Jongwoo Jung

List of Publications by Year
in descending order

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33	167	1477746 6	1281420 11
papers	citations	h-index	g-index

36	36	36	193
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	First record of the complete mitochondrial genome and phylogenetic analysis of Limnodrilus hoffmeisteri ClaparÃ¨de, 1862 (Annelida; Clitellata; Oligochaeta). Mitochondrial DNA Part B: Resources, 2022, 7, 177-179.	0.2	0
2	The complete mitochondrial genome of Nais communis Piguet, 1906 (Annelida; Clitellata; Naididae). Mitochondrial DNA Part B: Resources, 2022, 7, 199-201.	0.2	1
3	Population genetic analysis reveals secondary contact between Eriocheir sinensis and E. japonica in South Korea. Genes and Genomics, 2022, , 1.	0.5	1
4	First record of the complete mitochondrial genome of Tubifex tubifex (MÃ¼ller) 1774 (Annelida;) Tj ETQq0 0 0 rgBT /Overlock 10 1208-1210.	0.2	2
5	Complete mitochondrial genome of Aedes flavopictus (Yamada, 1921) (Diptera: Culicidae) collected in South Korea. Mitochondrial DNA Part B: Resources, 2021, 6, 265-267.	0.2	2
6	Complete mitochondrial genome of Cyclograpsus intermedius Ortmann, 1894 (Crustacea: Decapoda:) Tj ETQq0 0 0 rgBT /Overlock 10 T	0.2	1
7	First record of the complete mitochondrial genome of the mantis shrimp, Gonodactylaceus randalli (Manning, 1978) (Stomatopoda: Gonodactylidae). Mitochondrial DNA Part B: Resources, 2021, 6, 510-511.	0.2	2
8	Comparative population genetics of the invasive mosquito Aedes albopictus and the native mosquito Aedes flavopictus in the Korean peninsula. Parasites and Vectors, 2021, 14, 377.	1.0	3
9	Complete mitochondrial genome of the commensal scale worm, Arctonoe vittata (Grube, 1855) (Polychaeta: Polynoidae), collected from benthic habitat of the eastern coast of Korea. Mitochondrial DNA Part B: Resources, 2021, 6, 2455-2457.	0.2	1
10	The mitochondrial genome of Faughnia haani (Stomatopoda): novel organization of the control region and phylogenetic position of the superfamily Parasquilloidea. BMC Genomics, 2021, 22, 716.	1.2	1
11	Complete mitochondrial genome of the mantis shrimp Taku spinosocarinatus (Fukuda, 1909) (Stomatopoda: Gonodactyloidea: Takuidae) in South Korea. Mitochondrial DNA Part B: Resources, 2020, 5, 3609-3610.	0.2	2
12	The complete mitochondrial genome of an interstitial polychaete, Pharyngocirrus uchidai (Annelida,) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5	0.2	0
13	Complete mitochondrial genome of the ghost crab Ocypode stimpsoni Ortmann, 1897 (Brachyura:) Tj ETQq1 1 0.784314 rgBT /Overl Resources, 2020, 5, 1699-1700.	0.2	4
14	First record of the complete mitochondrial genome of Aedes koreicus (Edwards, 1917) (Diptera:) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 2	0.2	3
15	Complete mitochondrial genome of the mantis shrimp, Chorisquilla orientalis Hwang, Ahyong, & Kim, 2018 (Stomatopoda: Protosquillidae). Mitochondrial DNA Part B: Resources, 2020, 5, 3823-3824.	0.2	1
16	Phylogeographic studies on two shore crab species from East Asia: similar but different stories. Genes and Genomics, 2019, 41, 1127-1134.	0.5	3
17	Development of 17 polymorphic microsatellite loci from Jeju striped field mouse, Apodemus agrarius chejuensis (Rodentia: Muridae), by 454 pyrosequencing. Hereditas, 2018, 155, 30.	0.5	1
18	Environmental factors affecting population level genetic divergence of the striped field mouse (Apodemus agrarius) in South Korea. Ecological Research, 2018, 33, 989-999.	0.7	3

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19	Subdividing the Common Intertidal Hermit Crab Hess, 1865 (Decapoda: Anomura: Paguridae) Based on Molecular, Morphological and Coloration Analyses. Zoological Studies, 2018, 57, e61.	0.3	5
20	Out of Asia: mitochondrial evolutionary history of the globally introduced supralittoral isopodLigia exotica. PeerJ, 2018, 6, e4337.	0.9	9
21	Genetic differentiation of the Korean striped field mouse, Apodemus agrarius (Muridae, Rodentia), based on microsatellite polymorphism. Mammalia, 2017, 81, .	0.3	5
22	Characterization of 15 microsatellite loci and genetic analysis of Heterodera schachtii (Nematoda:) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50	0.6	9
23	Comparison of Population Genetic Structure of Two Seashore-Dwelling Animal Species, Periwinkle Littorina brevicula and Acorn Barnacle Fistulobalanus albicostatus from Korea. Animal Systematics, Evolution and Diversity, 2016, 32, 105-111.	0.2	3
24	Population Genetic Structure of the Malaria VectorAnopheles sinensis(Diptera: Culicidae) Sensu Stricto and Evidence for Possible Introgression in the Republic of Korea. Journal of Medical Entomology, 2015, 52, 1270-1281.	0.9	6
25	Six Korean New Records of the Nais Species (Annelida, Clitellata, Naididae). Hangug Hwangyeong Saengmul Haghoeji, 2015, 33, 153-159.	0.1	3
26	Korea Barcode of Life Database System (KBOL). Animal Cells and Systems, 2012, 16, 11-19.	0.8	5
27	The polymorphism and the geographical distribution of the knockdown resistance (kdr) of Anopheles sinensis in the Republic of Korea. Malaria Journal, 2012, 11, 151.	0.8	31
28	Isolation and characterization of microsatellite markers for the endangered crab species Sesarmops intermedius. Conservation Genetics Resources, 2012, 4, 183-185.	0.4	0
29	Isolation and characterization of microsatellite loci in the Korean crayfish,Cambaroides similisand application to natural population analysis. Animal Cells and Systems, 2011, 15, 37-43.	0.8	9
30	Microsatellite variation in the pinewood nematode, Bursaphelenchus xylophilus (Steiner and Buhrer) Nickle in South Korea. Genes and Genomics, 2010, 32, 151-158.	0.5	29
31	A preliminary study on the origin of Koreans based on Y-STR variation. Genes and Genomics, 2010, 32, 353-359.	0.5	4
32	Amplified fragment length polymorphism analysis and genetic variation of the pinewood nematodeBursaphelenchus xylophilusin South Korea. Animal Cells and Systems, 2010, 14, 31-36.	0.8	7
33	Two genetic lineages of sea slaters, Ligia (Crustacea: Isopoda) in South Korea: a population genetic approach. Molecules and Cells, 2008, 25, 523-30.	1.0	11