

# Bayesian Phylogenetic Analysis of Combined Data

Systematic Biology

53, 47-67

DOI: [10.1080/10635150490264699](https://doi.org/10.1080/10635150490264699)

Citation Report

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| 831 | <i>Nucleospora cyclopteri</i> n. sp., an intranuclear microsporidian infecting wild lumpfish, <i>Cyclopterus lumpus</i> L., in Icelandic waters. Parasites and Vectors, 2013, 6, 49.   | 1.0 | 32        |
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| 838 | The evolution of host associations in the parasitic wasp genus <i>Ichneumon</i> (Hymenoptera: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 582 Td   | 3.2 | 24        |
| 839 | Application of plastid and nuclear markers to DNA barcoding of Euro-Mediterranean oaks ( <i>Quercus</i> , Fagaceae): problems, prospects and phylogenetic implications. <i>Botanical Journal of the Linnean Society</i> , 2013, 172, 478-499. | 0.8 | 80        |
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| 862 | North American Transmission of Hemosporidian Parasites in the Swainson's Thrush ( <i>Catharus</i> )  | 0.8 | 19        |
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| 884 | Empirical evaluation of partitioning schemes for phylogenetic analyses of mitogenomic data: An avian case study. <i>Molecular Phylogenetics and Evolution</i> , 2013, 66, 69-79.   | 1.2 | 55        |
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| 895 | Molecular phylogenetics and phylogeographic structure of <i>Sumichrasti</i> 's harvest mouse ( <i>Reithrodontomys sumichrasti</i> : Cricetidae) based on mitochondrial and nuclear DNA sequences. <i>Molecular Phylogenetics and Evolution</i> , 2013, 68, 282-292. | 1.2 | 22        |
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| 897 | The first complete mitochondrial genome from <i>Bostrychus</i> genus ( <i>Bostrychus sinensis</i> ) and partitioned Bayesian analysis of Eleotridae fish phylogeny. <i>Journal of Genetics</i> , 2013, 92, 247-257.   | 0.4 | 22        |
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| 917 | A revision of Afrotropical <i>Quasimodo</i> flies (Diptera: Schizophora; Curtonotidae). Part IV—the continental Afrotropical species of <i>Curtonotum</i> Macquart, with descriptions of thirteen new species and a combined phylogenetic analysis of the Curtonotidae. <i>Zootaxa</i> , 2013, 3684, 1-166. | 0.2 | 3         |
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| 930 | Ochratoxin production and taxonomy of the yellow aspergilli ( <i>Aspergillus</i> section) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 502 Td (   | 4.5 | 117       |
| 931 | Fleas of Small Mammals on Reunion Island: Diversity, Distribution and Epidemiological Consequences. <i>PLoS Neglected Tropical Diseases</i> , 2014, 8, e3129.   | 1.3 | 23        |
| 932 | Beyond fossil calibrations: realities of molecular clock practices in evolutionary biology. <i>Frontiers in Genetics</i> , 2014, 5, 138.  | 1.1 | 124       |
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| 1178 | One species or four? Yes!...and, no. Or, arbitrary assignment of lineages to species obscures the diversification processes of Neotropical fishes. <i>PLoS ONE</i> , 2017, 12, e0172349.  | 1.1 | 38        |
| 1179 | Daily activity patterns influence retinal morphology, signatures of selection, and spectral tuning of opsin genes in colubrid snakes. <i>BMC Evolutionary Biology</i> , 2017, 17, 249.  | 3.2 | 23        |
| 1180 | Phylogenetics of subtribe Orchidinae s.l. (Orchidaceae; Orchidoideae) based on seven markers (plastid) <i>Tj ETQq0 0 0 rgBT /Overlock 10 Plant Biology</i> , 2017, 17, 222.   | 1.6 | 41        |
| 1181 | Polyphyly of the traditional family Flabellinidae affects a major group of Nudibranchia: aeolidacean taxonomic reassessment with descriptions of several new families, genera, and species (Mollusca,) <i>Tj ETQq1 1 0.784314 rgBT /Overlock</i>  |     |           |
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| 1202 | Prisutnost pijavice <i>Pontobdella muricata</i> (Hirudinea: Piscicolidae) na hrskavi Anja Āama u Tirenskom moru (Srednje Sredozemlje). <i>Acta Adriatica</i> , 2018, 58, 225-234.  | 0.2 | 4         |
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| 1204 | <i>Bysmatrum austrafum</i> sp. nov. (Dinophyceae), a novel tidal pool dinoflagellate from South Africa. <i>Phycologia</i> , 2018, 57, 169-178.   | 0.6 | 5         |
| 1205 | <i>Betaphycus gelatinus</i> and <i>B. philippinensis</i> (Gigartinales, Rhodophyta) are conspecific. <i>Phytotaxa</i> , 2018, 372, 22.   | 0.1 | 5         |
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| 1210 | Land Plant Molecular Phylogenetics: A Review with Comments on Evaluating Incongruence Among Phylogenies. <i>Critical Reviews in Plant Sciences</i> , 2018, 37, 113-127.  | 2.7 | 37        |
| 1211 | Studies of <i>Myxidium giardi</i> C p de, 1906 infections in Icelandic eels identifies a genetically diverse clade of myxosporeans that represents the <i>Paramyxidium</i> n. g. (Myxosporea: Myxidiidae). <i>Parasites and Vectors</i> , 2018, 11, 551. | 1.0 | 7         |

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| 1213 | Re-establishment of <i>Spodoptera teferii</i> Laporte Rougeot (Lepidoptera: Noctuidae.) <i>Tj ETQq1 1 0.784314 rgBT /Overlo</i><br><i>La Societe Entomologique De France</i> , 2018, 54, 497-510.  | 0.4  | 7         |
| 1214 | Big data analysis of human mitochondrial DNA substitution models: a regression approach. <i>BMC Genomics</i> , 2018, 19, 759.  | 1.2  | 5         |
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| 1226 | Nuclear genes, matK and the phylogeny of the Poales. <i>Taxon</i> , 2018, 67, 521-536.   | 0.4  | 20        |
| 1227 | Evaluating Model Performance in Evolutionary Biology. <i>Annual Review of Ecology, Evolution, and Systematics</i> , 2018, 49, 95-114.  | 3.8  | 39        |
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| 1234 | <i>Helicascus alatus</i> (Morosphaeriaceae), a new freshwater species from southwestern China. <i>Phytotaxa</i> , 2018, 351, 210.  | 0.1 | 3         |
| 1235 | <i>Plagiodinium ballux</i> sp. nov. (Dinophyceae), a deep (36‰) sand dwelling dinoflagellate from subtropical Japan. <i>Phycological Research</i> , 2019, 67, 12-20.   | 0.8 | 2         |
| 1236 | The complete mitochondrial genome of the Yunnan red-backed vole <i>Eothenomys miletus</i> (Rodentia: Cricetidae) and its phylogeny. <i>Mitochondrial DNA Part B: Resources</i> , 2019, 4, 1424-1425.   | 0.2 | 3         |
| 1237 | A Systematist's Guide to Estimating Bayesian Phylogenies From Morphological Data. <i>Insect Systematics and Diversity</i> , 2019, 3, 2.  | 0.7 | 28        |
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| 1241 | Taxonomy, phylogeny and biogeography of francolins: Galliformes, Phasianidae, Phasianinae, Gallini; <i>Francolinus</i> , <i>Ortygornis</i> , <i>Afrocolinus</i> gen. nov. <i>Peliperdix</i> and <i>Scleroptila</i> spp.. <i>Ostrich</i> , 2019, 90, 191-221. | 0.4 | 10        |
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| 1249 | Phylogeny of Anthospermeae of the Coffee Family Inferred Using Clock and Nonclock Models. <i>International Journal of Plant Sciences</i> , 2019, 180, 386-402.   | 0.6 | 7         |
| 1250 | A new section and a new species of <i>Alternaria</i> encountered from Oman. <i>Phytotaxa</i> , 2019, 405, 279.   | 0.1 | 20        |
| 1251 | Supergene validation: A model-based protocol for assessing the accuracy of non-model-based supergene methods. <i>MethodsX</i> , 2019, 6, 2181-2188.  | 0.7 | 1         |
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| 1256 | Taxonomy, phylogeny and biogeography of African spurfowls Galliformes, Phasianidae, Phasianinae, Coturnicini: <i>Pternistis</i> spp.. <i>Östrich</i> , 2019, 90, 145-172.  | 0.4 | 12        |
| 1257 | Two new species of freshwater crabs of the genus <i>Potamonautes</i> MacLeay, 1838 (Decapoda: Brachyura: Tj ETQq1 1 0.784314 rgBT / 2019, 39, 426-435.   | 0.3 | 8         |
| 1258 | Allopatric separation represents an overlooked cryptic species in the <i>Anania hortulata</i> species complex (Lepidoptera: Crambidae: Pyraustinae): congruence between genetic and morphological evidence. <i>Canadian Entomologist</i> , 2019, 151, 163-186. | 0.4 | 4         |
| 1259 | Phylogenetic classification and generic delineation of <i>Hydeomyces deserti</i> sp. nov., (Phaeosphaeriaceae) from Jebel Akhdar Mountain in Oman. <i>Phytotaxa</i> , 2019, 391, 28.   | 0.1 | 12        |
| 1260 | Phylogeny and morphology of <i>Phillipsia hydei</i> sp. nov. (Sarcoscyphaceae) from Thailand. <i>Phytotaxa</i> , 2019, 395, 277.   | 0.1 | 1         |
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| 1266 | Updated phylogenetic and systematics of the <i>Acrapex albivena</i> Hampson, 1910 and <i>A. stygiata</i> (Hampson, 1910) species groups (Lepidoptera, Noctuidae, Noctuinae, Apameini, Sesamiina), with the description of nine new species from the Afrotropics. <i>Annales De La Societe Entomologique De France</i> , 2019, 55, 219-248. | 0.4 | 4         |
| 1267 | A molecular phylogeny of chafers revisits the polyphyly of Tanyproctini (Scarabaeidae.) <i>Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 70</i>  | 0.7 | 12        |
| 1268 | Molecular-based assessments of tribal and generic limits and relationships in Rubiaceae (Gentianales): Polyphyly of Pomazoteae and paraphyly of Ophiorrhizeae and <i>Ophiorrhiza</i> . <i>Taxon</i> , 2019, 68, 72-91.   | 0.4 | 12        |
| 1269 | Validation and phylogenetic placement of the Placentophoraceae <i>fam. nov.</i> (Gigartinales.) <i>Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 70</i>  | 0.6 | 4         |
| 1270 | <i>Prunus sunhangii</i> : A new species of <i>Prunus</i> from central China. <i>Plant Diversity</i> , 2019, 41, 19-25.   | 1.8 | 4         |
| 1271 | Large-scale mitochondrial gene rearrangements in the hermit crab <i>Pagurus nigrofascia</i> and phylogenetic analysis of the Anomura. <i>Gene</i> , 2019, 695, 75-83.  | 1.0 | 21        |
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| 1273 | The extraordinary genus <i>Myja</i> is not a tergipedid, but related to the Facelinidae s. str. with the addition of two new species from Japan (Mollusca, Nudibranchia). <i>ZooKeys</i> , 2019, 818, 89-116.  | 0.5 | 16        |
| 1274 | One stop shop II: taxonomic update with molecular phylogeny for important phytopathogenic genera: 26 (2019). <i>Fungal Diversity</i> , 2019, 94, 41-129.   | 4.7 | 69        |
| 1275 | <i>Lobocriconema iranense</i> (Van den Berg, Eskandari, Tiedt & Karegar, 2010) n. comb. and description of <i>L. nokandense</i> n. sp. (Nematoda: Criconematidae) from Iran. <i>Nematology</i> , 2019, 21, 1043-1061.  | 0.2 | 8         |
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| 1277 | CuPhylo: A CUDA Based Application Program Interface and Library for Phylogenetic Analysis. , 2019, , .   |     | 0         |
| 1278 | Analyses of Plastome Sequences Improve Phylogenetic Resolution and Provide New Insight Into the Evolutionary History of Asian Sonerileae/Dissochaeteae. <i>Frontiers in Plant Science</i> , 2019, 10, 1477.  | 1.7 | 26        |
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| 1283 | Nuclear protein phylogenies support the monophyly of the three bryophyte groups (Bryophyta) <i>Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 70</i>  | 3.5 | 84        |



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| 1284 | Phylogeny of Mesoamerican freshwater mussels and a revised tribe-level classification of the Ambleminae. <i>Zoologica Scripta</i> , 2019, 48, 106-117.  | 0.7 | 15        |
| 1285 | Homoplasy-Based Partitioning Outperforms Alternatives in Bayesian Analysis of Discrete Morphological Data. <i>Systematic Biology</i> , 2019, 68, 657-671.   | 2.7 | 31        |
| 1286 | Phylogeny of hymenolepidid cestodes (Cestoda: Cyclophyllidea) from mammalian hosts based on partial 28S rDNA, with focus on parasites from shrews. <i>Parasitology Research</i> , 2019, 118, 73-88.   | 0.6 | 13        |
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