783) (Lepidoptera: Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 Part B: Resources, 2020, 5, 1542-1544.	0.2	7
128	The complete mitochondrial genome of <i>Alphitobius diaperinus</i> Panzer, 1797 (Coleoptera: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50	0.2	7
129	The complete mitochondrial genome of <i>Rhopalosiphum nymphaeae</i> (Linnaeus, 1761) (Hemiptera: Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50	0.2	7
130	The complete mitochondrial genome of <i>Aphis gossypii</i> Glover, 1877 (Hemiptera: Aphididae) isolated from <i>Leonurus japonicus</i> in Korea. Mitochondrial DNA Part B: Resources, 2021, 6, 62-65.	0.2	7
131	The complete mitochondrial genome of <i>Arabidopsis thaliana</i> (Brassicaceae) isolated in Korea. Korean Journal of Plant Taxonomy, 2021, 51, 176-180.	0.3	7
132	A New Mitochondrial Genome of <i>Sogatella furcifera</i> (Horváth) (Hemiptera: Delphacidae) and Mitogenome-Wide Investigation on Polymorphisms. Insects, 2021, 12, 1066.	1.0	7
133	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
134	The complete chloroplast genome of <i>Suaeda japonica</i> Makino (Amaranthaceae). Mitochondrial DNA Part B: Resources, 2019, 4, 1505-1507.	0.2	6
135	The complete mitochondrial genome of <i>Micromus angulatus</i> (Stephens, 1836) (Neuroptera: Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50	0.2	6
136	The complete chloroplast genome of <i>Potentilla Freyniana</i> Bornm. (Rosaceae). Mitochondrial DNA Part B: Resources, 2019, 4, 2420-2421.	0.2	6
137	The complete chloroplast genome of <i>Leucobryum juniperoideum</i> (brid.) C. MÅ¼ll. (Leucobryaceae, Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50	0.2	6
138	The complete chloroplast genome sequence of the <i>Pseudostellaria okamotoi</i> Ohwi (Caryophyllaceae). Mitochondrial DNA Part B: Resources, 2019, 4, 174-175.	0.2	6
139	The complete chloroplast genome sequence of <i>Pseudostellaria palibiniana</i> (Takeda) Ohwi (Caryophyllaceae). Mitochondrial DNA Part B: Resources, 2019, 4, 973-974.	0.2	6
140	The complete mitochondrial genome of <i>Rotunda rotundapex</i> (Miyata & Kishida, 1990) (Lepidoptera: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50	0.2	6
141	The complete mitochondrial genome of <i>Myzus persicae</i> (Sulzer, 1776; Hemiptera: Aphididae) isolated in Korea. Mitochondrial DNA Part B: Resources, 2021, 6, 10-12.	0.2	6
142	The complete chloroplast genome of <i>Zoysia matrella</i> (L.) Merr. isolated in Korea (Poaceae): investigation of intraspecific variations on chloroplast genomes. Mitochondrial DNA Part B: Resources, 2021, 6, 572-574.	0.2	6
143	The complete mitochondrial genome of <i>Scapania ampliata</i> Steph., 1897 (Scapaniaceae, Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50	0.2	6
144	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

#	ARTICLE	IF	CITATIONS
145	Complete Genome Sequence of the Endosymbiotic Bacterium <i>Candidatus Riesia pediculicola</i> . Microbiology Resource Announcements, 2021, 10, .	0.3	6
146	Complete Genome Sequence of <i>Lentilactobacillus parabuchneri</i> Strain KEM. Microbiology Resource Announcements, 2021, 10, .	0.3	6
147	The complete chloroplast genome sequence of new species candidate of <i>Plantago depressa</i> Willd. in Korea (Plantaginaceae). Mitochondrial DNA Part B: Resources, 2021, 6, 1961-1963.	0.2	6
148	The complete chloroplast genome of <i>Chrysanthemum zawadskii</i> Herbich (Asteraceae) isolated in Korea. Mitochondrial DNA Part B: Resources, 2021, 6, 1956-1958.	0.2	6
149	The complete chloroplast genome of <i>Castanopsis sieboldii</i> (Makino) Hatus (Fagaceae). Mitochondrial DNA Part B: Resources, 2021, 6, 2743-2745.	0.2	6
150	Comprehensive Analysis of the Effect of Probiotic Intake by the Mother on Human Breast Milk and Infant Fecal Microbiota. Journal of Korean Medical Science, 2021, 36, e58.	1.1	6
151	The complete mitochondrial genome of the millipede <i>Epanerchodus koreanus</i> Verhoeff, 1937 collected in limestone cave of Korea (Polydesmidae: Polydesmida). Mitochondrial DNA Part B: Resources, 2020, 5, 3845-3847.	0.2	6
152	New Lung Cancer Panel for High-Throughput Targeted Resequencing. Genomics and Informatics, 2014, 12, 50.	0.4	6
153	Functional genomics in the rice blast fungus to unravel the fungal pathogenicity. Journal of Zhejiang University: Science B, 2008, 9, 747-752.	1.3	5
154	The complete mitochondrial genome of the subterranean termite, <i>Reticulitermes kanmonensis</i> Takematsu, 1999 (Isoptera: Rhinotermitidae). Mitochondrial DNA Part B: Resources, 2017, 2, 508-509.	0.2	5
155	The chloroplast genome sequence of <i>Magnolia kobus</i> DC. (Magnoliaceae). Mitochondrial DNA Part B: Resources, 2018, 3, 342-343.	0.2	5
156	The complete chloroplast genome sequence of rose-gold pussy willow, <i>Salix gracilistyla</i> Miq. (Salicaceae). Mitochondrial DNA Part B: Resources, 2019, 4, 2118-2120.	0.2	5
157	Complete mitochondrial genome sequence of lettuce pathogenic fungus, <i>Fusarium oxysporum</i> f. sp. <i>lactucae</i> 09-002. Mitochondrial DNA Part B: Resources, 2019, 4, 3434-3436.	0.2	5
158	Complete mitochondrial genome sequence of lettuce pathogenic fungus, <i>Fusarium oxysporum</i> f. sp. <i>lactucae</i> 16-086. Mitochondrial DNA Part B: Resources, 2019, 4, 3227-3228.	0.2	5
159	The complete mitochondrial genome of <i>Ectomomyrmex javanus</i> Mayr, 1867 (Hymenoptera: Formicidae). Mitochondrial DNA Part B: Resources, 2019, 4, 1636-1637.	0.2	5
160	The complete mitochondrial genome sequence of Chinese minnow in Korea, <i>Rhynchocypris oxycephalus</i> (Sauvage and Dabry de Thiersant, 1874). Mitochondrial DNA Part B: Resources, 2019, 4, 662-663.	0.2	5
161	The complete chloroplast genome, <i>Duchesnea chrysantha</i> (Zoll. & Moritzi) Miq. (Rosoideae). Mitochondrial DNA Part B: Resources, 2019, 4, 951-952.	0.2	5
162	The Complete chloroplast genome sequence of the <i>Nymphaea lotus</i> L. (Nymphaeaceae). Mitochondrial DNA Part B: Resources, 2019, 4, 389-390.	0.2	5

#	ARTICLE	IF	CITATIONS
163	The complete chloroplast genome of <i>Pyrus ussuriensis</i> Maxim. (Rosaceae). Mitochondrial DNA Part B: Resources, 2019, 4, 1000-1001.	0.2	5
164	The complete chloroplast genome of <i>Potentilla centigrana</i> Maxim. (Rosaceae). Mitochondrial DNA Part B: Resources, 2019, 4, 688-689.	0.2	5
165	Complete mitochondrial genome sequence of Afla-Guard <sup>®</sup> , commercially available non-toxicogenic <i>Aspergillus flavus</i> . Mitochondrial DNA Part B: Resources, 2020, 5, 3572-3574.	0.2	5
166	The second complete mitochondrial genome of <i>Alphitobius diaperinus</i> Panzer, 1797 (Coleoptera: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50) Mitochondrial DNA Part B: Resources, 2020, 5, 2979-2981.	0.2	5
167	The complete chloroplast genomes of two cold hardness coffee trees, <i>Coffea arabica</i> L. (Rubiaceae). Mitochondrial DNA Part B: Resources, 2020, 5, 1619-1621.	0.2	5
168	The complete mitochondrial genome of <i>Nylanderia flavipes</i> (Smith, 1874) (Hymenoptera: Formicidae). Mitochondrial DNA Part B: Resources, 2020, 5, 420-421.	0.2	5
169	Complete mitochondrial genome of the acrobat ant <i>Crematogaster teranishii</i> Santschi, 1930 (Formicidae; Hymenoptera). Mitochondrial DNA Part B: Resources, 2021, 6, 593-595.	0.2	5
170	Complete mitochondrial genome of the H3 haplotype Argentine ant <i>Linepithema humile</i> (Mayr, 1868) (Formicidae; Hymenoptera). Mitochondrial DNA Part B: Resources, 2021, 6, 786-788.	0.2	5
171	The complete chloroplast genome of <i>Abeliophyllum distichum</i> f. <i>lilacinum</i> Nakai (Oleaceae) from the Chungbuk Province, Korea. Mitochondrial DNA Part B: Resources, 2021, 6, 1754-1756.	0.2	5
172	The complete chloroplast genome of <i>Zoysia macrostachya</i> (Poaceae): Insights into intraspecific variations and species delimitation of the <i>Zoysia</i> species. Korean Journal of Plant Taxonomy, 2021, 51, 326-331.	0.3	5
173	Genome-wide analysis of T-DNA integration into the chromosomes of <i>Magnaporthe oryzae</i> . Molecular Microbiology, 2007, 66, 826-826.	1.2	4
174	The complete chloroplast genome of <i>Potentilla stolonifera</i> var. <i>quelpaertensis</i> Nakai. Mitochondrial DNA Part B: Resources, 2019, 4, 1289-1291.	0.2	4
175	The complete chloroplast genome of Nepal Holly, <i>Ilex integra</i> Thunb. (Aquifoliaceae). Mitochondrial DNA Part B: Resources, 2019, 4, 1257-1258.	0.2	4
176	The complete chloroplast genome of Korean endemic species, <i>Cirsium rhinoceros</i> (H.L. & Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50)	0.2	4
177	The complete chloroplast genome of <i>Fissidens nobilis</i> Griff. (Fissidentaceae, Bryophyta). Mitochondrial DNA Part B: Resources, 2019, 4, 2225-2226.	0.2	4
178	The complete mitochondrial genome of <i>Ceutorhynchus obstructus</i> (Marsham, 1802) (Coleoptera: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50)	0.2	4
179	The complete chloroplast genome sequence of the <i>Nymphaea capensis</i> Thunb. (Nymphaeaceae). Mitochondrial DNA Part B: Resources, 2019, 4, 401-402.	0.2	4
180	The complete mitochondrial genome of <i>Wiesnerella denudata</i> (Mitt.) Steph. (Wiesnerellaceae, Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50) <i>denudata</i> . Mitochondrial DNA Part B: Resources, 2020, 5, 3351-3353.	0.2	4

#	ARTICLE	IF	CITATIONS
181	The first complete mitogenome of <i>Cervus canadensis nannodes</i> (Merriam, 1905). Mitochondrial DNA Part B: Resources, 2020, 5, 2294-2296.	0.2	4
182	Complete mitochondrial genome of the jet ant <i>Lasius spathepus</i> Wheeler, W.M., 1910 (Formicidae). Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5	0.2	4
183	The complete mitochondrial genome of <i>Douinia plicata</i> (Lindb.) Konstant. & Vilnet (Scapaniaceae). Tj ETQq1 1 0.784314 rgBT /Over	0.2	4
184	Complete Genome Sequence of a Blochmannia Endosymbiont of <i>Colobopsis nipponica</i> . Microbiology Resource Announcements, 2021, 10, .	0.3	4
185	Investigation of Nucleotide Diversity Based on 17 Sea Cucumber Mitochondrial Genomes and Assessment of Sea Cucumber Mitochondrial Gene Markers. Advances in Oceanography & Marine Biology, 2021, 2, .	0.1	4
186	The investigation of intraspecific characteristics and comparative analyses of the complete mitochondrial genome of <i>Stegobium paniceum</i> (Linnaeus, 1758) (Coleoptera: Ptinidae) assembled from public NGS raw reads of the black truffle, <i>Tuber melanosporum</i> . Science Progress, 2022, 105, 003685042110723.	1.0	4
187	The complete chloroplast genome of <i>Glycyrrhiza uralensis</i> Fisch. isolated in Korea (Fabaceae). Korean Journal of Plant Taxonomy, 2021, 51, 353-362.	0.3	4
188	The complete chloroplast genome of <i>Potentilla fragarioides</i> var. <i>major</i> Maxim. Mitochondrial DNA Part B: Resources, 2019, 4, 1265-1266.	0.2	3
189	The complete mitochondrial genome of <i>Neocaridina heteropoda koreana</i> Kubo, 1938 (Decapoda). Tj ETQq1 1 0.784314 rgBT /	0.2	3
190	The complete mitochondrial genome of <i>Cervus canadensis</i> (Erxleben, 1777), as a model species of Chronic Wasting Disease (CWD). Mitochondrial DNA Part B: Resources, 2020, 5, 2621-2623.	0.2	3
191	The complete chloroplast genome of <i>Douinia plicata</i> (Lindb.) Konstant. & Vilnet (Scapaniaceae, Jungermanniales). Mitochondrial DNA Part B: Resources, 2020, 5, 3680-3682.	0.2	3
192	Complete mitochondrial genome of the gate-keeper ant <i>Colobopsis nipponica</i> (Wheeler, W.M., 1928) (Formicidae: Hymenoptera). Mitochondrial DNA Part B: Resources, 2021, 6, 86-88.	0.2	3
193	Taxonomic studies of the tribe Potentilleae (Rosaceae) in Korea. Korean Journal of Plant Taxonomy, 2019, 49, 28-69.	0.3	3
194	The complete chloroplast genome of <i>Limonium tetragonum</i> (Plumbaginaceae) isolated in Korea. Korean Journal of Plant Taxonomy, 2021, 51, 337-344.	0.3	3
195	The complete mitochondrial genome of <i>Uroleucon erigeronense</i> (Thomas, 1878) (Hemiptera). Tj ETQq1 1 0.784314 rgBT /Over	0.2	3
196	The complete mitochondrial genomes from three body color variants of sea cucumbers, <i>Apostichopus japonicus</i> (Selenka, 1867). Mitochondrial DNA Part B: Resources, 2019, 4, 836-837.	0.2	2
197	First Record of the Complete Mitochondrial Genome of a Saprotrophic and Opportunistic Human Pathogenic Fungus, <i>Scopulariopsis brevicaulis</i> . Mycobiology, 2020, 48, 528-531.	0.6	2
198	The complete chloroplast genome of <i>Wiesnerella denudata</i> (Mitt.) Steph. (Wiesnerellaceae). Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 62 To	0.2	2

#	ARTICLE	IF	CITATIONS
199	The complete mitochondrial genome of the far Eastern myotis: <i>Myotis bombinus</i> Thomas, 1906 in mainland of Korea (Chiroptera, Vespertilionidae). Mitochondrial DNA Part B: Resources, 2021, 6, 615-616.	0.2	2
200	Assaying Mitochondrial COI Sequences and Their Molecular Studies in Hexapoda, PART I: From 2000 to 2009. Korean Journal of Applied Entomology, 2013, 52, 395-402.	0.3	2
201	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
202	Complete mitochondrial genome sequence of <i>Aspergillus flavus</i> SRRC1009: insight of intraspecific variations on <i>A. flavus</i> mitochondrial genomes. Mitochondrial DNA Part B: Resources, 2020, 5, 3567-3569.	0.2	1
203	The complete chloroplast genome of <i>Oxybasis glauca</i> (L.) S. Fuentes, Uotila & Borsch (Amaranthaceae) as the first chloroplast genome in genus <i>Oxybasis</i> . Mitochondrial DNA Part B: Resources, 2020, 5, 1410-1412.	0.2	1
204	PlantGPS: A Mobile Application for Collecting Biogeographical Information of Plants and for Their Monitoring. Korean Journal of Nature Conservation, 2016, 10, 9-17.	0.2	0
205	The complete chloroplast genome of <i>Atriplex gmelinii</i> C. A. Mey. ex Bong. (Amaranthaceae). Mitochondrial DNA Part B: Resources, 2022, 7, 541-543.	0.2	0
206	The complete chloroplast genome of <i>Utricularia tenuicaulis</i> Miki (Lentibulariaceae) isolated in Korea. Mitochondrial DNA Part B: Resources, 2022, 7, 1143-1145.	0.2	0