Naoko Takezaki

List of Publications by Year in descending order

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NAOKO TAKEZAKI

#	Article	IF	CITATIONS
1	Resolving the Early Divergence Pattern of Teleost Fish Using Genome-Scale Data. Genome Biology and Evolution, 2021, 13, .	2.5	7
2	Global Rate Variation in Bony Vertebrates. Genome Biology and Evolution, 2018, 10, 1803-1815.	2.5	28
3	CNVs and Microsatellite DNA Polymorphism. Evolutionary Studies, 2017, , 143-155.	0.1	2
4	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
5	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
6	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
7	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
8	Genomic Drift and Evolution of Microsatellite DNAs in Human Populations. Molecular Biology and Evolution, 2009, 26, 1835-1840.	8.9	12
9	Empirical Tests of the Reliability of Phylogenetic Trees Constructed With Microsatellite DNA. Genetics, 2008, 178, 385-392.	2.9	84
10	The <i>N</i> â€Acylethanolamineâ€Hydrolyzing Acid Amidase (NAAA). Chemistry and Biodiversity, 2007, 4, 1914-1925.	2.1	163
11	Mhc class I genes of the cichlid fish Oreochromis niloticus. Immunogenetics, 2006, 58, 917-928.	2.4	11
12	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
13	Identification and characterization of a TAP-family gene in the lamprey. Immunogenetics, 2003, 55, 38-48.	2.4	40
14	Ancestry and kinships of native Siberian populations: The HLA evidence. Evolutionary Anthropology, 2003, 12, 231-245.	3.4	9
15	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
16	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
17	Sequencing of amphioxus PSMB5 / 8 gene and phylogenetic position of agnathan sequences. Gene, 2002, 282, 179-187.	2.2	9
18	Origin and affinities of indigenous Siberian populations as revealed by HLA class II gene frequencies. Human Genetics, 2002, 110, 209-226.	3.8	37

#	Article	IF	CITATIONS
19	Mhc class II B gene evolution in East African cichlid fishes. Immunogenetics, 2000, 51, 556-575.	2.4	46
20	Genetic Distances and Reconstruction of Phylogenetic Trees From Microsatellite DNA. Genetics, 1996, 144, 389-399.	2.9	1,038