Mi-Sook Min

List of Publications by Year
in descending order

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52	676	686830 13	642321 23
papers	citations	h-index	g-index

53	53	53	911
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Spatiotemporal Diversification of the True Frogs (GenusRana): A Historical Framework for a Widely Studied Group of Model Organisms. Systematic Biology, 2016, 65, 824-842.	2.7	125
2	Genetic evidence for a high diversity and wide distribution of endemic strains of the pathogenic chytrid fungus <scp>B</scp>atrachochytrium dendrobatidis in wild <scp>A</scp>sian amphibians. Molecular Ecology, 2013, 22, 4196-4209.	2.0	113
3	Robust molecular phylogeny and palaeodistribution modelling resolve a complex evolutionary history: glacial cycling drove recurrent mt<scp>DNA</scp> introgression among Pelophylax frogs in East Asia. Journal of Biogeography, 2015, 42, 2159-2171.	1.4	37
4	Asia-wide phylogeography of wild boar (Sus scrofa) based on mitochondrial DNA and Y-chromosome: Revising the migration routes of wild boar in Asia. PLoS ONE, 2020, 15, e0238049.	1.1	23
5	Evolutionary and biogeographical implications of variation in skull morphology of raccoon dogs (Nyctereutes procyonoides, Mammalia: Carnivora). Biological Journal of the Linnean Society, 2015, 116, 856-872.	0.7	21
6	Diversity and phylogeography of Northeast Asian brown frogs allied to Rana dybowskii (Anura,) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 54	1.2	21
7	Yellow sea mediated segregation between North East Asian Dryophytes species. PLoS ONE, 2020, 15, e0234299.	1.1	21
8	Genetic diversity and genetic structure of the Siberian roe deer (Capreolus pygargus) populations from Asia. BMC Genetics, 2015, 16, 100.	2.7	20
9	Influence of geology and human activity on the genetic structure and demography of the Oriental fire-bellied toad (Bombina orientalis). Molecular Phylogenetics and Evolution, 2016, 97, 69-75.	1.2	20
10	Molecular phylogenetic status of the Korean goral and Japanese serow based on partial sequences of the mitochondrial cytochrome b gene. Molecules and Cells, 2004, 17, 365-72.	1.0	20
11	Individual identification and sex determination of Eurasian otters (Lutra lutra) in Daegu city based on genetic analysis of otter spraint. Genes and Genomics, 2011, 33, 653-657.	0.5	18
12	A new species of salamander of the genus HynobiusÂ(Amphibia, Caudata, Hynobiidae) from South Korea. Zootaxa, 2016, 4169, 475-503.	0.2	15
13	Population genetic study of the raccoon dog (<i>Nyctereutes procyonoides</i>) in South Korea using newly developed 12 microsatellite markers. Genes and Genetic Systems, 2013, 88, 69-76.	0.2	13
14	Genetic Diversity and Population Structure of East Asian Raccoon Dog (Nyctereutes) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 227 Td (p 249-259.	0.3	13
15	Disentangling the Impacts of Speciation, Sympatry and the Island Effect on the Morphology of Seven Hynobius sp. Salamanders. Animals, 2021, 11, 187.	1.0	12
16	Molecular and Morphological Evidence for Rana kunyuensis as a Junior Synonym of Rana coreana (Anura: Ranidae). Journal of Herpetology, 2015, 49, 302.	0.2	11
17	Evaluation of biochemical and haematological parameters and prevalence of selected pathogens in feral cats from urban and rural habitats in South Korea. Journal of Feline Medicine and Surgery, 2016, 18, 443-451.	0.6	11
18	Phylogenetic relationships of three representative sea krait species (genus Laticauda; elapidae;) Tj ETQq0 0 0 rgBT /Overlock 10 T Analysis, 2018, 29, 772-777.	0.7	11

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19	Isolation and characterization of 15 microsatellite loci in the Korean goral (Nemorhaedus caudatus). Molecular Ecology Notes, 2005, 5, 421-423.	1.7	10
20	Age structure and growth rates of two Korean salamander species (Hynobius) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 707 Td (yangi 14, 315-322.	0.8	10
21	Population genetic structure of endangered Mongolian racerunner (Eremias argus) from the Korean Peninsula. Molecular Biology Reports, 2014, 41, 7339-7347.	1.0	10
22	Phylogeography of the Asian lesser white-toothed shrew, Crocidura shantungensis, in East Asia: role of the Korean Peninsula as refugium for small mammals. Genetica, 2018, 146, 211-226.	0.5	9
23	Organization and variation of the mitochondrial DNA control region in five Caprinae species. Genes and Genomics, 2010, 32, 335-344.	0.5	8
24	Genetic origin identification of Siberian chipmunks (Tamias sibiricus) in pet shops of South Korea. Animal Cells and Systems, 2011, 15, 161-168.	0.8	8
25	Origin of sex chromosomes in six groups of Rana rugosa frogs inferred from a sexâ€linked DNA marker. Journal of Experimental Zoology Part A: Ecological and Integrative Physiology, 2017, 327, 444-452.	0.9	8
26	Phylogenetic structure and ancestry of Korean clawed salamander,Onychodactylus koreanus(Caudata: Hynobiidae). Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis, 2018, 29, 650-658.	0.7	6
27	The Asian plethodontid salamander preserves historical genetic imprints of recent northern expansion. Scientific Reports, 2021, 11, 9193.	1.6	6
28	Phylogeographic study of the Bufo gargarizans species complex, with emphasis on Northeast Asia. Animal Cells and Systems, 2021, 25, 434-444.	0.8	6
29	Isolation and characterization of 12 microsatellite loci from Korean water deer (Hydropotes inermis) Tj ETQq1 1 0.784314 rgBT /Overl	0.5	5
30	Korea Barcode of Life Database System (KBOL). Animal Cells and Systems, 2012, 16, 11-19.	0.8	5
31	Sequencing and analysis of the complete mitochondrial genome of Hyla suweonensis (Anura:) Tj ETQq1 1 0.784314 rgBT /Ove	0.2	5
32	What is the taxonomic status of East Asian otter species based on molecular evidence?: focus on the position of the Japanese otter holotype specimen from museum. Animal Cells and Systems, 2019, 23, 228-234.	0.8	5
33	Genetic and phylogenetic structure of Hynobius quelpaertensis, an endangered endemic salamander species on the Korean Peninsula. Genes and Genomics, 2020, 42, 165-178.	0.5	5
34	Development of 10 microsatellite loci from the Korean Ratsnake (Elaphe schrenckii) and its application across Elaphe species from South Korea, Russia, and China. Genes and Genomics, 2010, 32, 401-405.	0.5	4
35	Genetic diversity and population demography of narrowâ€ridged finless porpoises from South Korea on the basis of mitochondrial DNA variation: Implications for its conservation in East Asia. Marine Mammal Science, 2019, 35, 574-594.	0.9	4
36	Phylogenetic relationships between different raccoon dog (Nyctereutes procyonoides) populations based on four nuclear and Y genes. Genes and Genomics, 2020, 42, 1075-1085.	0.5	4

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37	Genetic diversity and inferred dispersal history of the Schlegelâ€™s Japanese Gecko (Gekko japonicus) in Northeast Asia based on population genetic analyses and paleo-species distribution modelling. Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis, 2020, 31, 120-130.	0.7	4
38	Phylogenetic study of extirpated Korean leopard using mitochondrial DNA from an old skin specimen in South Korea. PeerJ, 2020, 8, e8900.	0.9	4
39	Population structure of the raccoon dog (Nyctereutes procyonoides) using microsatellite loci analysis in South Korea: Implications for disease management. Journal of Veterinary Medical Science, 2018, 80, 1631-1638.	0.3	3
40	Genetic and morphologic diversity of the moles (Talpomorpha, Talpidae, Mogera) from the continental Far East. Journal of Zoological Systematics and Evolutionary Research, 2019, 57, 662.	0.6	3
41	The complete mitochondrial genome information of Rana uenoi (Amphibia, Anura, Ranidae) and the phylogenetic implication. Mitochondrial DNA Part B: Resources, 2021, 6, 689-690.	0.2	3
42	Whole genome survey of big cats (Genus: Panthera) identifies novel microsatellites of utility in conservation genetic study. Scientific Reports, 2021, 11, 14164.	1.6	3
43	The complete mitochondrial genome information of Hynobius unisacculus (Amphibia, Caudata,) Tj ETQq1 1 0.784314 rgBT /Ov	0.2	2
44	Development and characterization of nine microsatellite loci from the Korean hare (Lepus coreanus) and genetic diversity in South Korea. Animal Cells and Systems, 2012, 16, 230-236.	0.8	1
45	Genetic diversity and phylogeography of Jeju Orthohantavirus (Hantaviridae) in the Republic of Korea. Virology, 2020, 543, 13-19.	1.1	1
46	The complete mitochondrial genome of the Amur soft-shelled turtle (Pelodiscus maackii) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 38	0.2	1
47	Yellow sea mediated segregation between North East Asian Dryophytes species. , 2020, 15, e0234299.		0
48	Yellow sea mediated segregation between North East Asian Dryophytes species. , 2020, 15, e0234299.		0
49	Yellow sea mediated segregation between North East Asian Dryophytes species. , 2020, 15, e0234299.		0
50	Yellow sea mediated segregation between North East Asian Dryophytes species. , 2020, 15, e0234299.		0
51	Yellow sea mediated segregation between North East Asian Dryophytes species. , 2020, 15, e0234299.		0
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