Posterior Summarization in Bayesian Phylogenetics Usi

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Citation Report

#	Article	IF	Citations
1	Cryptic Diversity Hidden within the Leafminer Genus Liriomyza (Diptera: Agromyzidae). Genes, 2018, 9, 554.	1.0	8
2	Maintenance and reappearance of extremely divergent intra-host HIV-1 variants. Virus Evolution, 2018, 4, vey030.	2.2	5
3	Interspecies evolutionary divergence in Liriodendron, evidence from the nucleotide variations of LcDHN-like gene. BMC Evolutionary Biology, 2018, 18, 195.	3.2	10
4	Strong and complex host- and habitat-associated genetic differentiation in an apparently polyphagous leaf mining insect. Biological Journal of the Linnean Society, 2018, 125, 885-899.	0.7	12
5	Marine Macrodasyida (Gastrotricha) from Hokkaido, Northern Japan. Species Diversity, 2018, 23, 183-192.	0.1	5
6	Deep Intraspecific Divergence in the Endemic Herb Lancea tibetica (Mazaceae) Distributed Over the Qinghai-Tibetan Plateau. Frontiers in Genetics, 2018, 9, 492.	1.1	6
7	Modeling site-specific amino-acid preferences deepens phylogenetic estimates of viral sequence divergence. Virus Evolution, 2018, 4, vey033.	2.2	11
8	Genomic surveillance of Neisseria gonorrhoeae to investigate the distribution and evolution of antimicrobial-resistance determinants and lineages. Microbial Genomics, 2018, 4, .	1.0	47
9	Emergence of genotype Cosmopolitan of dengue virus type 2 and genotype III of dengue virus type 3 in Thailand. PLoS ONE, 2018, 13, e0207220.	1.1	38
10	Morphological and molecular characterization of Lobophora declerckii and L. variegata (Dictyotales,) Tj ETQq1 1 0.784314 rgBT /Over	0.1	5
11	The Gambian epauletted fruit bat shows increased genetic divergence in the Ethiopian highlands and in an area of rapid urbanization. Ecology and Evolution, 2018, 8, 12803-12820.	0.8	6
12	Phylogenomic and single nucleotide polymorphism analyses revealed the hybrid origin of Spondias bahiensis (family Anacardiaceae): de novo genome sequencing and comparative genomics. Genetics and Molecular Biology, 2018, 41, 878-883.	0.6	6
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14	Rediscovery of the Poorly KnownAmphisbaena bahianaVanzolini, 1964 (Squamata, Amphisbaenidae), with Data on Its Phylogenetic Placement, External Morphology and Natural History. South American Journal of Herpetology, 2018, 13, 238-248.	0.5	6
15	Dynamic biogeographic models and dinosaur origins. Earth and Environmental Science Transactions of the Royal Society of Edinburgh, 2018, 109, 325-332.	0.3	8
16	Molecular phylogeny and systematics of the centipede genus Ethmostigmus Pocock (Chilopoda :) Tj ETQq1 1 0.784314 rgBT /Overlock	0.5	10
17	Redescription and phylogenetic placement of Cirrhilabrus sanguineus Cornic (Teleostei: Labridae), with first documentation of the female formÂ. Zootaxa, 2018, 4526, 358.	0.2	6
18	Phylogeny, biogeography and taxonomic re-assessment of Multifurca (Russulaceae, Russulales) using three-locus data. PLoS ONE, 2018, 13, e0205840.	1.1	10

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19	Characterization of the complete mitochondrial genome of Hydrotaea spinigera (Diptera: Muscidae) with phylogenetic implications. Journal of Asia-Pacific Entomology, 2018, 21, 1416-1423.	0.4	3
20	Morphology and Phylogeny of Calyptospora paranaidji n. sp. (Eimeriorina: Calyptosporidae), an Apicomplexan Parasite of the Hepatic Tissue of Cichla piquiti Kullander & Ferreira, 2006, From a Reservoir in the Brazilian Amazon Region. Journal of Eukaryotic Microbiology, 2018, 66, 608-616.	0.8	3
21	Comparative analysis of two mitochondrial genomes of flesh flies (Sarcophaga antilope and) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 667 T of Biological Macromolecules, 2018, 120, 1955-1964.	3.6	9
22	High Throughput Identification of Antihypertensive Peptides from Fish Proteome Datasets. Marine Drugs, 2018, 16, 365.	2.2	12
23	A new species of Tanaecium (Bignonieae, Bignoniaceae) from the Brazilian Amazon and its phylogenetic placement. Plant Systematics and Evolution, 2018, 304, 1245-1253.	0.3	6
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25	Assessing genomic admixture between cryptic Plutella moth species following secondary contact. Genome Biology and Evolution, 2018, 10, 2973-2985.	1.1	5
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27	Genetic structure of the grey side-gilled sea slug (Pleurobranchaea maculata) in coastal waters of New Zealand. PLoS ONE, 2018, 13, e0202197.	1.1	7
28	Diversity and taxonomy of Tricholoma species from Yunnan, China, and notes on species from Europe and North America. Mycologia, 2018, 110, 1081-1109.	0.8	18
29	Characterization of Begomoviruses Sampled during Severe Epidemics in Tomato Cultivars Carrying the Ty-1 Gene. International Journal of Molecular Sciences, 2018, 19, 2614.	1.8	14
30	Origin and macroevolution of micro-moths on sunken Hawaiian Islands. Proceedings of the Royal Society B: Biological Sciences, 2018, 285, 20181047.	1.2	24
31	Phacidiaceae endophytes of Picea rubens in Eastern Canada. Botany, 2018, 96, 555-588.	0.5	11
32	Genomic Epidemiology Reconstructs the Introduction and Spread of Zika Virus in Central America and Mexico. Cell Host and Microbe, 2018, 23, 855-864.e7.	5.1	82
33	Allele Phasing Greatly Improves the Phylogenetic Utility of Ultraconserved Elements. Systematic Biology, 2019, 68, 32-46.	2.7	74
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35	The effect of dictionary omissions on phylogenies computationally inferred from lexical data. Language Dynamics and Change, 2018, 8, 78-107.	0.4	1
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37	Rapid evolutionary dynamics of pepper mild mottle virus. Virus Research, 2018, 256, 96-99.	1.1	7
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39	On the roles of landscape heterogeneity and environmental variation in determining population genomic structure in a dendritic system. Molecular Ecology, 2018, 27, 3484-3497.	2.0	52
40	Chronic and Occult Hepatitis B Virus Infection in Pregnant Women in Botswana. Genes, 2018, 9, 259.	1.0	21
41	Integrated genomic and fossil evidence illuminates lifeâ€™s early evolution and eukaryote origin. Nature Ecology and Evolution, 2018, 2, 1556-1562.	3.4	274
42	The unusual flagellar targeting mechanism and functions of the trypanosome orthologue of the ciliary GTPase Arl13b. Journal of Cell Science, 2018, 131, .	1.2	5
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46	Phylogenetic relationships of Neotropical lady ferns (Athyriaceae), with a description of Ephemeropteris , gen. nov.. Taxon, 2019, 68, 425-441.	0.4	4
47	Comparative genomics suggests loss of keratin K24 in three evolutionary lineages of mammals. Scientific Reports, 2019, 9, 10924.	1.6	10
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49	Ptychomitrium subcrispatum ThÃ©r. & P.de la Varde, an east southern African species excluded from the Cape Verde bryoflora. Journal of Bryology, 2019, 41, 281-284.	0.4	1
50	Molecular Characterization of HIV-1 Minority Subtypes in Hong Kong: A Recent Epidemic of CRF07_BC among the Men who have Sex with Men Population. Current HIV Research, 2019, 17, 53-64.	0.2	5
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53	Elucidating Viral Communities During a Phytoplankton Bloom on the West Antarctic Peninsula. Frontiers in Microbiology, 2019, 10, 1014.	1.5	28
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84	Boomeranging around Australia: Historical biogeography and population genomics of the antiâ€equatorial fish Microcanthus strigatus (Teleostei: Microcanthidae). Molecular Ecology, 2019, 28, 3771-3785.	2.0	17
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120	<p>Two new species of the Phimochirus holthuisi complex from the Gulf of Mexico, supported by morphology, color, and genetics (Crustacea: Anomura:) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 2		
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