Naoko Takezaki

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Genetic Distances and Reconstruction of Phylogenetic Trees From Microsatellite DNA. Genetics, 1996, 144, 389-399.	2.9	1,038
2	POPTREE2: Software for Constructing Population Trees from Allele Frequency Data and Computing Other Population Statistics with Windows Interface. Molecular Biology and Evolution, 2010, 27, 747-752.	8.9	587
3	The <i>N</i> â€Acylethanolamineâ€Hydrolyzing Acid Amidase (NAAA). Chemistry and Biodiversity, 2007, 4, 1914-1925.	2.1	163
4	POPTREEW: Web Version of POPTREE for Constructing Population Trees from Allele Frequency Data and Computing Some Other Quantities. Molecular Biology and Evolution, 2014, 31, 1622-1624.	8.9	156
5	Molecular Phylogeny of Early Vertebrates: Monophyly of the Agnathans as Revealed by Sequences of 35 Genes. Molecular Biology and Evolution, 2003, 20, 287-292.	8.9	141
6	The Phylogenetic Relationship of Tetrapod, Coelacanth, and Lungfish Revealed by the Sequences of Forty-Four Nuclear Genes. Molecular Biology and Evolution, 2004, 21, 1512-1524.	8.9	136
7	Empirical Tests of the Reliability of Phylogenetic Trees Constructed With Microsatellite DNA. Genetics, 2008, 178, 385-392.	2.9	84
8	Mhc class II B gene evolution in East African cichlid fishes. Immunogenetics, 2000, 51, 556-575.	2.4	46
9	Origin and Speciation of Haplochromine Fishes in East African Crater Lakes Investigated by the Analysis of Their mtDNA, Mhc Genes, and SINEs. Molecular Biology and Evolution, 2003, 20, 1448-1462.	8.9	41
10	Identification and characterization of a TAP-family gene in the lamprey. Immunogenetics, 2003, 55, 38-48.	2.4	40
11	Origin and affinities of indigenous Siberian populations as revealed by HLA class II gene frequencies. Human Genetics, 2002, 110, 209-226.	3.8	37
12	Resolving the Phylogenetic Position of Coelacanth: The Closest Relative Is Not Always the Most Appropriate Outgroup. Genome Biology and Evolution, 2016, 8, 1208-1221.	2.5	29
13	Global Rate Variation in Bony Vertebrates. Genome Biology and Evolution, 2018, 10, 1803-1815.	2.5	28
14	Genomic Drift and Evolution of Microsatellite DNAs in Human Populations. Molecular Biology and Evolution, 2009, 26, 1835-1840.	8.9	12
15	Mhc class I genes of the cichlid fish Oreochromis niloticus. Immunogenetics, 2006, 58, 917-928.	2.4	11
16	Support for Lungfish as the Closest Relative of Tetrapods by Using Slowly Evolving Ray-finned fish as the Outgroup. Genome Biology and Evolution, 2017, 9, evw288.	2.5	11
17	Sequencing of amphioxus PSMB5 / 8 gene and phylogenetic position of agnathan sequences. Gene, 2002, 282, 179-187.	2.2	9
18	Ancestry and kinships of native Siberian populations: The HLA evidence. Evolutionary Anthropology, 2003, 12, 231-245.	3.4	9

#	Article	IF	CITATIONS
19	Resolving the Early Divergence Pattern of Teleost Fish Using Genome-Scale Data. Genome Biology and Evolution, 2021, 13, .	2.5	7
20	CNVs and Microsatellite DNA Polymorphism. Evolutionary Studies, 2017, , 143-155.	0.1	2