

# Chang-Bae Kim

## List of Publications by Year in descending order

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Version: 2024-02-01

34  
papers

141  
citations

1478505

6  
h-index

1372567

10  
g-index

34  
all docs

34  
docs citations

34  
times ranked

140  
citing authors

#	ARTICLE	IF	CITATIONS
1	Comparative Transcriptome Analysis for Understanding Predator-Induced Polyphenism in the Water Flea <i>Daphnia pulex</i> . <i>International Journal of Molecular Sciences</i> , 2018, 19, 2110.	4.1	20
2	SortPred: The first machine learning based predictor to identify bacterial sortases and their classes using sequence-derived information. <i>Computational and Structural Biotechnology Journal</i> , 2022, 20, 165-174.	4.1	15
3	Molecular Characterization and Gene Expression of Glutathione Peroxidase 1 in <i>Tor tambroides</i> Exposed to Temperature Stress. <i>Evolutionary Bioinformatics</i> , 2019, 15, 117693431985358.	1.2	14
4	Role of Transcriptome in the Gills of Chinese Mitten Crabs in Response to Salinity Change: A Meta-Analysis of RNA-Seq Datasets. <i>Biology</i> , 2021, 10, 39.	2.8	13
5	The complete mitochondrial genome of <i>Phyllidiella pustulosa</i> (Cuvier, 1804) (Nudibranchia, Tj ETQq1 1 0.784314 rgBT /Overlock 0.4 5	0.4	5
6	The complete mitochondrial genome of a blowfly <i>Calliphora nigribarbis</i> (Vollenhoven, 1863) (Insecta: Diptera: Calliphoridae). <i>Mitochondrial DNA Part B: Resources</i> , 2019, 4, 2318-2319.	0.4	7
7	Complete mitogenome and phylogenetic analysis of hide beetle <i>Dermestes maculatus</i> (Insecta, Tj ETQq1 1 0.784314 rgBT /Overlock 0.4 6	0.4	6
8	Comparative analyses of the three complete mitochondrial genomes from forensic important beetle genus <i>Dermestes</i> with phylogenetic relationships. <i>Gene</i> , 2019, 706, 146-153.	2.2	6
9	Molecular Identification of Necrophagous <i>Dermestes</i> Species in South Korea Using Cytochrome c Oxidase Subunit I Nucleotide Sequences (Genus <i>Dermestes</i> ). <i>Journal of Forensic Sciences</i> , 2020, 65, 283-287.	1.6	6
10	Complete mitochondrial DNA analysis of <i>Hydrotaea ignava</i> (Insecta, Diptera, Muscidae). <i>Mitochondrial DNA Part B: Resources</i> , 2017, 2, 825-826.	0.4	5
11	The complete mitogenome of brownbranded moon jellyfish <i>Aurelia limbata</i> (Cnidaria, Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 302 1875-1876.	0.4	5
12	Complete mitochondrial genome analysis of <i>Lingula anatina</i> from Korea (Brachiopoda, Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 302 0.4 4	0.4	4
13	Analysis of complete mitochondrial genome of <i>Etisus anaglyptus</i> (Arthropoda, Decapoda, Tj ETQq1 1 0.784314 rgBT /Overlock 0.4 3	0.4	3
14	Characterization of the complete mitochondrial genome of <i>Hermisenda emurai</i> (Baba, 1937) (Nudibranchia, Facelinidae). <i>Mitochondrial DNA Part B: Resources</i> , 2019, 4, 860-861.	0.4	3
15	Seasonal Diversity of Microeukaryotes in the Han River, Korea Through 18S rRNA Gene Metabarcoding. <i>Evolutionary Bioinformatics</i> , 2022, 18, 117693432210746.	1.2	3
16	Complete mitochondrial genome of <i>Laqueus Japonicus</i> (Brachiopoda, Terebratulida, Laqueidae). <i>Mitochondrial DNA Part B: Resources</i> , 2017, 2, 883-884.	0.4	2
17	Complete mitochondrial genome of <i>Atergatis integerrimus</i> (Decapoda, Xanthidae) from the Philippines. <i>Mitochondrial DNA Part B: Resources</i> , 2018, 3, 205-206.	0.4	2
18	The complete mitogenome of ghost jellyfish <i>Cyanea nozakii</i> (Cnidaria, Semaestomeae, Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 62 0.4 2	0.4	2

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19	The complete mitochondrial genome of <i>Hemigrapsus penicillatus</i> (De Haan, 1835) (Decapoda, Tj ETQq1 1 0.784314 rgBT /Overlock 10 TF	0.4	2
20	Characterization and Comparative Analysis of Complete Mitogenomes of Three <i>Cacatua</i> Parrots (Psittaciformes: Cacatuidae). <i>Genes</i> , 2021, 12, 209.	2.4	2
21	An Investigation of the Variations in Complete Mitochondrial Genomes of <i>Lingula anatina</i> in the Western Pacific Region. <i>Biology</i> , 2021, 10, 367.	2.8	2
22	A report of a new species and new record of <i>Cadlina</i> (Nudibranchia, Cadlinidae) from South Korea. <i>ZooKeys</i> , 2020, 996, 1-18.	1.1	2
23	Complete mitochondrial genome of xanthid crab <i>Atergatis floridus</i> (Linnaeus, 1767). <i>Mitochondrial DNA Part B: Resources</i> , 2018, 3, 202-204.	0.4	1
24	Characterization of the complete mitochondrial genome of <i>Triceratopyga calliphoroides</i> (Rohdendorf, 1931) (Insecta: Diptera: Calliphoridae). <i>Mitochondrial DNA Part B: Resources</i> , 2019, 4, 2183-2184.	0.4	1
25	Sequencing and analysis of the complete mitogenome of <i>Doriprismatica atromarginata</i> (Cuvier, 1804). <i>Mitochondrial DNA Part B: Resources</i> , 2019, 4, 2894-2895.	0.4	1
26	Analysis of complete mitochondrial genome of <i>Matuta planipes</i> (Decapoda, Brachyura, Matutidae) from South Korea. <i>Mitochondrial DNA Part B: Resources</i> , 2019, 4, 906-907.	0.4	1
27	The mitochondrial genome analysis of <i>Protaeolidiella atra</i> Baba, 1955 from Korea. <i>Mitochondrial DNA Part B: Resources</i> , 2020, 5, 1277-1278.	0.4	1
28	Characterization of the complete mitochondrial genome of <i>Okenia hiroi</i> (Baba, 1938) (Nudibranchia, Tj ETQq0 0 0 rgBT /Overlock 10 TF	0.4	1
29	The complete mitochondrial genome of <i>Scyra compressipes</i> Stimpson, 1857 (Decapoda: Tj ETQq1 1 0.784314 rgBT /Overlock 10 TF	0.4	1
30	Development of polymorphic microsatellite markers for the <i>Trichoglossus haematodus</i> and cross-species amplification in <i>Trichoglossus moluccanus</i> . <i>Molecular Biology Reports</i> , 2021, 48, 5787-5793.	2.3	1
31	The complete mitochondrial genome of <i>Haloa japonica</i> (Pilsbry, 1895) (Cephalaspidea, Haminoeidae). <i>Mitochondrial DNA Part B: Resources</i> , 2020, 5, 1275-1276.	0.4	0
32	Image Identification of Multiple Parrot Species Belonging to CITES Using Deep Neural Networks. <i>Lecture Notes in Computer Science</i> , 2021, , 248-253.	1.3	0
33	Sequencing and analysis of the mitochondrial genome of <i>Pennahia anea</i> (Bloch, 1793) (Perciformes: Sciaenidae). <i>Mitochondrial DNA Part B: Resources</i> , 2021, 6, 1389-1391.	0.4	0
34	Development of the new microsatellite markers of <i>Lucilia sericata</i> (Diptera: Calliphoridae) from Korea. <i>Molecular Biology Reports</i> , 2021, 48, 8245-8248.	2.3	0