

Sequence-Based Species Delimitation for the DNA Taxo

Systematic Biology

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Citation Report

#	ARTICLE	IF	CITATIONS
1	The evolutionary nature of diversification in sexuals and asexuals. , 2001, , 29-45.		6
2	Temporal patterns in diversification rates. , 2001, , 278-300.		16
3	Theoretical outlook. , 2001, , 345-373.		0
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5	Rapid and convergent evolution of parental care in hydrobiid gastropods from New Zealand. Journal of Evolutionary Biology, 2005, 18, 1076-1086.	0.8	33
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7	Phylogeny of Mysis (Crustacea, Mysida): history of continental invasions inferred from molecular and morphological data. Cladistics, 2005, 21, 575-596.	1.5	39
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22	Does the DNA barcoding gap exist? â€” a case study in blue butterflies (Lepidoptera: Lycaenidae). <i>Frontiers in Zoology</i> , 2007, 4, 8.	0.9	405
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39	The parasitic polychaete known as <i>Asetocalamyzas laoncola</i> (Calamyzidae) is in fact the dwarf male of the spionid <i>Scolelepis laoncola</i> (comb. nov.). <i>Invertebrate Biology</i> , 2008, 127, 403-416.	0.3	42
40	Distribution and Host Range of the Microsporidian <i>Pleistophora mulleri</i> . <i>Journal of Eukaryotic Microbiology</i> , 2008, 55, 355-362.	0.8	14
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284	Goldilocks Meets Santa Rosalia: An Ephemeral Speciation Model Explains Patterns of Diversification Across Time Scales. <i>Evolutionary Biology</i> , 2012, 39, 255-261.	0.5	195
285	Evolving Concepts of Bacterial Species. <i>Evolutionary Biology</i> , 2012, 39, 148-157.	0.5	33
286	An integrative morphological and molecular diagnostics for <i>Typhlodromus pyri</i> (Acari: Tj ETQq1 1 0.784314 rgBT /Overlock 10	0.7	24
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295	Identification of molecular markers for DNA barcoding in the Aphidiinae (Hym. Braconidae). Molecular Ecology Resources, 2012, 12, 197-208.	2.2	62
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298	Lichen myco- and photobiont diversity and their relationships at the edge of life (McMurdo Dry) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50	1.3	66
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312	Speciation and dietary specialization in <i>Drupa</i> , a genus of predatory marine snails (Gastropoda: Tj ETQq1 1 0,784314,rgBT /Over	0.7	16
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326	Functional rare males in diploid parthenogenetic <i>Asteremia</i> . <i>Journal of Evolutionary Biology</i> , 2013, 26, 1934-1948.	0.8	28
327	Bayesian species delimitation reveals generalist and specialist parasitic wasps on <i>Galerucella</i> beetles (Chrysomelidae): sorting by herbivore or plant host. <i>BMC Evolutionary Biology</i> , 2013, 13, 92.	3.2	43
328	Mitochondrial lineage sorting in action – historical biogeography of the <i>Hyles euphorbiae</i> complex (Sphingidae, Lepidoptera) in Italy. <i>BMC Evolutionary Biology</i> , 2013, 13, 83.	3.2	28

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330	Evolution of microgastropods (Ellobioidea, Carychiidae): integrating taxonomic, phylogenetic and evolutionary hypotheses. <i>BMC Evolutionary Biology</i> , 2013, 13, 18.	3.2	54
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341	Integration of cytochrome <i>c</i> oxidase I barcodes and geometric morphometrics to delimit species in the genus <i>Gnopharmia</i> (Lepidoptera: Geometridae, Ennominae). <i>Zoological Journal of the Linnean Society</i> , 2013, 169, 70-83.	1.0	9
342	Evolutionary history of nematodes associated with sweat bees. <i>Molecular Phylogenetics and Evolution</i> , 2013, 66, 847-856.	1.2	14
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350	Uncovering historical signature of mitochondrial DNA hidden in the nuclear genome: the biogeography of <i>Schistocerca</i> revisited. <i>Cladistics</i> , 2013, 29, 643-662.	1.5	16
351	Holarctic phylogeography of the testate amoeba <i>Halosphenia papilio</i> (Amoebozoa: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 667) limitation. <i>Molecular Ecology</i> , 2013, 22, 5172-5184.	2.0	67
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355	Ancient lineage, young troglobites: recent colonization of caves by <i>Nesticella</i> spiders. <i>BMC Evolutionary Biology</i> , 2013, 13, 183.	3.2	32
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363	Molecular phylogeny and historical biogeography of the Anatolian lizard <i>Apathya</i> (<i>Squamata</i> , Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 667) 1.2 56	1.2	56
364	Nonlinear projection methods for visualizing Barcode data and application on two data sets. <i>Molecular Ecology Resources</i> , 2013, 13, 976-990.	2.2	6

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366	An exploration of species boundaries in turret-building tarantulas of the Mojave Desert (Araneae). <i>Tj ETQq1 1 0.784314 rgBT /Overl</i> 327-340.	1.2	68
367	A phylogenetic analysis of relationships among genera of Conopidae (Diptera) based on molecular and morphological data. <i>Cladistics</i> , 2013, 29, 193-226.	1.5	8
368	Assessing model fit in phylogeographical investigations: an example from the North American sandbar willow <i>Saxifraga melanopsis</i> . <i>Journal of Biogeography</i> , 2013, 40, 131-141.	1.4	35
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371	Timetree of Aselloidea Reveals Species Diversification Dynamics in Groundwater. <i>Systematic Biology</i> , 2013, 62, 512-522.	2.7	55
372	Endemism and diversification in freshwater insects of Madagascar revealed by coalescent and phylogenetic analysis of museum and field collections. <i>Molecular Phylogenetics and Evolution</i> , 2013, 66, 979-991.	1.2	50
373	Two apparently unrelated groups of symbiotic annelids, Nautiliniellidae and Calamyzidae (Phyllodocida, Annelida), are a clade of derived chrysopetalid polychaetes. <i>Cladistics</i> , 2013, 29, 610-628.	1.5	18
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376	Integrative taxonomy and conservation of cryptic beetles in the Mediterranean region (Hydrophilidae). <i>Zoologica Scripta</i> , 2013, 42, 182-200.	0.7	34
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378	Species Diversity, Phylogeny and Large Scale Biogeographic Patterns of the Genus <i>Padina</i> (Phaeophyceae, Dictyotales). <i>Journal of Phycology</i> , 2013, 49, 130-142.	1.0	53
379	Genetic testing reveals some mislabeling but general compliance with a ban on herbivorous fish harvesting in Belize. <i>Conservation Letters</i> , 2013, 6, 132-140.	2.8	35
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381	DNA barcoding species inventory of Microgastrinae wasps (Hymenoptera, Braconidae) from a Mexican tropical dry forest. <i>Molecular Ecology Resources</i> , 2013, 13, 1146-1150.	2.2	18
382	Whole-community DNA barcoding reveals a spatio-temporal continuum of biodiversity at species and genetic levels. <i>Nature Communications</i> , 2013, 4, 1892.	5.8	71

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384	Integrative analysis of DNA phylogeography and morphology of the European rose chafer (<i>Cetonia</i>) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 507 Td (Phylogenetics and Evolution, 2013, 69, 83-94.	1.2	22
385	A skeleton-less sponge of Caribbean mangroves: invasive or undescribed?. <i>Invertebrate Biology</i> , 2013, 132, 81-94.	0.3	4
386	The peculiar nemertean larva <i>pilidium recurvatum</i> belongs to <i>Riserius</i> sp., a basal heteronemertean that eats <i>Carcinonemertes errans</i> , a hoplonemertean parasite of Dungeness crab. <i>Invertebrate Biology</i> , 2013, 132, 207-225.	0.3	14
387	Beta diversity at multiple hierarchical levels: explaining the high diversity of scarab beetles in tropical montane forests. <i>Journal of Biogeography</i> , 2013, 40, 2134-2145.	1.4	18
388	A rapid diagnostic technique of <i>Bactrocera cucurbitae</i> and <i>Bactrocera zonata</i> (Diptera:) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 507 Td (Phylogenetic relationships of <i>Hymenodryas</i> (<i>Nymphalidae</i>) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 507 Td (Cladistics, 2013, 29, 629-642.	1.7	10
389	Phylogenetic relationships of <i>Hymenodryas</i> (<i>Nymphalidae</i>) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 507 Td (Cladistics, 2013, 29, 629-642.	1.5	9
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391	Molecular phylogeny of the genus <i>Margotrema</i> (Digenea: Allocreadiidae), parasitic flatworms of goodeid freshwater fishes across central Mexico: species boundaries, host-specificity, and geographical congruence. <i>Zoological Journal of the Linnean Society</i> , 2013, 168, 1-16.	1.0	19
392	Genetic differentiation of the African dwarf crocodile <i>Osteolaemus tetraspis</i> Cope, 1861 (Crocodylia) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 507 Td (255-266.	0.7	12
393	Species inflation and taxonomic artefacts – A critical comment on recent trends in mammalian classification. <i>Mammalian Biology</i> , 2013, 78, 1-6.	0.8	161
394	Reliable, verifiable and efficient monitoring of biodiversity via metabarcoding. <i>Ecology Letters</i> , 2013, 16, 1245-1257.	3.0	514
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396	Forms of water mites (Acari: Hydrachnidia): intraspecific variation or valid species?. <i>Ecology and Evolution</i> , 2013, 3, 3415-3435.	0.8	35
397	<i>Bombus cullumanus</i> – an extinct European bumblebee species?. <i>Apidologie</i> , 2013, 44, 121-132.	0.9	20
398	Species diversity and reticulate evolution in the <i>Asplenium normale</i> complex (Aspleniaceae) in China and adjacent areas. <i>Taxon</i> , 2013, 62, 673-687.	0.4	40
399	Taxonomic revision of Madagascan <i>Rhantus</i> (Coleoptera, Dytiscidae, Colymbetinae) with an emphasis on <i>Manjakatampo</i> as a conservation priority. <i>ZooKeys</i> , 2013, 350, 21-45.	0.5	10
400	Integrative taxonomy and preliminary assessment of species limits in the <i>Liolaemus walkeri</i> complex (Squamata, Liolaemidae) with descriptions of three new species from Peru. <i>ZooKeys</i> , 2013, 364, 47-91.	0.5	40

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402	Extensive cryptic species diversity and fine-scale endemism in the marine red alga <i>Portieria</i> in the Philippines. Proceedings of the Royal Society B: Biological Sciences, 2013, 280, 20122660.	1.2	93
403	Thermal tolerance breadths among groundwater crustaceans living in a thermally constant environment. Journal of Experimental Biology, 2013, 216, 1683-94.	0.8	38
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672	Phylogeny of the symphytan grade of Hymenoptera: new pieces into the old jigsaw(fly) puzzle. Cladistics, 2015, 31, 1-17.	1.5	108

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674	Capacity of Japanese <i>A. sobara</i> species (Hymenoptera; Braconidae) to parasitize a fruit pest <i>Drosophila suzukii</i> (Diptera; Drosophilidae). <i>Journal of Applied Entomology</i> , 2015, 139, 105-113.	0.8	46
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681	Community assembly on remote islands: a comparison of Hawaiian and Mascarene spiders. <i>Journal of Biogeography</i> , 2015, 42, 39-50.	1.4	16
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893	<i>Torix</i> group <i>Rickettsia</i> are widespread in <i>Culicoides</i> biting midges (Diptera: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 587 Td (Microbiology, 2017, 19, 4238-4255.	1.8	41
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904	Morphological and molecular divergence between <i>Crenidens crenidens</i> (Forsskl, 1775) and <i>C. indicus</i> Day, 1873 (Perciformes: Sparidae) and notes on a Red Sea endemic lineage of <i>C. crenidens</i> . <i>Marine Biodiversity</i> , 2017, 47, 1273-1285.	0.3	6
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916	Toward a methodical framework for comprehensively assessing forest multifunctionality. <i>Ecology and Evolution</i> , 2017, 7, 10652-10674.	0.8	41
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927	Examining metrics and magnitudes of molecular genetic differentiation used to delimit cetacean subspecies based on mitochondrial <i>cytb</i> control region sequences. <i>Marine Mammal Science</i> , 2017, 33, 76-100.	0.9	28
928	Evolutionary history of <i>Podarcis tiliguerta</i> on Corsica and Sardinia. <i>BMC Evolutionary Biology</i> , 2017, 17, 27.	3.2	6

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934	BarcodingR: an integrated <i>scpr</i> package for species identification using <i>DNA</i> barcodes. <i>Methods in Ecology and Evolution</i> , 2017, 8, 627-634.	2.2	38
935	From Alaska to Antarctica: Species boundaries and genetic diversity of <i>Prasiola</i> (Trebouxiophyceae), a foliose chlorophyte associated with the bipolar lichen-forming fungus <i>Mastodia tessellata</i> . <i>Molecular Phylogenetics and Evolution</i> , 2017, 107, 117-131.	1.2	57
936	Molecular and morphological delimitation of Australian Triops species (Crustacea: Branchiopoda: Tj ETQq1 1 0.784314 rgBT /Overlock 0.7 10	0.7	10
937	Sympatric lineage divergence in cryptic Neotropical sweat bees (Hymenoptera: Halictidae: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 422 Td 0.7 4	0.7	4
938	Predation by generalist arthropod predators on <i>Apolygus lucorum</i> (Hemiptera: Miridae): molecular gut content analysis and field cage assessment. <i>Pest Management Science</i> , 2017, 73, 628-635.	1.7	17
939	<i>Copromyxa laresi</i> n. sp. (Amoebozoa: Tubulinea) and Transfer of <i>Cashia limacoides</i> (Page,) Tj ETQq0 0 0 rgBT /Overlock 10 5 0.8	0.8	5
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951	Diversity and specificity of sap-feeding herbivores and their parasitoids on Australian fig trees. <i>Insect Conservation and Diversity</i> , 2017, 10, 107-119.	1.4	4
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981	CLADES: A classification-based machine learning method for species delimitation from population genetic data. <i>Molecular Ecology Resources</i> , 2018, 18, 1144-1156.	2.2	13
982	More diverse than expected: distributional patterns of <i>Oecidiobanchus</i> Hessler, 1970 (Isopoda, Tj ETQq1 1 0.784314 rgBT /Overlock 10	0.3	10

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1031	Selected results of DNA-based species identification on animal foods. <i>Journal of the Science of Food and Agriculture</i> , 2018, 98, 2437-2439.	1.7	0
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1051	Species delimitation, global phylogeny and historical biogeography of the parasitoid wasp genus <i>Spathius</i> (Braconidae: Doryctinae) reveal multiple Oligocene-Miocene intercontinental dispersal events. <i>Zoological Journal of the Linnean Society</i> , 2018, 182, 723-734.	1.0	7
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1053	Concordance between DNA-based species boundaries and reproductive isolating barriers in the <i>Scytosiphon lomentaria</i> species complex (Ectocarpales, Phaeophyceae). <i>Phycologia</i> , 2018, 57, 232-242.	0.6	13
1054	DNA barcoding a nightmare taxon: assessing barcode index numbers and barcode gaps for sweat bees. <i>Genome</i> , 2018, 61, 21-31.	0.9	54

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1055	Contrasting phylogeographic pattern among <i>Eudyptes</i> penguins around the Southern Ocean. <i>Scientific Reports</i> , 2018, 8, 17481.	1.6	29
1056	Evidence supporting cryptic species within two sessile microinvertebrates, <i>Limnias melicerta</i> and <i>L. ceratophylli</i> (Rotifera, Gnesiotrocha). <i>PLoS ONE</i> , 2018, 13, e0205203.	1.1	12
1057	Species identification and connectivity of marine amphipods in Canada's three oceans. <i>PLoS ONE</i> , 2018, 13, e0197174.	1.1	22
1058	Molecular and anatomical analyses reveal that <i>Peronia verruculata</i> (Gastropoda: Onchidiidae) is a cryptic species complex. <i>Contributions To Zoology</i> , 2018, 87, 149-165.	0.2	10
1059	Inferring species boundaries using acoustic and morphological data in the ground cricket genus <i>Gymnogryllus</i> (Orthoptera: Grylloidea: Gryllinae). <i>Systematics and Biodiversity</i> , 2018, 16, 731-742.	0.5	4
1061	<i>Betaphycus gelatinus</i> and <i>B. philippinensis</i> (Gigartinales, Rhodophyta) are conspecific. <i>Phytotaxa</i> , 2018, 372, 22.	0.1	5
1062	Speciation in a biodiversity hotspot: Phylogenetic relationships, species delimitation, and divergence times of Patagonian ground frogs from the <i>Eupsophus roseus</i> group (Alsodidae). <i>PLoS ONE</i> , 2018, 13, e0204968.	1.1	9
1063	Integrative Taxonomy and Its Implications for Species-Level Systematics of Parasitoid Hymenoptera. <i>Entomological Review</i> , 2018, 98, 834-864.	0.1	15
1064	Integrative taxonomy reveals hidden species within a common fungal parasite of ladybirds. <i>Scientific Reports</i> , 2018, 8, 15966.	1.6	52
1065	Molecular phylogeny and systematics of the centipede genus <i>Ethmostigmus</i> Pocock (Chilopoda: Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 0.5	0.5	10
1066	Integrative taxonomic methods reveal an incorrect synonymisation of the South African <i>Pseudonereis podocirra</i> (Schmarda) as the widespread <i>Pseudonereis variegata</i> (Grube) from Chile. <i>Invertebrate Systematics</i> , 2018, 32, 1282.	0.5	14
1067	Compensatory gastric stretching following subtotal gastric resection due to gastric adenocarcinoma in a diamond python (<scp><i>Morelia spilota spilota</i></scp>). <i>Australian Veterinary Journal</i> , 2018, 96, 481-486.	0.5	3
1068	A re-description of <i>Cyrtodactylus chrysopylos</i> Bauer (Squamata: Gekkonidae) with comments on the adaptive significance of orange coloration in hatchlings and descriptions of two new species from eastern Myanmar (Burma). <i>Zootaxa</i> , 2018, 4527, 151.	0.2	16
1069	Genetic diversity in the camellia weevil, <i>Curculio chinensis</i> Chevrolat (Coleoptera: Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 0.3	0.3	10
1070	Delimiting species of marine gastropods (Turridae, Conoidea) using <scp>RAD</scp> sequencing in an integrative taxonomy framework. <i>Molecular Ecology</i> , 2018, 27, 4591-4611.	2.0	20
1071	Hierarchies of evolutionary radiation in the world's most species rich vertebrate group, the Neotropical <i>Pristimantis</i> leaf litter frogs. <i>Systematics and Biodiversity</i> , 2018, 16, 807-819.	0.5	17
1072	Integrating coalescent species delimitation with analysis of host specificity reveals extensive cryptic diversity despite minimal mitochondrial divergence in the malaria parasite genus <i>Leucocytozoon</i> . <i>BMC Evolutionary Biology</i> , 2018, 18, 128.	3.2	49
1073	Utility of molecular-assisted alpha taxonomy of the genus <i>Cystophora</i> (Fucales, Phaeophyceae) from New Zealand and Australia. <i>Phycologia</i> , 2018, 57, 374-384.	0.6	2

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1075	Reverse taxonomy applied to the <i>Brachionus calyciflorus</i> cryptic species complex: Morphometric analysis confirms species delimitations revealed by molecular phylogenetic analysis and allows the (re)description of four species. <i>PLoS ONE</i> , 2018, 13, e0203168.	1.1	66
1076	DNA barcode sheds light on systematics and evolution of neotropical freshwater trahiras. <i>Genetica</i> , 2018, 146, 505-515.	0.5	11
1077	A two-step DNA barcoding approach for delimiting moth species: moths of Dongling Mountain (Beijing, China). <i>Tj ETQq1 1 0.784314 16 BT / Overl</i>	1.6	16
1078	A new species of Crocodile Newt <i>Tylototriton</i> (Caudata: Salamandridae) from Shan State, Myanmar (Burma). <i>Zootaxa</i> , 2018, 4500, 553-573.	0.2	13
1079	Species delimitation at a global scale reveals high species richness with complex biogeography and patterns of symbiont association in <i>Peltigera</i> section <i>Peltigera</i> (lichenized Ascomycota). <i>Tj ETQq1 1 0.784314 16 BT / Overl</i>	1.6	16
1080	Evaluation of six regions for their potential as <i>DNA</i> barcodes in epiphyllous liverworts from Thailand. <i>Applications in Plant Sciences</i> , 2018, 6, e011174.	0.8	4
1081	Two new species of <i>Atlantoscia</i> Ferrara & Taiti, 1981 (Isopoda: Oniscidea: Philosciidae) from southern Brazil described in the light of integrative taxonomy. <i>Zootaxa</i> , 2018, 4482, 551-565.	0.2	9
1082	The evolutionary history of Mediterranean Batoidea (Chondrichthyes: Neoselachii). <i>Zoologica Scripta</i> , 2018, 47, 686-698.	0.7	12
1083	Two more new species of <i>Hemiphyllodactylus</i> Bleeker (Squamata: Gekkonidae) from the Shan Hills of eastern Myanmar (Burma). <i>Zootaxa</i> , 2018, 4483, 295-316.	0.2	16
1084	Host specificity of parasitoids (Encyrtidae) toward armored scale insects (Diaspididae): Untangling the effect of cryptic species on quantitative food webs. <i>Ecology and Evolution</i> , 2018, 8, 7879-7893.	0.8	10
1085	Phylogeography of the Red Algal <i>Laurencia</i> Complex in the Macaronesia Region and Nearby Coastal Areas: Recent Advances and Future Perspectives. <i>Diversity</i> , 2018, 10, 10.	0.7	10
1086	The <i>Lordiphosa denticeps</i> species group (Diptera: Drosophilidae) in China, with redescriptions of four known species and descriptions of nine new species. <i>Zootaxa</i> , 2018, 4471, 37.	0.2	23
1087	Combining molecular and landscape tools for targeting evolutionary processes in reserve design: An approach for islands. <i>PLoS ONE</i> , 2018, 13, e0200830.	1.1	7
1088	Integrative species delimitation in practice: Revealing cryptic lineages within the short-nosed skink <i>Plestiodon brevirostris</i> (Squamata: Scincidae). <i>Molecular Phylogenetics and Evolution</i> , 2018, 129, 242-257.	1.2	17
1089	Detection of sister-species in invasive populations of the fall armyworm <i>Spodoptera frugiperda</i> (Lepidoptera: Noctuidae) from Uganda. <i>PLoS ONE</i> , 2018, 13, e0194571.	1.1	82
1090	Instant taxonomy: choosing adequate characters for species delimitation and description through congruence between molecular data and quantitative shape analysis. <i>Invertebrate Systematics</i> , 2018, 32, 551.	0.5	20
1091	Genetic diversity of marine oligochaetous clitellates in selected areas of the South Atlantic as revealed by DNA barcoding. <i>Invertebrate Systematics</i> , 2018, 32, 524.	0.5	8

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1093	How many names for a beloved genus? â€œ Coalescent-based species delimitation in <i>Xanthium</i> L. (<i>Ambrosiinae</i> , <i>Asteraceae</i>). <i>Molecular Phylogenetics and Evolution</i> , 2018, 127, 135-145.	1.2	24
1094	Paleoceneâ€Eocene and Plioceneâ€Pleistocene sea-level changes as â€œspecies pumpsâ€ in Southeast Asia: Evidence from <i>Altheopus</i> spiders. <i>Molecular Phylogenetics and Evolution</i> , 2018, 127, 545-555.	1.2	33
1095	A hyper-diverse genus of acanthocephalans revealed by tree-based and non-tree-based species delimitation methods: Ten cryptic species of <i>Neoechinorhynchus</i> in Middle American freshwater fishes. <i>Molecular Phylogenetics and Evolution</i> , 2018, 127, 30-45.	1.2	43
1096	Unexpected species diversity within Japanese <i>Mundochthonius</i> pseudoscorpions (<i>Pseudoscorpiones</i>): Tj ETQq0 0 0 rgBT /Overlock 10 T morphological examination. <i>Invertebrate Systematics</i> , 2018, 32, 259.	0.5	14
1097	Plant Species Complexes as Models to Understand Speciation and Evolution: A Review of South American Studies. <i>Critical Reviews in Plant Sciences</i> , 2018, 37, 54-80.	2.7	60
1098	First genetic data for band-winged grasshoppers (<i>Orthoptera</i> : <i>Acrididae</i> : <i>Oedipodinae</i>) of the Biskra region of Algeria with new records for the country. <i>African Zoology</i> , 2018, 53, 31-40.	0.2	7
1099	Exploring the diversity of Asian <i>Cryptocercus</i> (<i>Blattodea</i> : <i>Cryptocercidae</i>): species delimitation based on chromosome numbers, morphology and molecular analysis. <i>Invertebrate Systematics</i> , 2018, 32, 69.	0.5	16
1100	One thousand DNA barcodes of piranhas and pacus reveal geographic structure and unrecognized diversity in the Amazon. <i>Scientific Reports</i> , 2018, 8, 8387.	1.6	47
1101	Taxonomic revision and insights into the speciation mode of the spider <i>Dysdera erythrina</i> species-complex (<i>Araneae</i> : <i>Dysderidae</i>): sibling species with sympatric distributions. <i>Invertebrate Systematics</i> , 2018, 32, 10.	0.5	17
1102	The hitchhiker's guide to single-locus species delimitation. <i>Molecular Ecology Resources</i> , 2018, 18, 1234-1246.	2.2	131
1103	A mega-cryptic species complex hidden among one of the most common annelids in the North East Atlantic. <i>PLoS ONE</i> , 2018, 13, e0198356.	1.1	63
1104	A global analysis of bats using automated comparative phylogeography uncovers a surprising impact of Pleistocene glaciation. <i>Journal of Biogeography</i> , 2018, 45, 1795-1805.	1.4	26
1105	Taxonomic revision of the Square-tailed Drongo species complex (<i>Passeriformes</i> : <i>Dicruridae</i>) with description of a new species from western Africa. <i>Zootaxa</i> , 2018, 4438, 105.	0.2	8
1106	An integrative species delimitation approach reveals fine-scale endemism and substantial unrecognized avian diversity in the Philippine Archipelago. <i>Conservation Genetics</i> , 2018, 19, 1153-1168.	0.8	16
1107	Molecular Phylogenetic Analysis and Species Delimitation in the Pine Needle-feeding Aphid Genus <i>Essigella</i> (<i>Hemiptera</i> , <i>Sternorrhyncha</i> , <i>Aphididae</i>). <i>Insect Systematics and Diversity</i> , 2018, 2, .	0.7	3
1108	Cryptic diversity, low connectivity and suspected human-mediated dispersal among 17 widespread Indo-Pacific hydroid species of the south-western Indian Ocean. <i>Journal of Biogeography</i> , 2018, 45, 2104-2117.	1.4	19
1109	Exploring the Udoteaceae diversity (<i>Bryopsidales</i> , <i>Chlorophyta</i>) in the Caribbean region based on molecular and morphological data. <i>Molecular Phylogenetics and Evolution</i> , 2018, 127, 758-769.	1.2	8

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1110	Molecular and morphological data of the freshwater fish <i>Glandulocauda melanopleura</i> (Characiformes: Characidae) provide evidences of river captures and local differentiation in the Brazilian Atlantic Forest. <i>PLoS ONE</i> , 2018, 13, e0194247.	1.1	19
1111	Diversity of the green algal genus <i>Spirogyra</i> (Conjugatophyceae) in the Hawaiian Islands. <i>Phycologia</i> , 2018, 57, 331-344.	0.6	7
1112	More Than Meets the Eye: Cryptic Diversity and Contrasting Patterns of Host-Specificity in Feather Mites Inhabiting Seabirds. <i>Frontiers in Ecology and Evolution</i> , 2018, 6, .	1.1	14
1113	Molecular association and morphological characterisation of <i>Himalopsyche</i> larval types (Trichoptera, Rhyacophilidae). <i>ZooKeys</i> , 2018, 773, 79-108.	0.5	7
1114	Speciation patterns in complex subterranean environments: a case study using short-tailed whipscorpions (Schizomida: Hubbardiidae). <i>Biological Journal of the Linnean Society</i> , 2018, 125, 355-367.	0.7	16
1115	Phylogeny and taxonomic reassessment of jerboa, <i>Dipus</i> (Rodentia, Dipodinae), in inland Asia. <i>Zoologica Scripta</i> , 2018, 47, 630-644.	0.7	4
1116	Patterns and drivers of species diversity in the Indo-Pacific red seaweed <i>Portieria</i> . <i>Journal of Biogeography</i> , 2018, 45, 2299-2313.	1.4	46
1117	The effects of contemporary selection and dispersal limitation on the community assembly of acidophilic microalgae. <i>Journal of Phycology</i> , 2018, 54, 720-733.	1.0	18
1118	Molecular Evidences of a Hidden Complex Scenario in <i>Leporinus cf. friderici</i> . <i>Frontiers in Genetics</i> , 2018, 9, 47.	1.1	17
1119	Genetic Variation of the Endangered Neotropical Catfish <i>Steindachneridion scriptum</i> (Siluriformes: Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf_50 342 Td	1.1	10
1120	Genetic and Morphological Analyses Demonstrate That <i>Schizolecis guntheri</i> (Siluriformes: Tj ETQq0 0 0 rgBT /Overlock 10 Tf_50 342 Td	1.1	5
1121	Little Divergence Among Mitochondrial Lineages of <i>Prochilodus</i> (Teleostei, Characiformes). <i>Frontiers in Genetics</i> , 2018, 9, 107.	1.1	22
1122	Species Delimitation, Phylogenetic Relationships, and Temporal Divergence Model in the Genus <i>Aeromonas</i> . <i>Frontiers in Microbiology</i> , 2018, 9, 770.	1.5	6
1123	Cross-Contamination Explains Inter and Intraspecific Horizontal Genetic Transfers between Asexual Bdelloid Rotifers. <i>Current Biology</i> , 2018, 28, 2436-2444.e14.	1.8	30
1124	DNA Barcoding Highlights Cryptic Diversity in the New Zealand Psylloidea (Hemiptera: Tj ETQq0 0 0 rgBT /Overlock 10 Tf_50 182 Td (St	0.7	26
1125	Isolation drives increased diversification rates in freshwater amphipods. <i>Molecular Phylogenetics and Evolution</i> , 2018, 127, 746-757.	1.2	17
1126	Reappraisal of the systematics of <i>Microglanis cottoides</i> (Siluriformes, Pseudopimelodidae), a catfish from southern Brazil. <i>PLoS ONE</i> , 2018, 13, e0199963.	1.1	4
1127	Species delimitation and mitogenome phylogenetics in the subterranean genus <i>Pseudoniphargus</i> (Crustacea: Amphipoda). <i>Molecular Phylogenetics and Evolution</i> , 2018, 127, 988-999.	1.2	25

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1129	<i>Lyrodus mersinensis</i> sp. nov. (Bivalvia: Teredinidae) another cryptic species in the <i>Lyrodus pedicellatus</i> (Quatrefages, 1849) complex. <i>Zootaxa</i> , 2018, 4442, 441-457.	0.2	12
1130	Taxonomy of <i>Cladonia angustiloba</i> and related species. <i>Lichenologist</i> , 2018, 50, 267-282.	0.5	15
1131	A new <i>Bunodophoron</i> species (<i>Sphaerophoraceae</i> , <i>Lecanorales</i>) from the Neotropics. <i>Lichenologist</i> , 2018, 50, 255-266.	0.5	6
1132	A genomic evaluation of taxonomic trends through time in coast horned lizards (genus <i>Phrynosoma</i>)	2.0	14
1133	Molecular and morphometric analyses reveal cryptic diversity within freshwater mussels (Bivalvia: <i>Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf</i>) <i>Society</i> , 2018, 124, 261-277.	0.7	30
1134	Biodiversity estimates and ecological interpretations of meiofaunal communities are biased by the taxonomic approach. <i>Communications Biology</i> , 2018, 1, 112.	2.0	28
1135	Phylogenetic systematics of the shelled sea slug genus <i>Oxynoe</i> Rafinesque, 1814 (Heterobranchia :)	0.5	12
1136	A Multi-Gene Analysis and Potential Spatial Distribution of Species of the Strodei Subgroup of the Genus <i>Nyssorhynchus</i> (Diptera: Culicidae). <i>Journal of Medical Entomology</i> , 2018, 55, 1486-1495.	0.9	4
1137	Exceptional biodiversity of the cryptofaunal decapods in the Chagos Archipelago, central Indian Ocean. <i>Marine Pollution Bulletin</i> , 2018, 135, 636-647.	2.3	7
1138	Evolutionary History. , 2018, , 45-75.		14
1139	Identity and genetic structure of eggplant fruit and shoot borer, <i>Leucinodes orbonalis</i> Guenée (Lepidoptera:Crambidae) populations in the Philippines inferred from morphological traits and COI sequence data. <i>Journal of Asia-Pacific Entomology</i> , 2018, 21, 1009-1019.	0.4	4
1140	Monotypic status of the South American relictual marsupial <i>Dromiciops gliroides</i> (Microbiotheria). <i>Journal of Mammalogy</i> , 2018, 99, 803-812.	0.6	26
1141	Genetics of lineage diversification and the evolution of host usage in the economically important wheat curl mite, <i>Aceria tosichella</i> Keifer, 1969. <i>BMC Evolutionary Biology</i> , 2018, 18, 122.	3.2	25
1142	DNA barcode based delineation of freshwater fishes from northern Western Ghats of India, one of the world's biodiversity hotspots. <i>Biodiversity and Conservation</i> , 2018, 27, 3349-3371.	1.2	15
1143	Why the COI barcode should be the community <i>scp>DNA</scp></i> metabarcode for the metazoa. <i>Molecular Ecology</i> , 2018, 27, 3968-3975.	2.0	131
1144	The Pleurobemini (Bivalvia : Unionida) revisited: molecular species delineation using a mitochondrial DNA gene reveals multiple conspecifics and undescribed species. <i>Invertebrate Systematics</i> , 2018, 32, 689.	0.5	21
1145	DNA Barcoding of Freshwater Fishes of Indo-Myanmar Biodiversity Hotspot. <i>Scientific Reports</i> , 2018, 8, 8579.	1.6	33

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1147	Wide distributions and cryptic diversity within a <i>Microstomum</i> (Platyhelminthes) species complex. Zoologica Scripta, 2018, 47, 486-498.	0.7	7
1148	Detection of Hermetia illucens by real-time PCR. Journal of Insects As Food and Feed, 2018, 4, 115-122.	2.1	16
1149	Prevalence of cryptic species in morphologically uniform taxa "Fast speciation and evolutionary radiation in Asian frogs. Molecular Phylogenetics and Evolution, 2018, 127, 723-731.	1.2	45
1150	Mitochondrial Divergence between Western and Eastern Great Bustards: Implications for Conservation and Species Status. Journal of Heredity, 2018, 109, 641-652.	1.0	5
1151	How to achieve internal fertilization without a vagina: the study case of the genus Archilina Ax, 1959 (Platyhelminthes, Proseriata) from Canary Islands. Marine Biodiversity, 2019, 49, 2057-2073.	0.3	16
1152	Taxonomy of the long-tailed mouse <i>Oligoryzomys destructor</i> (Sigmodontinae: Oryzomyini) with the designation of neotypes for <i>Hesperomys destructor</i> Tschudi, 1844 and <i>Hesperomys melanostoma</i> Tschudi, 1844. Journal of Zoological Systematics and Evolutionary Research, 2019, 57, 127-144.	0.6	11
1153	First molecular phylogeny of <i>Agrilus</i> (Coleoptera: Buprestidae), the largest genus on Earth, with DNA barcode database for forestry pest diagnostics. Bulletin of Entomological Research, 2019, 109, 200-211.	0.5	32
1154	Molecular species delimitation of the Asian chestnut gall wasp biocontrol agent released in Italy. Insect Systematics and Evolution, 2019, 50, 327-345.	0.2	9
1155	DNA Barcoding: Bioinformatics Workflows for Beginners. , 2019, , 985-995.		5
1156	A comparative evolutionary study reveals radically different scales of genetic structuring within two atyid shrimp species (Crustacea: Decapoda: Atyidae). Zoological Journal of the Linnean Society, 2019, 186, 200-212.	1.0	5
1157	A comprehensive molecular phylogeny of tiger beetles (Coleoptera, Carabidae, Cicindelinae). Systematic Entomology, 2019, 44, 305-321.	1.7	31
1158	Molecular phylogeny of the scorpionflies Panorpidae (Insecta: Mecoptera) and chromosomal evolution. Cladistics, 2019, 35, 385-400.	1.5	31
1159	Species diversity in Gymnogeophagus (Teleostei: Cichlidae) and comparative biogeography of cichlids in the Middle Paran� basin, an emerging hotspot of fish endemism. Hydrobiologia, 2019, 832, 331-354.	1.0	15
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1161	Phylogenetic relationships and biogeographic history of the Australian trapdoor spider genus Conothele (Araneae: Mygalomorphae: Halonoproctidae): diversification into arid habitats in an otherwise tropical radiation. Invertebrate Systematics, 2019, , .	0.5	10
1162	Phylogenetic relationships within the flatworm genus Matuxia (Platyhelminthes, Tricladida,) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 107 T genus. Organisms Diversity and Evolution, 2019, 19, 377-390.	0.7	0
1163	Living on the edge: Assessing the diversity of South African Pocillopora on the margins of the Southwestern Indian Ocean. PLoS ONE, 2019, 14, e0220477.	1.1	4

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1164	Geographic structure of genetic variation in the Parachute Gecko <i>Ptychozoon lionotum</i> Annandale, 1905 across Indochina and Sundaland with descriptions of three new species. <i>Zootaxa</i> , 2019, 4638, 151-198.	0.2	9
1165	Is reproductive strategy a key factor in understanding the evolutionary history of Southern Ocean Asteroidea (Echinodermata)? <i>Ecology and Evolution</i> , 2019, 9, 8465-8478.	0.8	14
1166	The Prevalence of Single-Specimen/Locality Species in Insect Taxonomy: An Empirical Analysis. <i>Diversity</i> , 2019, 11, 106.	0.7	16
1167	Fungal species boundaries in the genomics era. <i>Fungal Genetics and Biology</i> , 2019, 131, 103249.	0.9	66
1168	Phylogeny, divergence times and species delimitation of <i>Tonicia</i> (Polyplacophora: Chitonidae) from the eastern Pacific Ocean. <i>Zoological Journal of the Linnean Society</i> , 2019, 186, 915-933.	1.0	9
1169	Species composition, diversity, and distribution of the genus <i>Ulva</i> along the coast of Jeju Island, Korea based on molecular phylogenetic analysis. <i>PLoS ONE</i> , 2019, 14, e0219958.	1.1	23
1170	Genomic data reveal deep genetic structure but no support for current taxonomic designation in a grasshopper species complex. <i>Molecular Ecology</i> , 2019, 28, 3869-3886.	2.0	23
1171	Molecular and morphological recognition of species boundaries in the neglected ant genus <i>Brachymyrmex</i> (Hymenoptera: Formicidae): toward a taxonomic revision. <i>Organisms Diversity and Evolution</i> , 2019, 19, 447-542.	0.7	15
1172	Systematics, biogeography, and evolution of <i>Pristurus minimus</i> (Squamata, Sphaerodactylidae) with the discovery of the smallest Arabian vertebrate. <i>Systematics and Biodiversity</i> , 2019, 17, 349-366.	0.5	7
1173	The Caribbean enigma: the presence of unusual cryptic diversity in intertidal mites (Arachnida, Acari). <i>Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 18</i>	0.7	13
1174	Efficient Identification of <i>Pulsatilla</i> (Ranunculaceae) Using DNA Barcodes and Micro-Morphological Characters. <i>Frontiers in Plant Science</i> , 2019, 10, 1196.	1.7	18
1175	Historical museum collections clarify the evolutionary history of cryptic species radiation in the world's largest amphibians. <i>Ecology and Evolution</i> , 2019, 9, 10070-10084.	0.8	36
1176	Clonal diversity and substantial genetic divergence of the <i>Daphnia similis</i> species complex in Chinese lakes: Possible adaptations to the uplift of the Qinghai-Tibetan Plateau. <i>Limnology and Oceanography</i> , 2019, 64, 2725-2737.	1.6	8
1177	Arid zone island hopping: the impact of dispersal on endemism in hydraenid beetles (Coleoptera:). <i>Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 18</i>	0.8	4
1178	Molecular species delimitation of reef-building coral genera, <i>Porites</i> and <i>Turbinaria</i> (Anthozoa): diversity. <i>Systematics and Biodiversity</i> , 2019, 17, 541-557.	0.5	4
1179	DNA barcodes expose unexpected diversity in Canadian mites. <i>Molecular Ecology</i> , 2019, 28, 5347-5359.	2.0	40
1180	Cryptic species and co-diversification in sand scorpions from the Karakum and Kyzylkum deserts of Central Asia. <i>Zoologica Scripta</i> , 2019, 48, 801-812.	0.7	11
1181	Molecular phylogeny and species delimitation of the genus <i>Dicerapanorpa</i> (Mecoptera: Panorpidae). <i>Zoological Journal of the Linnean Society</i> , 2019, 187, 1173-1195.	1.0	8

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1182	Diversification in the mountains: a generic reappraisal of the Western Ghats endemic gecko genus <i>Dravidogecko</i> Smith, 1933 (Squamata: Gekkonidae) with descriptions of six new species. <i>Zootaxa</i> , 2019, 4688, zootaxa.4688.1.1.	0.2	20
1183	Evaluating methodologies for species delimitation: the mismatch between phenotypes and genotypes in lichenized fungi (<i>Bryoria</i> sect. <i>Implexae</i> , <i>Parmeliaceae</i>). <i>Persoonia: Molecular Phylogeny and Evolution of Fungi</i> , 2019, 42, 75-100.	1.6	44
1184	DNA Barcoding Reveals High Levels of Divergence among Mitochondrial Lineages of Brycon (Characiformes, Bryconidae). <i>Genes</i> , 2019, 10, 639.	1.0	9
1185	Do mosses really exhibit so large distribution ranges? Insights from the integrative taxonomic study of the <i>Lewinskya affinis</i> complex (Orthotrichaceae, Bryopsida). <i>Molecular Phylogenetics and Evolution</i> , 2019, 140, 106598.	1.2	23
1186	Integrating patterns of thermal tolerance and phenotypic plasticity with population genetics to improve understanding of vulnerability to warming in a widespread copepod. <i>Global Change Biology</i> , 2019, 25, 4147-4164.	4.2	49
1187	Morphology and molecules reveal two new species of <i>Porites</i> (Scleractinia, Poritidae) from the Red Sea and the Gulf of Aden. <i>Systematics and Biodiversity</i> , 2019, 17, 491-508.	0.5	12
1188	Phylogenetic analysis of the <i>Baikalodrilus</i> species flock (Annelida: Clitellata: Naididae), an endemic genus to Lake Baikal (Russia). <i>Zoological Journal of the Linnean Society</i> , 2019, 187, 987-1015.	1.0	5
1189	Multilocus approach reveals a complex evolutionary history of the invasive mile-a-minute plant, <i>Mikania micrantha</i> (Asteraceae), in its natural habitat. <i>Botanical Journal of the Linnean Society</i> , 2019, 191, 188-215.	0.8	1
1190	Unexpected species diversity in electric eels with a description of the strongest living bioelectricity generator. <i>Nature Communications</i> , 2019, 10, 4000.	5.8	45
1191	The <i>Hirtodrosophila melanderi</i> species group (Diptera: Drosophilidae) from the Huanglong National Nature Reserve, Sichuan, China. <i>Zootaxa</i> , 2019, 4623, zootaxa.4623.1.7.	0.2	1
1192	Genetic and chromatic variation of <i>Coprophanaeus</i> (<i>Megaphanaeus</i>) <i>ensifer</i> (Germar, 1821) (Coleoptera: Scarabaeidae). <i>Zoologischer Anzeiger</i> , 2019, 283, 150-160.	0.4	6
1193	Cryptic Diversity and Database Errors Challenge Non-indigenous Species Surveys: An Illustration With <i>Botrylloides</i> spp. in the English Channel and Mediterranean Sea. <i>Frontiers in Marine Science</i> , 2019, 6, .	1.2	44
1194	<i>Ophelimus migdanorum</i> Molina-Mercader sp. nov. (Hymenoptera: Eulophidae): Application of Integrative Taxonomy for Disentangling a Polyphenism Case in <i>Eucalyptus globulus</i> Labill Forest in Chile. <i>Forests</i> , 2019, 10, 720.	0.9	8
1195	Species Delimitation and Analysis of Cryptic Species Diversity in the XXI Century. <i>Entomological Review</i> , 2019, 99, 463-472.	0.1	28
1196	Diversity and distribution of Ischnomesidae (Crustacea: Isopoda: Asellota) along the Kuril-Kamchatka Trench – A genetic perspective. <i>Progress in Oceanography</i> , 2019, 178, 102174.	1.5	19
1197	Chromosome number diversity in Asian <i>Cryptocercus</i> (Blattodea, Cryptocercidae) and implications for karyotype evolution and geographic distribution on the Western Sichuan Plateau. <i>Systematics and Biodiversity</i> , 2019, 17, 594-608.	0.5	4
1198	Emergence of sympatry in a radiation of subterranean amphipods. <i>Journal of Biogeography</i> , 2019, 46, 657-669.	1.4	19
1199	An integrative taxonomic and phylogenetic approach reveals a complex of cryptic species in the “peppermint” shrimp <i>Lysmata wurdemanni sensu stricto</i> . <i>Zoological Journal of the Linnean Society</i> , 2019, 185, 1018-1038.	1.0	7

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1200	Diversification of bent-toed geckos (<i>Cyrtodactylus</i>) on Sumatra and west Java. <i>Molecular Phylogenetics and Evolution</i> , 2019, 134, 1-11.	1.2	18
1201	Assessing eukaryotic biodiversity in the Florida Keys National Marine Sanctuary through environmental DNA metabarcoding. <i>Ecology and Evolution</i> , 2019, 9, 1029-1040.	0.8	69
1202	A new <i>Epeorus</i> (Caucasiron) (Ephemeroptera: Heptageniidae) species from Turkey based on molecular and morphological evidence. <i>Zootaxa</i> , 2019, 4550, 58-70.	0.2	13
1203	Multiplex real-time PCR for the detection of insect DNA and determination of contents of <i>Tenebrio molitor</i> , <i>Locusta migratoria</i> and <i>Achaeta domestica</i> in food. <i>European Food Research and Technology</i> , 2019, 245, 559-567.	1.6	17
1204	A molecular approach to the identification of marine fish of the Dongsha Islands (South China Sea). <i>Fisheries Research</i> , 2019, 213, 105-112.	0.9	17
1205	Phylogeny and molecular species delimitation of long-nosed armadillos (<i>Dasybus: Cingulata</i>) supports morphology-based taxonomy. <i>Zoological Journal of the Linnean Society</i> , 2019, 186, 813-825.	1.0	27
1206	Using multilocus approach to uncover cryptic diversity within <i>Pseudotrapelus</i> lizards from Saudi Arabia. <i>Saudi Journal of Biological Sciences</i> , 2019, 26, 1442-1449.	1.8	7
1207	An assessment of species limits of the South American mouse genus <i>Oligoryzomys</i> (Rodentia, Tj ETQq1 1 0.784314,rgBT /Oyerlock 10 0.7	0.7	10
1208	Establishment of six new <i>Rhabdoblatta</i> species (Blattodea, Blaberidae, Epilamprinae) from China. <i>ZooKeys</i> , 2019, 851, 27-69.	0.5	10
1209	Diversity and relationships of <i>Ampedini Gistel, 1848</i> (Coleoptera : Elateridae) in Switzerland and Europe. <i>Invertebrate Systematics</i> , 2019, , .	0.5	2
1210	Phylogeography of <i>Simulium</i> Subgenus <i>Wilhelmia</i> (Diptera: Simuliidae)â€™Insights From Balkan Populations. <i>Journal of Medical Entomology</i> , 2019, 56, 967-978.	0.9	8
1211	For common community phylogenetic analyses, go ahead and use synthesis phylogenies. <i>Ecology</i> , 2019, 100, e02788.	1.5	80
1212	Testing and using complete plastomes and ribosomal DNA sequences as the next generation DNA barcodes in <i>Panax</i> (Araliaceae). <i>Molecular Ecology Resources</i> , 2019, 19, 1333-1345.	2.2	45
1213	Molecular phylogeny and historical biogeography of <i>Parnara</i> butterflies (Lepidoptera: Hesperidae). <i>Molecular Phylogenetics and Evolution</i> , 2019, 139, 106545.	1.2	5
1214	Untangling a mess of worms: Species delimitations reveal morphological crypsis and variability in Southeast Asian semi-aquatic earthworms (Almidae, Glyphidrilus). <i>Molecular Phylogenetics and Evolution</i> , 2019, 139, 106531.	1.2	12
1215	Does DNA barcoding offer meaningful insights into the diversity of the parrotfish of the genus <i>Sparisoma</i> (Scaridae)?. <i>Journal of Applied Ichthyology</i> , 2019, 35, 1029.	0.3	2
1216	Speciation across mountains: Phylogenomics, species delimitation and taxonomy of the <i>Liolaemus leopardinus</i> clade (Squamata, Liolaemidae). <i>Molecular Phylogenetics and Evolution</i> , 2019, 139, 106524.	1.2	28
1217	Species delimitation of neotropical Characins (Stevardiinae): Implications for taxonomy of complex groups. <i>PLoS ONE</i> , 2019, 14, e0216786.	1.1	31

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1218	Extensive regional endemism and cryptic diversity in the Tennessee and Kentucky, USA populations of the burrowing crayfish <i>Cambarus deweesae</i> (Bouchard & Etnier, 1979) (Decapoda: Astacidea). <i>Tj ETQq0 0 0 rgBT/Overlock 10 Tf 5</i>		
1219	Studies of Laboulbeniales on <i>Myrmica</i> ants (IV): host-related diversity and thallus distribution patterns of <i>Rickia wasmannii</i> . <i>Parasite</i> , 2019, 26, 29.	0.8	10
1220	A taxonomic review and revisions of Microstomidae (Platyhelminthes: Macrostomorpha). <i>PLoS ONE</i> , 2019, 14, e0212073.	1.1	11
1221	Coalescent-based species delimitation in the sand lizards of the <i>Liolaemus wiegmanni</i> complex (Squamata: Liolaemidae). <i>Molecular Phylogenetics and Evolution</i> , 2019, 138, 89-101.	1.2	16
1222	Towards a global DNA barcode reference library for quarantine identifications of lepidopteran stemborers, with an emphasis on sugarcane pests. <i>Scientific Reports</i> , 2019, 9, 7039.	1.6	16
1223	The roles of morphological traits, resource variation and resource partitioning associated with the dietary niche expansion in the fish-eating bat <i>Myotis pilosus</i> . <i>Molecular Ecology</i> , 2019, 28, 2944-2954.	2.0	22
1224	Ancient landscapes of the Namib Desert harbor high levels of genetic variability and deeply divergent lineages for Collembola. <i>Ecology and Evolution</i> , 2019, 9, 4969-4979.	0.8	10
1225	Phylogenetics and species delimitations of the operculated land snail <i>Cyclophorus volvulus</i> (Gastropoda: Cyclophoridae) reveal cryptic diversity and new species in Thailand. <i>Scientific Reports</i> , 2019, 9, 7041.	1.6	21
1226	Taxonomic and systematic implications of the revision of the phylogenetic relations in the genus <i>Ectinogonia</i> Spinola 1837 (Coleoptera: Buprestidae) of Central Chile. <i>Zootaxa</i> , 2019, 4603, 159.	0.2	3
1227	Hidden species diversity in <i>Pachyhynobius</i> : A multiple approaches species delimitation with mitogenomes. <i>Molecular Phylogenetics and Evolution</i> , 2019, 137, 138-145.	1.2	18
1228	Revision of <i>Bairdiella</i> (Sciaenidae: Perciformes) from the western South Atlantic, with insights into its diversity and biogeography. <i>Neotropical Ichthyology</i> , 2019, 17, .	0.5	11
1229	A new species of crocodile newt <i>Tylototriton</i> (Caudata: Salamandridae) from northern Myanmar (Burma). <i>Journal of Natural History</i> , 2019, 53, 475-495.	0.2	10
1230	DNA barcoding uncovers extensive cryptic diversity in the African long-fin tetra <i>Bryconalestes longipinnis</i> (Alestidae: Characiformes). <i>Journal of Fish Biology</i> , 2019, 95, 379-392.	0.7	12
1231	Species diversity of <i>Marmosa</i> subgenus <i>Micoureus</i> (Didelphimorphia, Didelphidae) and taxonomic evaluation of the white-bellied woolly mouse opossum, <i>Marmosa constantiae</i> . <i>Zoological Journal of the Linnean Society</i> , 2019, 187, 240-277.	1.0	27
1232	A taxonomic reassessment of <i>Caulerpa</i> (Chlorophyta, Caulerpaceae) in southern Australia, based on <i>tuf</i> A and <i>rbc</i> L sequence data. <i>Phycologia</i> , 2019, 58, 234-253.	0.6	10
1233	Categorization of species as native or nonnative using DNA sequence signatures without a complete reference library. <i>Ecological Applications</i> , 2019, 29, e01914.	1.8	14
1234	Biogeography of <i>Mesalina</i> (Reptilia: Lacertidae), with special emphasis on the <i>Mesalina adramitana</i> group from Arabia and the Socotra Archipelago. <i>Molecular Phylogenetics and Evolution</i> , 2019, 137, 300-312.	1.2	12
1235	Species limits in butterflies (Lepidoptera: Nymphalidae): reconciling classical taxonomy with the multispecies coalescent. <i>Systematic Entomology</i> , 2019, 44, 745-756.	1.7	23

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1237	Complete subspecies-level phylogeny of the Oriolidae (Aves: Passeriformes): Out of Australasia and return. <i>Molecular Phylogenetics and Evolution</i> , 2019, 137, 200-209.	1.2	18
1238	Singleton molecular species delimitation based on COI-5P barcode sequences revealed high cryptic/undescribed diversity for Chinese katydids (Orthoptera: Tettigoniidae). <i>BMC Evolutionary Biology</i> , 2019, 19, 79.	3.2	24
1239	The systematics of Lobophora (Dictyotales, Phaeophyceae) in the western Atlantic and eastern Pacific oceans: eight new species. <i>Journal of Phycology</i> , 2019, 55, 611-624.	1.0	21
1240	The roles of geography, climate and sexual selection in driving divergence among insect populations on mountaintops. <i>Journal of Biogeography</i> , 2019, 46, 784-795.	1.4	12
1241	Genetic and morphological variations of the lichenized fungus <i>Steinera intricata</i> (Arctomiaceae, Tj ETQq1 1 0.784314 rgBT /Overlock 0 907-918.	0.5	0
1242	The Klingon batbugs: Morphological adaptations in the primitive bat bugs, <i>Bucimex chilensis</i> and <i>Primicimex cavernis</i> , including updated phylogeny of Cimicidae. <i>Ecology and Evolution</i> , 2019, 9, 1736-1749.	0.8	13
1243	First records of the parthenogenetic Surinam cockroach <i>Pycnoscelus surinamensis</i> (Insecta: Tj ETQq1 1 0.784314 rgBT /Overlock 0 4	0.8	4
1244	Demographic Expansion of the Predominant Bemisia tabaci (Gennadius) (Hemiptera: Aleyrodidae) Mitotypes Associated With the Cotton Leaf Curl Virus Epidemic in Pakistan. <i>Annals of the Entomological Society of America</i> , 2019, 112, 265-280.	1.3	19
1245	Cryptic Diversity, but to What Extent? Discordance Between Single-Locus Species Delimitation Methods Within Mainland Anoles (Squamata: Dactyloidae) of Northern Central America. <i>Frontiers in Genetics</i> , 2019, 10, 11.	1.1	34
1246	Species delimitation and multi-locus species tree solve an old taxonomic problem for European squat lobsters of the genus Munida Leach, 1820. <i>Marine Biodiversity</i> , 2019, 49, 1751-1773.	0.3	16
1247	Molecular discrimination of <i>Ancistrus</i> lineages (Siluriformes: Loricariidae) using barcode DNA tool. <i>Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis</i> , 2019, 30, 602-608.	0.7	6
1248	Dark offshoot: Phylogenomic data sheds light on the evolutionary history of a new species of cave brittle star. <i>Molecular Phylogenetics and Evolution</i> , 2019, 136, 151-163.	1.2	4
1249	Description of ten new Lobophora species from the Bismarck Sea (Papua New Guinea). <i>Phycological Research</i> , 2019, 67, 228-238.	0.8	11
1250	Two more new species of the Cyrtodactylus peguensis group (Squamata: Gekkonidae) from the fringes of the Ayeyarwady Basin, Myanmar. <i>Zootaxa</i> , 2019, 4577, 274.	0.2	11
1251	DNA barcodes and their characteristic diagnostic sites analysis of Schizothoracinae fishes in Qinghai province. <i>Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis</i> , 2019, 30, 592-601.	0.7	2
1252	Phylogenetic relationships of rock-inhabiting black fungi belonging to the widespread genera <i>Lichenothelia</i> and <i>Saxomyces</i> . <i>Mycologia</i> , 2019, 111, 127-160.	0.8	13
1253	Relationship between contrasting morphotypes and the phylogeny of the marine gastropod genus <i>Tegula</i> (Vetigastropoda: Tegulidae) in East Asia. <i>Journal of Molluscan Studies</i> , 2019, 85, 24-34.	0.4	10

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1254	Incomplete estimates of genetic diversity within species: Implications for <sc>DNA</sc> barcoding. Ecology and Evolution, 2019, 9, 2996-3010.	0.8	72
1255	New lineages and old species: Lineage diversity and regional distribution of <i>Moina</i> (Crustacea: Tj ETQq1 1 0.784314 rgBT /Ovgrlock 10 Tf 50	1.2	19
1256	Six new species of the <i>Cyrtodactylus intermedius</i> complex (Squamata: Gekkonidae) from the Cardamom Mountains and associated highlands of Southeast Asia. Zootaxa, 2019, 4554, 1-62.	0.2	31
1257	Two new species of <i>Heptapterus</i> (Siluriformes: Heptapteridae) from the Uruguay River basin, Brazil. Journal of Fish Biology, 2019, 94, 352-373.	0.7	7
1258	Extensive Variations in Diurnal Growth Patterns and Metabolism Among <i>Ulva</i> spp. Strains. Plant Physiology, 2019, 180, 109-123.	2.3	37
1259	Revisiting species boundaries and distribution ranges of <i>Nemacheilus</i> spp. (Cypriniformes: Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 barcodes: implications for conservation in a biodiversity hotspot. Conservation Genetics, 2019, 20, 517-529.	0.8	17
1260	Fin whale (<i>Balaenoptera physalus</i>) mitogenomics: A cautionary tale of defining sub-species from mitochondrial sequence monophyly. Molecular Phylogenetics and Evolution, 2019, 135, 86-97.	1.2	11
1261	Multilocus phylogeny, species delimitation and biogeography of Iberian valvatiform springsnails (Caenogastropoda: Hydrobiidae), with the description of a new genus. Zoological Journal of the Linnean Society, 2019, 186, 892-914.	1.0	29
1262	Description of a new species of the Neotropical cichlid genus <i>Gymnogeophagus</i> Miranda Ribeiro, 1918 (Teleostei: Cichliformes) from the Middle Paran� basin, Misiones, Argentina. PLoS ONE, 2019, 14, e0210166.	1.1	3
1263	Cryptic diversity and dynamic chromosome evolution in Alpine scorpions (Euscorpiidae: Euscorpius). Molecular Phylogenetics and Evolution, 2019, 134, 152-163.	1.2	26
1264	Deep segregation in the open ocean: Macaronesia� as an evolutionary hotspot for low dispersal marine invertebrates. Molecular Ecology, 2019, 28, 1784-1800.	2.0	20
1265	Morphological Species of <i>Gloeandromyces</i> (Ascomycota, Laboulbeniales) Evaluated Using Single-locus Species Delimitation Methods. Fungal Systematics and Evolution, 2019, 3, 19-34.	0.9	15
1266	Systematic position and composition of <i>Merodon nigratarsis</i> and <i>M. avidus</i> groups (Diptera, Syrphidae) with a description of four new hoverflies species. Contributions To Zoology, 2019, 89, 74-125.	0.2	11
1267	A New Green Salamander in the Southern Appalachians: Evolutionary History of <i>Aneides aeneus</i> and Implications for Management and Conservation with the Description of a Cryptic Microendemic Species. Copeia, 2019, 107, 748.	1.4	10
1268	<p>Multi-instar descriptions of cave dwelling Erythraeidae (Trombidiformes: Parasitengona) employing an integrative approach</p>. Zootaxa, 2019, 4717, 137-184.	0.2	7
1269	Biochemical and molecular differentiation of <i>Anacroneuria</i> species (Plecoptera, Insecta) in Andean National Park, Venezuela. Systematics and Biodiversity, 2019, 17, 669-678.	0.5	2
1270	Move to stay: genetic structure and demographic history of a wolf spider inhabiting coastal sand dunes of southern South America. Systematics and Biodiversity, 2019, 17, 635-649.	0.5	2
1271	Phylogeography, Population Structure, and Species Delimitation in Rockhopper Penguins (<i>Eudyptes</i>) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50	1.0	3

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1272	BIN overlap confirms transcontinental distribution of pest aphids (Hemiptera: Aphididae). PLoS ONE, 2019, 14, e0220426.	1.1	2
1273	High diversity of coralline algae in New Zealand revealed: Knowledge gaps and implications for future research. PLoS ONE, 2019, 14, e0225645.	1.1	37
1274	Integrative systematics and ecology of a new deep-sea family of tanaidacean crustaceans. Scientific Reports, 2019, 9, 18720.	1.6	13
1275	Unravelling the phylogeny, cryptic diversity and morphological evolution of Diptilomiopus mites (Acari: Eriophyoidea). Experimental and Applied Acarology, 2019, 79, 323-344.	0.7	9
1276	Cryptic diversity in the North American Dromochorus tiger beetles (Coleoptera: Carabidae:). Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 587 To Society, 2019, 186, 250-285.	1.0	18
1277	Longitudinal monitoring of environmental factors at Culicidae larval habitats in urban areas and their association with various mosquito species using an innovative strategy. Pest Management Science, 2019, 75, 923-934.	1.7	6
1278	Game of clones: Is Wolbachia inducing speciation in a weevil with a mixed reproductive mode?. Molecular Phylogenetics and Evolution, 2019, 133, 42-53.	1.2	16
1279	Population genetic structure and species delimitation of a widespread, Neotropical dwarf gecko. Molecular Phylogenetics and Evolution, 2019, 133, 54-66.	1.2	29
1280	DNA barcoding fishes from the Congo and the Lower Guinean provinces: Assembling a reference library for poorly inventoried fauna. Molecular Ecology Resources, 2019, 19, 728-743.	2.2	19
1281	Species delimitation methods reveal cryptic diversity in the <i>Hypnea cornuta</i> complex (Cystocloniaceae, Rhodophyta). European Journal of Phycology, 2019, 54, 135-153.	0.9	28
1282	Integrating multilocus DNA data and 3D geometric morphometrics to elucidate species boundaries in the case of Pyrenaearia (Pulmonata: Hygromiidae). Molecular Phylogenetics and Evolution, 2019, 132, 194-206.	1.2	8
1283	DNA barcoding identification and genetic diversity of bamboo shoot wireworms (Coleoptera:). Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 587 To 0.4	0.4	6
1284	Genetic and morphological differentiation of <i>Porphyra</i> and <i>Pyropia</i> species (Bangiales,). Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 587 To 1.0	1.0	9
1285	Conspecificity of the model organism <i>Ulva mutabilis</i> and <i>Ulva compressa</i> (Ulvophyceae,). Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 587 To 1.0	1.0	36
1286	Real-time PCR assays for rapid detection of <i>Zeugodacus cucumis</i> and <i>Bactrocera jarvisi</i> (Diptera: Tephritidae) for quarantine application. Journal of Applied Entomology, 2019, 143, 155-163.	0.8	3
1287	An Antarctic flock under the Thorson's rule: Diversity and larval development of Antarctic Velutinidae (Mollusca: Gastropoda). Molecular Phylogenetics and Evolution, 2019, 132, 1-13.	1.2	13
1288	Phylogeography and Molecular Species Delimitation of <i>Pratylenchus capsici</i> n. sp., a New Root-Lesion Nematode in Israel on Pepper (<i>Capsicum annum</i>). Phytopathology, 2019, 109, 847-858.	1.1	22
1289	Persistence of phylogeographic footprints helps to understand cryptic diversity detected in two marine amphipods widespread in the Mediterranean basin. Molecular Phylogenetics and Evolution, 2019, 132, 53-66.	1.2	22

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1290	Intragenomic <i>internal transcribed spacer 2</i> variation in a genus of parasitoid wasps (Hymenoptera: Braconidae): implications for accurate species delimitation and phylogenetic analysis. <i>Insect Molecular Biology</i> , 2019, 28, 485-498.	1.0	4
1291	Cox1 barcoding versus multilocus species delimitation: validation of two mite species with contrasting effective population sizes. <i>Parasites and Vectors</i> , 2019, 12, 8.	1.0	25
1292	Unmasking Antarctic mollusc lineages: novel evidence from philinoid snails (Gastropoda: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 662 Td (1.5	12
1293	An integrative study of island diversification: Insights from the endemic <i>Haemodracon</i> geckos of the Socotra Archipelago. <i>Molecular Phylogenetics and Evolution</i> , 2019, 133, 166-175.	1.2	8
1294	Diversifying into the branches: Species boundaries in African green and bush snakes, <i>Philothamnus</i> (Serpentes: Colubridae). <i>Molecular Phylogenetics and Evolution</i> , 2019, 130, 357-365.	1.2	15
1295	Phylogeography of Atlantic Forest glassfrogs (<i>Vitreorana</i>): when geography, climate dynamics and rivers matter. <i>Heredity</i> , 2019, 122, 545-557.	1.2	21
1296	Sex-Biased Dispersal Obscures Species Boundaries in Integrative Species Delimitation Approaches. <i>Systematic Biology</i> , 2019, 68, 441-459.	2.7	21
1297	New insights into the <i>Usnea cornuta</i> aggregate (Parmeliaceae, lichenized Ascomycota): Molecular analysis reveals high genetic diversity correlated with chemistry. <i>Molecular Phylogenetics and Evolution</i> , 2019, 131, 125-137.	1.2	13
1298	Invisible apple snail invasions: importance of continued vigilance and rigorous taxonomic assessments. <i>Pest Management Science</i> , 2019, 75, 1277-1286.	1.7	15
1299	Unparalleled disjunction or unexpected relationships? Molecular phylogeny and biogeography of Melanopsidae (Caenogastropoda: Cerithioidea), with the description of a new family and a new genus from the ancient continent Zealandia. <i>Cladistics</i> , 2019, 35, 401-425.	1.5	9
1300	DNA barcoding of large oak-living cerambycids: diagnostic tool, phylogenetic insights and natural hybridization between <i>Cerambyx cerdo</i> and <i>Cerambyx welensii</i> (Coleoptera: Cerambycidae). <i>Bulletin of Entomological Research</i> , 2019, 109, 583-594.	0.5	16
1301	A polyphasic approach to the delimitation of diatom species: a case study for the genus <i>Pinnularia</i> (Bacillariophyta). <i>Journal of Phycology</i> , 2019, 55, 365-379.	1.0	15
1302	Cryptic diversity revealed in the leaf-toed gecko <i>Asaccus montanus</i> (Squamata, Phyllodactylidae) from the Hajar Mountains of Arabia. <i>Journal of Zoological Systematics and Evolutionary Research</i> , 2019, 57, 369-382.	0.6	9
1303	Species delimitation in Neotropical fishes of the genus <i>Characidium</i> (Teleostei, Characiformes). <i>Zoologica Scripta</i> , 2019, 48, 69-80.	0.7	22
1304	Phylogeographic structure in the chromosomally polymorphic rodent <i>Cricetulus barabensis</i> sensu lato (Mammalia, Cricetidae). <i>Journal of Zoological Systematics and Evolutionary Research</i> , 2019, 57, 679-694.	0.6	9
1305	New insights on the genus <i>Otoplana</i> Du Plessis, 1889 (Platyhelminthes: Proseriata), with description of two new species from the Canary Islands. <i>Marine Biodiversity</i> , 2019, 49, 2075-2087.	0.3	15
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1307	Dynamics of global institutional collaboration in insect taxonomy reveal imbalance of taxonomic effort. <i>Insect Conservation and Diversity</i> , 2019, 12, 18-28.	1.4	4

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1309	Phylogenetics and species delimitation of two hydrozoa (phylum Cnidaria): <i>Turritopsis</i> (McCrary, 1857) and <i>Pennaria</i> (Goldfuss, 1820). <i>Marine Biodiversity</i> , 2019, 49, 1085-1100.	0.3	23
1310	Morphometrical and molecular evidence suggests cryptic diversity among hookworms (Nematoda: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5 2020, 94, e8.	0.4	1
1311	Development of a <i>Yersinia entomophaga</i> bait for control of larvae of the porina moth (<i>Wiseana</i> spp.), a pest of New Zealand improved grassland systems. <i>Pest Management Science</i> , 2020, 76, 350-359.	1.7	5
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1313	Cryptic species delineation in freshwater planarians of the genus <i>Dugesia</i> (Platyhelminthes, Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 5 variability. <i>Molecular Phylogenetics and Evolution</i> , 2020, 143, 106496.	1.2	24
1314	Coinfection of the secondary symbionts, <i>Hamiltonella defensa</i> and <i>Arsenophonus</i> sp. contribute to the performance of the major aphid pest, <i>Aphis gossypii</i> (Hemiptera: Aphididae). <i>Insect Science</i> , 2020, 27, 86-98.	1.5	37
1315	Marine heat waves, climate change, and failed spawning by coastal invertebrates. <i>Limnology and Oceanography</i> , 2020, 65, 627-636.	1.6	30
1316	Tracking mite trophic interactions by multiplex PCR. <i>Pest Management Science</i> , 2020, 76, 597-608.	1.7	6
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1318	Genetic evidence for a species complex within the piranha <i>Serrasalmus maculatus</i> (Characiformes, Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5 and <i>Molecular Biology</i> , 2020, 43, e20190131.	0.6	4
1319	Survey of infection and determination of the transmission vector of <i>Onchocerca fasciata</i> in camels () Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 5	0.7	3
1320	Deer keds on wild ungulates in northern Italy, with a taxonomic key for the identification of <i>Lipoptena</i> spp. of Europe. <i>Medical and Veterinary Entomology</i> , 2020, 34, 74-85.	0.7	19
1321	Population structure and species delimitation of rice white tip nematode, <i>Aphelenchoides besseyi</i> (Nematoda: Aphelenchoididae), in China. <i>Plant Pathology</i> , 2020, 69, 159-167.	1.2	16
1322	A Comparative Characterization of the Mitochondrial Genomes of <i>Paramoeba</i> aparasomata and <i>Neoparamoeba pemaquidensis</i> (Amoebozoa, Paramoebidae). <i>Journal of Eukaryotic Microbiology</i> , 2020, 67, 167-175.	0.8	2
1323	<i>Pseudoparamoeba garorimi</i> n. sp., with Notes on Species Distinctions within the Genus. <i>Journal of Eukaryotic Microbiology</i> , 2020, 67, 132-139.	0.8	3
1324	Integrative species delimitation in Nearctic ambush bugs (Heteroptera: Reduviidae: Phymatinae): insights from molecules, geometric morphometrics and ecological associations. <i>Systematic Entomology</i> , 2020, 45, 205-223.	1.7	15
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1327	Exploring community assembly among Javanese and Balinese freshwater shrimps (<i>Atyidae</i> , Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 747 1	1.0	4
1328	Species delimitation based on integrative approach suggests reallocation of genus in Hypostomini catfish (Siluriformes, Loricariidae). <i>Hydrobiologia</i> , 2020, 847, 563-578.	1.0	16
1329	Molecular identification of forensically important fly species in Spain using COI barcodes. <i>Science and Justice - Journal of the Forensic Science Society</i> , 2020, 60, 293-302.	1.3	7
1330	Invasive lumbricid earthworms in North Americaâ€”Different life histories but common dispersal?. <i>Journal of Biogeography</i> , 2020, 47, 674-685.	1.4	7
1331	Integrative species delimitation of desmosomatid and nannoniscid isopods from the Kuril-Kamchatka trench, with description of a hadal species. <i>Progress in Oceanography</i> , 2020, 182, 102236.	1.5	15
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1333	Unraveling the diversification and systematic puzzle of the highly polymorphic <i>Psammobates tentorius</i> (Bell, 1828) complex (Reptilia: Testudinidae) through phylogenetic analyses and species delimitation approaches. <i>Journal of Zoological Systematics and Evolutionary Research</i> , 2020, 58, 308-326.	0.6	7
1334	Phylogeography, speciation and demographic history: Contrasting evidence from mitochondrial and nuclear markers of the <i>Odorrana graminea sensu lato</i> (Anura, Ranidae) in China. <i>Molecular Phylogenetics and Evolution</i> , 2020, 144, 106701.	1.2	14
1335	LIMES: a tool for comparing species partition. <i>Bioinformatics</i> , 2020, 36, 2282-2283.	1.8	18
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1337	Dwarf geckos and giant rivers: the role of the São Francisco River in the evolution of <i>Lygodactylus klugei</i> (Squamata: Gekkonidae) in the semi-arid Caatinga of north-eastern Brazil. <i>Biological Journal of the Linnean Society</i> , 2020, 129, 88-98.	0.7	16
1338	Integrative taxonomy uncovers hidden species diversity in the rheophilic genus <i>Potamometra</i> (Hemiptera: Gerridae). <i>Zoologica Scripta</i> , 2020, 49, 174-186.	0.7	26
1339	Invasion of <i>Trichoferus campestris</i> (Coleoptera: Cerambycidae) into the United States characterized by high levels of genetic diversity and recurrent introductions. <i>Biological Invasions</i> , 2020, 22, 1309-1323.	1.2	6
1340	Morphological and molecular analyses of Leafhopper, <i>Amrasca biguttula</i> (Ishida) (Hemiptera: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 187 Asia-Pacific Entomology, 2020, 23, 260-268.	0.4	1
1341	Phylogeny and species delimitation of the genus <i>Longgenacris</i> and <i>Fruhstorferiola viridifemorata</i> species group (Orthoptera: Acrididae: Melanoplinae) based on molecular evidence. <i>PLoS ONE</i> , 2020, 15, e0237882.	1.1	6
1342	Phylogeography and potential glacial refugia of terrestrial gastropod <i>Faustina faustina</i> (RossmÄssler,) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 187 models. <i>Organisms Diversity and Evolution</i> , 2020, 20, 747-762.	0.7	8
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1345	Phylogenomics from transcriptomic "bycatch" clarify the origins and diversity of avian trypanosomes in North America. <i>PLoS ONE</i> , 2020, 15, e0240062.	1.1	8
1346	Evolution and biogeography of the <i>Zanclaea-Scleractinia</i> symbiosis. <i>Coral Reefs</i> , 2022, 41, 779-795.	0.9	18
1347	Molecular taxonomy, phylogeny and biogeography of the <i>Niphargus tatrensis</i> species complex (Amphipoda, Niphargidae) in Austria. <i>Organisms Diversity and Evolution</i> , 2020, 20, 701-722.	0.7	9
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1349	Using multiple lines of evidence to delimit protogynes and deutogynes of four-legged mites: a case study on <i>Epitrimerus sabinae</i> s.l. (Acari : Eriophyidae). <i>Invertebrate Systematics</i> , 2020, , .	0.5	5
1350	Species Diversity in the Braconid Wasp Genus <i>Allorhogas</i> (Doryctinae) Associated With Cynipid Galls on Live Oaks (<i>Quercus</i> : Fagaceae) Using Natural History, Phylogenetics, and Morphology. <i>Insect Systematics and Diversity</i> , 2020, 4, .	0.7	7
1351	Phylogeny-based species delimitation and integrative taxonomic revision of the <i>Hyalinobatrachium fleischmanni</i> species complex, with resurrection of <i>H. viridissimum</i> (Taylor, 1942). <i>Systematics and Biodiversity</i> , 2020, 18, 464-484.	0.5	12
1352	Amphibian taxonomy: early 21st century case studies. <i>Journal of Natural History</i> , 2020, 54, 1-13.	0.2	10
1353	A Species delimitation approach to uncover cryptic species in the South American fire ant decapitating flies (Diptera: Phoridae: Pseudacteon). <i>PLoS ONE</i> , 2020, 15, e0236086.	1.1	8
1354	HaplowebMaker and CoMa: Two web tools to delimit species using haplowebs and conspecificity matrices. <i>Methods in Ecology and Evolution</i> , 2020, 11, 1434-1438.	2.2	32
1355	Molecular markers reveal diversity in composition of <i>Megastigmus</i> (Hymenoptera: Megastigmidae) from eucalypt galls. <i>Ecology and Evolution</i> , 2020, 10, 11565-11578.	0.8	1
1356	DNA Barcoding for the Assessment of the Taxonomy and Conservation Status of the Fish Bycatch of the Northern Brazilian Shrimp Trawl Fishery. <i>Frontiers in Marine Science</i> , 2020, 7, .	1.2	11
1357	Global phylogeography suggests extensive eucosmopolitanism in Mesopelagic Fishes (Maurolicus: Tj ETQq1 1 0.784314 rgBJ_10	1.6	10
1358	New insights into and limitations of the molecular phylogeny in the taxon-rich land snail genus <i>Montenegrina</i> (Mollusca: Gastropoda: Clausiliidae). <i>Journal of Zoological Systematics and Evolutionary Research</i> , 2020, 58, 662-690.	0.6	6
1359	Molecular data suggest multiple origins and diversification times of freshwater gammarids on the Aegean archipelago. <i>Scientific Reports</i> , 2020, 10, 19813.	1.6	10
1360	Multilocus species-delimitation in the <i>Xerotyphlops vermicularis</i> (Reptilia: Typhlopidae) species complex. <i>Molecular Phylogenetics and Evolution</i> , 2020, 152, 106922.	1.2	8
1361	Phylogeography of the "cosmopolitan" orb-weaver <i>Argiope trifasciata</i> (Araneae: Araneidae). <i>Biological Journal of the Linnean Society</i> , 2020, 131, 61-75.	0.7	6

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1363	Hidden diversity in Antarctica: Molecular and morphological evidence of two different species within one of the most conspicuous ascidian species. <i>Ecology and Evolution</i> , 2020, 10, 8127-8143.	0.8	6
1364	Historical biogeography identifies a possible role of Miocene wetlands in the diversification of the Amazonian rocket frogs (Aromobatidae: <i>Allobates</i>). <i>Journal of Biogeography</i> , 2020, 47, 2472-2482.	1.4	31
1365	Molecular phylogeny and trait evolution of Madeiran land snails: radiation of the Geomitriini (Stylommatophora: Helicoidea: Geomitridae). <i>Cladistics</i> , 2020, 36, 594-616.	1.5	6
1366	Biogeography and integrative taxonomy of <i>Epipterygium</i> (Mniaceae, Bryophyta). <i>Taxon</i> , 2020, 69, 1150-1171.	0.4	9
1367	Foliose species of New Zealand red algae: diversity in the genus <i>Tsengia</i> (Tsengiaceae). <i>Tj ETQq1 1 0.784314 rgBT /Overlock 10</i>	0.8	4
1368	Assessing species number and genetic diversity of the Mountainsnails (Oreohelicidae). <i>Conservation Genetics</i> , 2020, 21, 971-985.	0.8	10
1369	Montane regions shape patterns of diversification in small mammals and reptiles from Madagascar's moist evergreen forest. <i>Journal of Biogeography</i> , 2020, 47, 2059-2072.	1.4	10
1370	Species complex diversification by host plant use in an herbivorous insect: The source of Puerto Rican cactus mealybug pest and implications for biological control. <i>Ecology and Evolution</i> , 2020, 10, 10463-10480.	0.8	9
1371	Multilocus and morphological analysis of south-eastern Iberian Wall lizards (Squamata, Podarcis). <i>Zoologica Scripta</i> , 2020, 49, 668-683.	0.7	7
1372	Molecular prospecting for cryptic species of the <i>Hypholoma fasciculare</i> complex: toward the effective and practical delimitation of cryptic macrofungal species. <i>Scientific Reports</i> , 2020, 10, 13224.	1.6	8
1373	DNA Barcoding and geographical scale effect: The problems of undersampling genetic diversity hotspots. <i>Ecology and Evolution</i> , 2020, 10, 10754-10772.	0.8	26
1374	Hidden diversity in <i>Prochilodus nigricans</i> : A new genetic lineage within the Tapaj's River basin. <i>PLoS ONE</i> , 2020, 15, e0237916.	1.1	1
1375	Oviposition and development response of perilla seed bugs (<i>Nysius</i> sp.) (Heteroptera: Lygaeidae) to five crop seeds. <i>Journal of Applied Entomology</i> , 2020, 144, 806-816.	0.8	5
1376	Disparate dispersal limitation in <i>Geomalacoccus</i> slugs unveiled by the shape and slope of the genetic-spatial distance relationship. <i>Ecography</i> , 2020, 43, 1229-1240.	2.1	5
1377	An integrative approach assesses the intraspecific variations of <i>Procamallanus</i> (<i>Spirocamallanus</i>) <i>inopinatus</i> , a common parasite in Neotropical freshwater fishes, and the phylogenetic patterns of Camallanidae. <i>Parasitology</i> , 2020, 147, 1752-1764.	0.7	12
1378	Elucidating species richness in lichen fungi: The genus <i>Stictia</i> (Ascomycota: Peltigeraceae) in Puerto Rico. <i>Taxon</i> , 2020, 69, 851-891.	0.4	11
1379	The unknown diversity of the genus <i>Characidium</i> (Characiformes: Crenuchidae) in the Choc's biogeographic region, Colombian Andes: Two new species supported by morphological and molecular data. <i>Journal of Fish Biology</i> , 2020, 97, 1662-1675.	0.7	6

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1381	Taxonomic revision of the <i>Menticirrhus americanus</i> (Linnaeus, 1758) and <i>M. littoralis</i> (Holbrook, 1847) (Percomorphacea: Sciaenidae) species complexes from the western Atlantic. <i>Zootaxa</i> , 2020, 4822, zootaxa.4822.3.1.	0.2	7
1382	Diversity and host associations of <i>Myrsidea</i> chewing lice (Phthiraptera: Menoponidae) in the tropical rainforest of Malaysian Borneo. <i>International Journal for Parasitology: Parasites and Wildlife</i> , 2020, 13, 231-247.	0.6	3
1383	Phylogenomics supported by geometric morphometrics reveals delimitation of sexual species within the polyploid apomictic <i>Ranunculus auricomus</i> complex (Ranunculaceae). <i>Taxon</i> , 2020, 69, 1191-1220.	0.4	22
1384	Revisiting the past – novel insight into intraspecific molecular diversity and phylogeny of widespread <i>Echinogammarus berilloni</i> . <i>Knowledge and Management of Aquatic Ecosystems</i> , 2020, , 46.	0.5	0
1385	Cryptic Diversity in the Monotypic Neotropical Micromoth Genus <i>Angelabella</i> (Lepidoptera: Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 262 Td	1.0	3
1386	Recent <i>Apareiodon</i> species evolutionary divergence (Characiformes: Parodontidae) evidenced by chromosomal and molecular inference. <i>Zoologischer Anzeiger</i> , 2020, 289, 166-176.	0.4	5
1387	Naming the untouchable – environmental sequences and niche partitioning as taxonomical evidence in fungi. <i>IMA Fungus</i> , 2020, 11, 23.	1.7	15
1388	Morphological and molecular species boundaries in the <i>Hyaella</i> species flock of Lake Titicaca (Crustacea: Amphipoda). <i>Contributions To Zoology</i> , 2020, 89, 353-372.	0.2	10
1389	Lineage diversity, morphological and genetic divergence in <i>Daphnia magna</i> (Crustacea) among Chinese lakes at different altitudes. <i>Contributions To Zoology</i> , 2020, 89, 450-470.	0.2	10
1390	Integrative taxonomic and geographic variation analyses in <i>Cyrtodactylus aequalis</i> (Squamata: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 34 Ecology and Evolution, 2020, 66, 151-179.	0.2	5
1391	New insights into the genetic diversity of the stone crayfish: taxonomic and conservation implications. <i>BMC Evolutionary Biology</i> , 2020, 20, 146.	3.2	25
1392	Phylogenetic relationships and cryptic species in the genus <i>Sthenoteuthis</i> (Cephalopoda: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 262 Td	1.2	8
1393	Species delimitation and phylogenetic relationships in the genus <i>Trypoxylon</i> (Hymenoptera: Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 5 Biodiversity, 2020, 18, 315-327.	0.5	5
1394	Large-scale DNA-based survey of frogs in Amazonia suggests a vast underestimation of species richness and endemism. <i>Journal of Biogeography</i> , 2020, 47, 1781-1791.	1.4	60
1395	Species delimitation through DNA barcoding of freshwater leeches of the <i>Glossiphonia</i> genus (Hirudinea: Glossiphoniidae) from Eastern Siberia, Russia. <i>Journal of Zoological Systematics and Evolutionary Research</i> , 2020, 58, 1437-1446.	0.6	10
1396	Unraveling the species diversity and relationships in the <i>Leptodactylus mystaceus</i> complex (Anura: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5 zootaxa.4779.2.1.	0.2	11
1397	Taxonomic revision of <i>Aegla lata</i> Bond-Buckup & Buckup, 1994 (Decapoda: Anomura: Aegliidae), with the description of a new species of <i>Aegla</i> Leach, 1820 from the Upper Paraná Ecoregion, Brazil. <i>Journal of Crustacean Biology</i> , 2020, 40, 425-434.	0.3	8

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1399	Phylogeny and classification of the leafhopper subfamily Eurymelinae (Hemiptera: Cicadellidae) inferred from molecules and morphology. <i>Systematic Entomology</i> , 2020, 45, 687-702.	1.7	18
1400	Unsupervised biodiversity estimation using proteomic fingerprints from MALDIâ€TOF MS data. <i>Limnology and Oceanography: Methods</i> , 2020, 18, 183-195.	1.0	4
1401	Global radiation in a rare biosphere soil diatom. <i>Nature Communications</i> , 2020, 11, 2382.	5.8	43
1402	Species delimitation analyses of NE Atlantic Chaetozoa (Annelida, Cirratulidae) reveals hidden diversity among a common and abundant marine annelid. <i>Molecular Phylogenetics and Evolution</i> , 2020, 149, 106852.	1.2	14
1403	Cryptic diversity and species boundaries within the <i>Paragalago zanzibaricus</i> species complex. <i>Molecular Phylogenetics and Evolution</i> , 2020, 150, 106887.	1.2	7
1404	Conservative plumage masks extraordinary phylogenetic diversity in the <i>Grallaria rufula</i> (Rufous) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 5	0.7	14
1405	Species diversity revealed in <i>Sigmella</i> Hebard, 1929 (Blattodea, ectobiidae) based on morphology and four molecular species delimitation methods. <i>PLoS ONE</i> , 2020, 15, e0232821.	1.1	5
1406	Unexpected but unsurprising lineage diversity within the most widespread Neotropical crocodylian genus <i>Caiman</i> (Crocodylia, Alligatoridae). <i>Systematics and Biodiversity</i> , 2020, 18, 377-395.	0.5	36
1407	Cryptic species and host specificity in the bryozoan-associated hydrozoan <i>Zanclaea divergens</i> (Hydrozoa, Zanclidae). <i>Molecular Phylogenetics and Evolution</i> , 2020, 151, 106893.	1.2	15
1408	An integrative DNA barcoding framework of ladybird beetles (Coleoptera: Coccinellidae). <i>Scientific Reports</i> , 2020, 10, 10063.	1.6	19
1409	Molecular delimitation and taxonomic revision of the wimple piranha <i>Catoprion</i> (Characiformes: Serrasalminidae) with the description of a new species. <i>Journal of Fish Biology</i> , 2020, 97, 668-685.	0.7	9
1410	Substantial differences in genetic diversity and spatial structuring among (cryptic) amphipod species in a mountainous river basin. <i>Freshwater Biology</i> , 2020, 65, 1641-1656.	1.2	14
1411	Phylogenetic structure of Neotropical annual fish of the genus <i>Cynopoecilus</i> (Cyprinodontiformes) Tj ETQq1 1 0.784314 rgBT /Overlock 0.6 3	0.6	3
1412	Unexpected high abyssal ophiuroid diversity in polymetallic nodule fields of the northeast Pacific Ocean and implications for conservation. <i>Biogeosciences</i> , 2020, 17, 1845-1876.	1.3	35
1413	Coalescence-based species delimitation using genome-wide data reveals hidden diversity in a cosmopolitan group of lichens. <i>Organisms Diversity and Evolution</i> , 2020, 20, 189-218.	0.7	7
1414	The utility of wing morphometrics for assigning type specimens to cryptic bumblebee species. <i>Systematic Entomology</i> , 2020, 45, 849-856.	1.7	5
1415	Global biodiversity of the genus <i>Ommastrephes</i> (Ommastrephidae: Cephalopoda): an allopatric cryptic species complex. <i>Zoological Journal of the Linnean Society</i> , 2020, 190, 460-482.	1.0	24

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1416	Evidence of strong small-scale population structure in the Antarctic freshwater copepod <i>Boeckella poppei</i> in lakes on Signy Island, South Orkney Islands. <i>Limnology and Oceanography</i> , 2020, 65, 2024-2040.	1.6	11
1417	Ant-eating spider maintains specialist diet throughout its ontogeny. <i>Journal of Zoology</i> , 2020, 311, 155-163.	0.8	6
1418	High diversity and pan-oceanic distribution of deep-sea polychaetes: <i>Prionospio</i> and <i>Aurospio</i> (Annelida: Spionidae) in the Atlantic and Pacific Ocean. <i>Organisms Diversity and Evolution</i> , 2020, 20, 171-187.	0.7	16
1419	Description of a new <i>Peronia</i> species (Gastropoda: Eupulmonata: Onchidiidae) from Iran, Persian Gulf. <i>Zootaxa</i> , 2020, 4758, zootaxa.4758.3.5.	0.2	3
1420	COI mtDNA barcoding and morphology for species delimitation in the spider genus <i>Ixchela</i> Huber (Araneae: Pholcidae), with the description of two new species from Mexico. <i>Zootaxa</i> , 2020, 4747, zootaxa.4747.1.2.	0.2	9
1421	Phylogeny and classification of Leptophlebiidae (Ephemeroptera) with an emphasis on Neotropical fauna. <i>Systematic Entomology</i> , 2020, 45, 415-429.	1.7	15
1422	Phylogenetic evidence for an expanded circumscription of <i>Gabura</i> (Arctomiaceae). <i>Lichenologist</i> , 2020, 52, 3-15.	0.5	2
1423	Single-locus <i>scp</i> DNA barcoding and species delimitation of the sandfly subgenus <i>Evandromyia</i> (<i>Aldamyia</i>). <i>Medical and Veterinary Entomology</i> , 2020, 34, 420-431.	0.7	10
1424	In silico and empirical evaluation of twelve metabarcoding primer sets for insectivorous diet analyses. <i>Ecology and Evolution</i> , 2020, 10, 6310-6332.	0.8	28
1425	Miocene Diversification in the Savannahs Precedes Tetraploid Rainforest Radiation in the African Tree Genus <i>Azelia</i> (Detarioideae, Fabaceae). <i>Frontiers in Plant Science</i> , 2020, 11, 798.	1.7	5
1426	Molecular species delimitation in the primitively segmented spider genus <i>Heptathela</i> endemic to Japanese islands. <i>Molecular Phylogenetics and Evolution</i> , 2020, 151, 106900.	1.2	10
1427	Sweet or salty? The origin of freshwater gastrotrichs (Gastrotricha, Chaetonotida) revealed by molecular phylogenetic analysis. <i>Cladistics</i> , 2020, 36, 458-480.	1.5	7
1428	First record of <i>Nassarius fuscus</i> (Hombron & Jacquinet, 1848) from the west coast of India, with the description of its sister species <i>Nassarius arewarensis</i> n. sp. (Buccinoidea: Nassariidae). <i>Journal of Molluscan Studies</i> , 2020, 86, 240-248.	0.4	1
1429	Intragenic evolutionary timing and hidden genetic diversity of <i>Paramecium</i> lineages (Ciliophora: Oligohymenophorea). <i>Systematics and Biodiversity</i> , 2020, 18, 662-674.	0.5	9
1430	Looks can be deceiving: <i>Didemnum pseudovexillum</i> sp. nov. (Asciadiacea) in European harbours. <i>Marine Biodiversity</i> , 2020, 50, .	0.3	10
1431	Make every species count: <i>scp</i> fastachar software for rapid determination of molecular diagnostic characters to describe species. <i>Molecular Ecology Resources</i> , 2020, 20, 1761-1768.	2.2	20
1432	Museomics reveals extensive cryptic diversity of Australian prionine longhorn beetles with implications for their classification and conservation. <i>Systematic Entomology</i> , 2020, 45, 745-770.	1.7	25
1433	A new endangered species of <i>Megaleporinus</i> (Characiformes: Anostomidae) from the Rio de Contas basin, eastern Brazil. <i>Journal of Fish Biology</i> , 2020, 96, 1349-1359.	0.7	7

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1434	Toward an understanding of the systematics and evolution of the genus <i>Acrapex</i> Hampson, 1894 (Lepidoptera: Noctuidae: Apameini: Sesamiina): molecular phylogenetics of the genus and review of the species-rich <i>Acrapex</i> aenigma group. <i>Annales De La Societe Entomologique De France</i> , 2020, 56, 29-91.	0.4	4
1435	The challenge of integrative taxonomy of rare, deep-water gastropods: the genus <i>Exilia</i> (Neogastropoda: Turbinelloidea: Ptychatractidae). <i>Journal of Molluscan Studies</i> , 2020, 86, 120-138.	0.4	0
1436	Disentangling the taxonomy of the subfamily Rasborinae (Cypriniformes, Danionidae) in Sundaland using DNA barcodes. <i>Scientific Reports</i> , 2020, 10, 2818.	1.6	28
1437	Extensive cryptic diversity of giant clams (Cardiidae: Tridacninae) revealed by DNA-sequence-based species delimitation approaches with new data from Hainan Island, South China Sea. <i>Journal of Molluscan Studies</i> , 2020, 86, 56-63.	0.4	9
1438	Multilocus phylogeny and cryptic diversity of white-toothed shrews (Mammalia, Eulipotyphla.) <i>Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 582</i>	3.2	17
1439	Mycoheterotrophic plants living on arbuscular mycorrhizal fungi are generally enriched in ¹³ C, ¹⁵ N and ² H isotopes. <i>Journal of Ecology</i> , 2020, 108, 1250-1261.	1.9	15
1440	A comparative approach for species delimitation based on multiple methods of multi-locus DNA sequence analysis: A case study of the genus <i>Giraffa</i> (Mammalia, Cetartiodactyla). <i>PLoS ONE</i> , 2020, 15, e0217956.	1.1	37
1441	A Plea for Standardized Nuclear Markers in Metazoan DNA Taxonomy. <i>Trends in Ecology and Evolution</i> , 2020, 35, 336-345.	4.2	53
1442	Multiple evolutionary lineages detected in giant reed (<i>Arundo donax</i> L.): Applied and evolutionary perspectives. <i>Annals of Applied Biology</i> , 2020, 176, 285-295.	1.3	3
1443	Integrative analyses on Western Palearctic <i>Lasiommata</i> reveal a mosaic of nascent butterfly species. <i>Journal of Zoological Systematics and Evolutionary Research</i> , 2020, 58, 809-822.	0.6	12
1444	Morpho-molecular delineation of structurally important reef species, the fire corals, <i>Millepora</i> spp., at RÅ©union Island, Southwestern Indian Ocean. <i>Hydrobiologia</i> , 2020, 847, 1237-1255.	1.0	7
1445	Patterns of morphological diversification in giant Berberomeloe blister beetles (Coleoptera:) <i>Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 582</i> <i>Zoological Journal of the Linnean Society</i> , 2020, 189, 1249-1312.	1.0	13
1446	Biodiversity inventory of the grey mullets (Actinopterygii: Mugilidae) of the Indo-Å©Australian Archipelago through the iterative use of DNA-Å©based species delimitation and specimen assignment methods. <i>Evolutionary Applications</i> , 2020, 13, 1451-1467.	1.5	23
1447	The impact of Miocene orogeny for the diversification of Caucasian <i>Epeorus</i> (Caucasiron) mayflies (Ephemeroptera: Heptageniidae). <i>Molecular Phylogenetics and Evolution</i> , 2020, 146, 106735.	1.2	17
1448	Cryptic diversity in ant-Å©mimic <i>Micaria</i> spiders (Araneae, Gnaphosidae) and a tribute to early naturalists. <i>Zoologica Scripta</i> , 2020, 49, 197-209.	0.7	14
1449	Eocene-Å©Oligocene sea-Å©level fall drove amphipod habitat shift from marine to freshwater in the Far East. <i>Zoologica Scripta</i> , 2020, 49, 357-365.	0.7	3
1450	Cryptic diversity and non-adaptive radiation of montane New Guinea skinks (Papuascincus; Scincidae). <i>Molecular Phylogenetics and Evolution</i> , 2020, 146, 106749.	1.2	19
1451	A taxonomic conundrum: Characterizing a cryptic radiation of Asian gracile skinks (Squamata:) <i>Tj ETQq1 1 0.784314 rgBT /Oyerlock 10 Tf 50 582</i>	1.2	10

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1452	DNA barcoding reveals cryptic diversity and peculiar phylogeographic patterns in mojarras (Perciformes: Gerreidae) from the Caribbean and South-western Atlantic. <i>Journal of the Marine Biological Association of the United Kingdom</i> , 2020, 100, 277-283.	0.4	8
1453	A taxonomic revision of <i>Lemna</i> sect. <i>Uninerves</i> (Lemnaceae). <i>Taxon</i> , 2020, 69, 56-66.	0.4	46
1454	Ultra-Barcoding Discovers a Cryptic Species in <i>Paris yunnanensis</i> (Melanthiaceae), a Medicinally Important Plant. <i>Frontiers in Plant Science</i> , 2020, 11, 411.	1.7	32
1455	Multilocus phylogeny of Paratelmatobiinae (Anura: Leptodactylidae) reveals strong spatial structure and previously unknown diversity in the Atlantic Forest hotspot. <i>Molecular Phylogenetics and Evolution</i> , 2020, 148, 106819.	1.2	22
1456	Biotope and biocenosis of cryoconite hole ecosystems on Ecology Glacier in the maritime Antarctic. <i>Science of the Total Environment</i> , 2020, 724, 138112.	3.9	22
1457	Phylogeography and genetic diversity of the copepod family Cyclopidae (Crustacea: Cyclopoida) from freshwater ecosystems of Southeast Nigeria. <i>BMC Evolutionary Biology</i> , 2020, 20, 45.	3.2	2
1458	A comprehensive molecular phylogeny of Afrotropical white-eyes (Aves: Zosteropidae) highlights prior underestimation of mainland diversity and complex colonisation history. <i>Molecular Phylogenetics and Evolution</i> , 2020, 149, 106843.	1.2	13
1459	Genome-informed integrative taxonomic description of three cryptic species in the earthworm genus <i>Carpetania</i> (Oligochaeta, Hormogastridae). <i>Systematics and Biodiversity</i> , 2020, 18, 203-215.	0.5	16
1460	Repositories for Taxonomic Data: Where We Are and What is Missing. <i>Systematic Biology</i> , 2020, 69, 1231-1253.	2.7	38
1461	Intraspecific variation in sensitivity to food availability and temperature-induced phenotypic plasticity in the rotifer <i>Keratella cochlearis</i> . <i>Journal of Experimental Biology</i> , 2020, 223, .	0.8	7
1462	Hidden diversity within the <i>Diopatra cuprea</i> complex (Annelida: Onuphidae): morphological and genetics analyses reveal four new species in the south-west Atlantic. <i>Zoological Journal of the Linnean Society</i> , 2021, 191, 637-671.	1.0	13
1463	Molecular diversity and species delimitation in the family Gasteruptionidae (Hymenoptera: Evanoidea). <i>Genome</i> , 2021, 64, 253-264.	0.9	8
1464	A new species group from the <i>Daphnia curvirostris</i> species complex (Cladocera: Anomopoda) from the eastern Palaearctic: taxonomy, phylogeny and phylogeography. <i>Zoological Journal of the Linnean Society</i> , 2021, 191, 772-822.	1.0	25
1465	Regional-scale aquifer hydrogeology as a driver of phylogeographic structure in the Neotropical catfish <i>Rhamdia guatemalensis</i> (Siluriformes: Heptapteridae) from cenotes of the Yucatán Peninsula, Mexico. <i>Freshwater Biology</i> , 2021, 66, 332-348.	1.2	5
1466	Lineage diversity and gene introgression in freshwater cladoceran crustaceans of the <i>Chydorus sphaericus</i> species complex. <i>Limnology and Oceanography</i> , 2021, 66, 95-107.	1.6	12
1467	Phylogeny of Lithobiidae Newport, 1844, with emphasis on the megadiverse genus <i>Lithobius</i> Leach, 1814 (Myriapoda, Chilopoda). <i>Cladistics</i> , 2021, 37, 162-184.	1.5	5
1468	Unexpected cryptic species diversity of parasites of the family Xenidae (Strepsiptera) with a constant diversification rate over time. <i>Systematic Entomology</i> , 2021, 46, 252-265.	1.7	7
1469	Phylogeographical history of the Olive Woodpecker <i>Dendropicos griseocephalus</i> , a species widely distributed across Africa. <i>Ibis</i> , 2021, 163, 417-428.	1.0	2

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1470	The omission of critical data in the pursuit of "revolutionary"™ methods to accelerate the description of species. <i>Systematic Entomology</i> , 2021, 46, 1-4.	1.7	28
1471	Italian odonates in the Pandora's box: A comprehensive DNA barcoding inventory shows taxonomic warnings at the Holarctic scale. <i>Molecular Ecology Resources</i> , 2021, 21, 183-200.	2.2	37
1472	Genetic diversity and morphological stasis in the Ceylon Snakehead, <i>Channa orientalis</i> (Teleostei). <i>Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50</i>	0.5	8
1473	Revision shock in Pacific oysters taxonomy: the genus <i>Magallana</i> (formerly <i>Crassostrea</i> in) <i>Tj ETQq1 1 0,784314 rgBT /Overlock 18</i>	1.0	18
1474	Foliose <i>Ulva</i> Species Show Considerable Inter-specific Genetic Diversity, Low Intra-specific Genetic Variation, and the Rare Occurrence of Inter-specific Hybrids in the Wild. <i>Journal of Phycology</i> , 2021, 57, 219-233.	1.0	24
1475	Extended phenotypes on coral reefs: cryptic phenotypes modulate coral-vermetid interactions. <i>Ecology</i> , 2021, 102, e03215.	1.5	1
1476	Multiple species delimitation approaches applied to the avian lark genus <i>Alaudala</i> . <i>Molecular Phylogenetics and Evolution</i> , 2021, 154, 106994.	1.2	14
1477	Assessment of fish diversity in the South China Sea using DNA taxonomy. <i>Fisheries Research</i> , 2021, 233, 105771.	0.9	16
1478	ASAP: assemble species by automatic partitioning. <i>Molecular Ecology Resources</i> , 2021, 21, 609-620.	2.2	575
1479	Another stripe on the tiger makes no difference? Unexpected diversity in the widespread tiger tarantula <i>Davus pentaloris</i> (Araneae: Theraphosidae: Theraphosinae). <i>Zoological Journal of the Linnean Society</i> , 2021, 192, 75-104.	1.0	8
1480	Integrative taxonomy reveals hidden cestode diversity in <i>Pimelodus</i> catfishes in the Neotropics. <i>Zoologica Scripta</i> , 2021, 50, 210-224.	0.7	10
1481	High diversity and strong variation in host specificity of seed parasitic acorn weevils. <i>Insect Conservation and Diversity</i> , 2021, 14, 367-376.	1.4	5
1482	Phylogeography of a widespread Palearctic forest bird species: The White-backed Woodpecker (Aves.) <i>Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50</i>	0.7	6
1483	Phylogenetics, historical biogeography and molecular species delimitation of <i>Gnaptorina</i> Reitter (Coleoptera: Tenebrionidae: Blaptini). <i>Systematic Entomology</i> , 2021, 46, 239-251.	1.7	8
1484	Contrasting biogeographical patterns in <i>Margarella</i> (Gastropoda: Calliostomatidae: Margarellinae) across the Antarctic Polar Front. <i>Molecular Phylogenetics and Evolution</i> , 2021, 156, 107039.	1.2	14
1485	P2C2M.GMYC: An R package for assessing the utility of the Generalized Mixed Yule Coalescent model. <i>Methods in Ecology and Evolution</i> , 2021, 12, 487-493.	2.2	10
1486	DNA-based and taxonomic identification of forensically important Sarcophagidae (Diptera) in southeastern Spain. <i>Science and Justice - Journal of the Forensic Science Society</i> , 2021, 61, 150-159.	1.3	6
1487	New records of <i>Chondracanthus saundersii</i> and <i>Schottera koreana</i> (Gigartinales, Rhodophyta) from Japan based on molecular and morphological analyses. <i>Phycological Research</i> , 2021, 69, 81-87.	0.8	2

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1488	Overlooked biodiversity of mitochondrial lineages in Hemiodus (Ostariophysi, Characiformes). <i>Zoologica Scripta</i> , 2021, 50, 337-351.	0.7	5
1489	Hidden introductions of freshwater red algae via the aquarium trade exposed by DNA barcodes. <i>Environmental DNA</i> , 2021, 3, 481-491.	3.1	4
1490	Revisiting the phylogeny of the genus <i>Lolliguncula</i> Steenstrup 1881 improves understanding of their biogeography and proves the validity of <i>Lolliguncula argus</i> Brakoniecki & Roper, 1985. <i>Molecular Phylogenetics and Evolution</i> , 2021, 154, 106968.	1.2	13
1491	Molecular Characterization of Mosquito Diversity in the Balearic Islands. <i>Journal of Medical Entomology</i> , 2021, 58, 608-615.	0.9	9
1492	New Distribution and Molecular Diversity of the Reniform Nematode <i>Rotylenchulus macrosoma</i> (Nematoda: Rotylenchulinae) in Europe. <i>Phytopathology</i> , 2021, 111, 720-730.	1.1	4
1493	Leeches as the intermediate host for strigeid trematodes: genetic diversity and taxonomy of the genera <i>Australapatemon</i> Sudarikov, 1959 and <i>Cotylurus</i> Szidat, 1928. <i>Parasites and Vectors</i> , 2021, 14, 44.	1.0	7
1494	A novel dataset to identify the endemic herpetofauna of the New Caledonia biodiversity hotspot with DNA barcodes. <i>Pacific Conservation Biology</i> , 2022, 28, 36-47.	0.5	6
1495	Morphological and Molecular Assessments of Bobtail Squids (Cephalopoda: Sepiolidae) Reveal a Hidden History of Biodiversity. <i>Frontiers in Marine Science</i> , 2021, 7, .	1.2	16
1496	Molecular Data Reveal the Presence of Three <i>Plocamium</i> Lamouroux Species with Complex Patterns of Distribution in Southern Chile. <i>Cryptogamie, Algologie</i> , 2021, 42, .	0.3	4
1497	Genetic analyses reveal cryptic diversity in the widely distributed. <i>Invertebrate Systematics</i> , 2021, 35, 298-311.	0.5	6
1498	Microbiota associated with <i>Mollitrichosiphum</i> aphids (Hemiptera: Aphididae: Greenideinae): diversity, host species specificity and phyllosymbiosis. <i>Environmental Microbiology</i> , 2021, 23, 2184-2198.	1.8	14
1499	Cryptic Clitellata: Molecular Species Delimitation of Clitellate Worms (Annelida): An Overview. <i>Diversity</i> , 2021, 13, 36.	0.7	11
1500	Molecular identification of <i>Brachygenys</i> and <i>Haemulon</i> species (Perciformes: Haemulidae) from the Brazilian coast. <i>Neotropical Ichthyology</i> , 2021, 19, .	0.5	2
1501	A new <i>Hyphessobrycon</i> (Characiformes: Characidae) of the <i>Hyphessobrycon heterorhabdus</i> species-group from the lower Amazon basin, Brazil. <i>Neotropical Ichthyology</i> , 2021, 19, .	0.5	6
1502	Beyond the comfort zone: amphibian diversity and distribution in the West Sahara-Sahel using mtDNA and nuDNA barcoding and spatial modelling. <i>Conservation Genetics</i> , 2021, 22, 233-248.	0.8	2
1503	Integrative taxonomy reveals disjunct distribution and first record of <i>Hoplias misionera</i> (Characiformes: Erythrinidae) in the Amazon River basin: morphological, DNA barcoding and cytogenetic considerations. <i>Neotropical Ichthyology</i> , 2021, 19, .	0.5	5
1504	Species boundaries and phylogeographic patterns in new species of <i>Nannoniscus</i> (Janiroidea): Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5 <i>Zoological Journal of the Linnean Society</i> , 2021, 193, 1020-1071.	1.0	9
1505	Impact of Pleistocene Eustatic Fluctuations on Evolutionary Dynamics in Southeast Asian Biodiversity Hotspots. <i>Systematic Biology</i> , 2021, 70, 940-960.	2.7	25

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1506	Late Pliocene population divergence and persistence despite Pleistocene climatic fluctuations in the Rio Doce snouted Treefrog (<i>Oloolygon carnevallii</i>). <i>Journal of Zoological Systematics and Evolutionary Research</i> , 2021, 59, 680-690.	0.6	1
1507	Use of species delimitation approaches to tackle the cryptic diversity of an assemblage of high Andean butterflies (Lepidoptera: Papilionoidea). <i>Genome</i> , 2021, 64, 937-949.	0.9	3
1508	Genetic structure and population demographics of <i>Hypnoidus bicolor</i> (Coleoptera: Tj ETQq0 0 0 rgBT /Overlock 10 2021, 77, 2282-2291.	1.7	6
1509	<sc>DNA</sc> barcoding: A different perspective to introducing undergraduate students to <sc>DNA</sc> sequence analysis. <i>Biochemistry and Molecular Biology Education</i> , 2021, 49, 416-421.	0.5	3
1510	Island colonization by a rheophilic fish: the phylogeography of <i>Garra ceylonensis</i> (Teleostei: Tj ETQq0 0 0 rgBT /Overlock 10 2021, 77, 2282-2291.	0.7	11
1511	Molecular systematics and biogeography of an Australian soil-burrowing cockroach with polymorphic males, <i>Geoscapheus dilatatus</i> (Blattodea: Blaberidae). <i>Austral Entomology</i> , 2021, 60, 317-329.	0.8	0
1512	Ecological and spatial patterns associated with diversification of South American Physaria (Brassicaceae) through the general concept of species. <i>Organisms Diversity and Evolution</i> , 2021, 21, 161-188.	0.7	3
1513	Molecular species delimitation reveals hidden specific diversity within a freshwater burrowing crayfish (Decapoda: Parastacidae) from southern Chile. <i>Systematics and Biodiversity</i> , 2021, 19, 237-251.	0.5	6
1514	Unexpected mitochondrial lineage diversity within the genus <i>Alonella</i> Sars, 1862 (Crustacea: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 2021, 77, 2282-2291.	0.9	14
1515	Integrative Taxonomy and Molecular Phylogeny of the Plant-Parasitic Nematode Genus <i>Paratylenchus</i> (Nematoda: Paratylenchinae): Linking Species with Molecular Barcodes. <i>Plants</i> , 2021, 10, 408.	1.6	22
1516	Cytochrome oxidase subunit I barcode species delineation methods imply critically underestimated diversity in common <i>Hermeuptychia</i> butterflies (Lepidoptera: Nymphalidae: Tj ETQq0 0 0 rgBT /Overlock 10 2021, 77, 2282-2291.	1.0	5
1517	Copepods and ostracods associated with bromeliads in the Yucatán Peninsula, Mexico. <i>PLoS ONE</i> , 2021, 16, e0248863.	1.1	3
1519	Systematics and biogeography of the <i>Boana albopunctata</i> species group (Anura, Hylidae), with the description of two new species from Amazonia. <i>Systematics and Biodiversity</i> , 2021, 19, 375-399.	0.5	20
1520	Molecular biogeography of the Mediterranean <i>Buthus</i> species complex (Scorpiones: Buthidae) at its southern Palaearctic margin. <i>Biological Journal of the Linnean Society</i> , 2021, 133, 166-178.	0.7	9
1521	Phylogeography and population genetic structure of red muntjacs: evidence of enigmatic Himalayan red muntjac from India. <i>Bmc Ecology and Evolution</i> , 2021, 21, 49.	0.7	3
1522	Morphological and phylogenetic analyses of <i>Toniniopsis subincompta</i> s. lat. (<i>Ramalinaceae</i> , <i>Lecanorales</i>) in Eurasia. <i>Lichenologist</i> , 2021, 53, 171-183.	0.5	4
1523	Comparative phylogeography of bromeliad species: effects of historical processes and mating system on genetic diversity and structure. <i>Botanical Journal of the Linnean Society</i> , 2021, 197, 263-276.	0.8	3
1525	Species delimitation with distinct methods based on molecular data to elucidate species boundaries in the <i>Cycas taiwaniana</i> complex (Cycadaceae). <i>Taxon</i> , 2021, 70, 477-491.	0.4	8

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1526	Accounting for species interactions is necessary for predicting how arctic arthropod communities respond to climate change. <i>Ecography</i> , 2021, 44, 885-896.	2.1	24
1527	Neotype designation and re-description of Forsskål's reticulate whipray <i>Himantura uarnak</i> . <i>Marine Biodiversity</i> , 2021, 51, 1.	0.3	2
1529	New <i>Cladosporium</i> Species from Normal and Galled Flowers of Lamiaceae. <i>Pathogens</i> , 2021, 10, 369.	1.2	11
1530	A multilocus assessment reveals two new synonymies for East Asian <i>Cyclommatus</i> stag beetles (Coleoptera, Lucanidae). <i>ZooKeys</i> , 2021, 1021, 65-79.	0.5	2
1532	Factors affecting the efficiency of molecular species delimitation in a species-rich insect family. <i>Molecular Ecology Resources</i> , 2021, 21, 1475-1489.	2.2	28
1533	Morphological and molecular evidence support the taxonomic separation of the medically important Neotropical spiders <i>Phoneutria depilata</i> (Strand, 1909) and <i>P. boliviensis</i> (F.O. Pickard-Cambridge, 1897) (Araneae, Ctenidae). <i>ZooKeys</i> , 2021, 1022, 13-50.	0.5	11
1534	Pond chironomid communities revealed by molecular species delimitation reflect eutrophication. <i>Ecology and Evolution</i> , 2021, 11, 4193-4204.	0.8	6
1535	Sareomycetes: more diverse than meets the eye. <i>IMA Fungus</i> , 2021, 12, 6.	1.7	8
1536	Entangled Aeglidæ (Decapoda, Anomura): Additional evidence for cryptic species. <i>Zoologica Scripta</i> , 2021, 50, 473-484.	0.7	4
1537	Molecular Signatures of Reticulate Evolution within the Complex of European Pine Taxa. <i>Forests</i> , 2021, 12, 489.	0.9	1
1538	An integrative approach to address species limits in the southernmost members of the <i>Liolaemus kingii</i> group (Squamata: Liolaemini). <i>Molecular Phylogenetics and Evolution</i> , 2021, 157, 107046.	1.2	5
1539	<i>Gymnotus paraguensis</i> , a Good Example of Phenotypic Plasticity in the Pantanal Biome, Brazil. <i>Zebrafish</i> , 2021, 18, 162-173.	0.5	1
1540	<i>Bosminopsis deitersi</i> (Crustacea: Cladocera) as an ancient species group: a revision. <i>PeerJ</i> , 2021, 9, e11310.	0.9	9
1541	An integrative taxonomic review of the <i>Natalanatalis</i> mountain catfish, <i>Amphilius natalensis</i> Boulenger 1917 (<i>Siluriformes</i> , <i>Amphiliidae</i>), with description of four new species. <i>Journal of Fish Biology</i> , 2021, 99, 219-239.	0.7	8
1542	A double-edged sword: Unrecognized cryptic diversity and taxonomic impediment in <i>Eois</i> (Lepidoptera, Geometridae). <i>Zoologica Scripta</i> , 2021, 50, 633-646.	0.7	3
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1544	Genetic diversity, population structure and rapid early detection of the parasitoid <i>Anastatus orientalis</i> (Hymenoptera: Eupelmidae) inside eggs of spotted lanternfly (Hemiptera: Fulgoridae). <i>Annals of Applied Biology</i> , 2021, 179, 12-20.	1.3	2
1545	A review of the taxonomy of spiny-backed orb-weaving spiders of the subfamily <i>Gasteracanthinae</i> (Araneae, Araneidae) in Thailand. <i>ZooKeys</i> , 2021, 1032, 17-62.	0.5	2

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1549	Molecular phylogeny and species delimitation of the genus <i>Tonkinacris</i> (Orthoptera, Acrididae). <i>Tj ETQq1 1 0.784314 rgBT /Overlock 10</i>	1.1	4
1550	The elephant in the room: first record of invasive gregarious species of serpulids (calcareous tube) <i>Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50</i>	0.3	10
1551	Disentangling the taxonomy of <i>Belostoma Latreille</i> (Hemiptera: Belostomatidae) with the aid of DNA barcoding approaches. <i>Austral Entomology</i> , 2021, 60, 514-524.	0.8	1
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1555	Evaluating species origins within tropical sky-islands arthropod communities. <i>Journal of Biogeography</i> , 2021, 48, 2199-2210.	1.4	7
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1557	Assessing the suitability of mitochondrial and nuclear DNA genetic markers for molecular systematics and species identification of helminths. <i>Parasites and Vectors</i> , 2021, 14, 233.	1.0	33
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1559	Species delimitation of <i>Margattea</i> cockroaches from China, with seven new species (Blattodea). <i>Tj ETQq1 1 0.784314 rgBT /Overlock 10</i>	0.5	3
1560	Integrative species delimitation based on COI, ITS, and morphological evidence illustrates a unique evolutionary history of the genus <i>Paracercion</i> (Odonata: Coenagrionidae). <i>PeerJ</i> , 2021, 9, e11459.	0.9	10
1561	Species Delimitation of Scavenger Flies in the Valley of Mexico. <i>Journal of Medical Entomology</i> , 2021, 58, 2206-2215.	0.9	2
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1563	Using Genealogical Concordance and Coalescent-Based Species Delimitation to Assess Species Boundaries in the <i>Diaporthe eres</i> Complex. <i>Journal of Fungi</i> (Basel, Switzerland), 2021, 7, 507.	1.5	19

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1565	Effectiveness of DNA Barcoding in discriminating <i>Daniellia ogea</i> (Harms) Rolfe ex Holland and <i>Daniellia oliveri</i> (Rolfe) Hutch. & Dalziel. <i>Trees, Forests and People</i> , 2021, 4, 100067.	0.8	1
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1574	Morphological and molecular characterisation of <i>Trichodorus hellalae</i> n. sp. (Nematoda: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 342 Td (T	0.8	2
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1576	The hidden diversity of the genus <i>Lyurella</i> Derzhavin, 1939 (Crustacea: Amphipoda: Crangonyctidae): four new species from the subterranean habitats of the northwestern Caucasus, Russia. <i>Zootaxa</i> , 2021, 5006, 127-168.	0.2	7
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1578	Species-specific <i>COI</i> primers for rapid identification of <i>Bemisia tabaci</i> Mediterranean (MED) species. <i>Journal of Applied Entomology</i> , 2021, 145, 1029-1038.	0.8	4
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1598	Assessment of putative species diversity of insect larvae occurring in bamboo galls by a DNA barcoding analysis. <i>Entomological Science</i> , 2021, 24, 338-344.	0.3	0
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1688	Subtle Ecological Gradient in the Tropics Triggers High Species-Turnover in a Local Geographical Scale. <i>PLoS ONE</i> , 2016, 11, e0156840.	1.1	7
1689	Hidden Genetic Diversity in an Asexually Reproducing Lichen Forming Fungal Group. <i>PLoS ONE</i> , 2016, 11, e0161031.	1.1	23
1690	Complex within a Complex: Integrative Taxonomy Reveals Hidden Diversity in <i>Cicadetta brevipennis</i> (Hemiptera: Cicadidae) and Unexpected Relationships with a Song Divergent Relative. <i>PLoS ONE</i> , 2016, 11, e0165562.	1.1	11
1691	High Species Richness of <i>Scinax</i> Treefrogs (Hylidae) in a Threatened Amazonian Landscape Revealed by an Integrative Approach. <i>PLoS ONE</i> , 2016, 11, e0165679.	1.1	39
1692	Species Delimitation and Phylogenetic Relationships in Ectobiid Cockroaches (Dictyoptera, Blattodea) from China. <i>PLoS ONE</i> , 2017, 12, e0169006.	1.1	25
1693	Genetic structure and distribution of <i>Parisotoma notabilis</i> (Collembola) in Europe: Cryptic diversity, split of lineages and colonization patterns. <i>PLoS ONE</i> , 2017, 12, e0170909.	1.1	16
1694	Species delimitation in asexual insects of economic importance: The case of black scale (<i>Parasaissetia</i>) Tj ETQq1 1 0.784314 ggBT /Overl	1.1	17
1695	Evolutionary history and phylogeographic relationships of shrews from <i>Sorex araneus</i> group. <i>PLoS ONE</i> , 2017, 12, e0179760.	1.1	24
1696	Genetic diversity and stock identification of small abalone (<i>Haliotis diversicolor</i>) in Taiwan and Japan. <i>PLoS ONE</i> , 2017, 12, e0179818.	1.1	12
1697	Species delimitation in frogs from South American temperate forests: The case of <i>Eupsophus</i> , a taxonomically complex genus with high phenotypic variation. <i>PLoS ONE</i> , 2017, 12, e0181026.	1.1	26
1698	Barcoding snakeheads (Teleostei, Channidae) revisited: Discovering greater species diversity and resolving perpetuated taxonomic confusions. <i>PLoS ONE</i> , 2017, 12, e0184017.	1.1	44
1699	Phylogeography and DNA-based species delimitation provide insight into the taxonomy of the polymorphic rose chafer <i>Protaetia</i> (<i>Potosia</i>) <i>cuprea</i> species complex (Coleoptera: Scarabaeidae): Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 2	1.1	17
1700	An integrative approach reveals five new species of highland papayas (Caricaceae, <i>Vasconcellea</i>) from northern Peru. <i>PLoS ONE</i> , 2020, 15, e0242469.	1.1	10
1701	How many species are there in the subgenus <i>Bursaphis</i> (Hemiptera: Sternorrhyncha: Aphididae)? CO-I evidence. <i>European Journal of Entomology</i> , 2011, 108, 469-479.	1.2	11
1702	A review of taxonomy and flower-breeding ecology of the <i>Colocasiomyia toshiokai</i> species group (Diptera: Drosophilidae), with description of a new species from Indonesia. <i>European Journal of Entomology</i> , 0, 116, 341-361.	1.2	10
1703	Ecological and molecular diversity of <i>Eulachnini</i> aphids (Hemiptera: Aphididae: Lachninae) on coniferous plants in Lithuania. <i>European Journal of Entomology</i> , 0, 117, 199-209.	1.2	2
1704	Three new species of the <i>Eigenmannia trilineata</i> species group (Gymnotiformes: Sternopygidae) from northwestern South America. <i>Neotropical Ichthyology</i> , 2020, 18, .	0.5	4

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1706	Phylogeny and Biogeography of the Amazonian <i>Pachyptera</i> (Bignoniaceae). <i>Systematic Botany</i> , 2020, 45, 361-374.	0.2	6
1707	Taxonomic Review of South American Butter Frogs: Phylogeny, Geographic Patterns, and Species Delimitation in the <i>Leptodactylus latrans</i> Species Group (Anura: Leptodactylidae). <i>Herpetological Monographs</i> , 2020, 34, .	1.1	15
1708	Morphological and DNA barcode species identifications of leafhoppers, planthoppers and treehoppers (Hemiptera: Auchenorrhyncha) at Barrow Island. <i>Records of the Western Australian Museum, Supplement</i> , 2013, 83, 253.	0.5	17
1709	Species Delimitation and Description of <i>Mesocriconema nebraskense</i> n. sp. (Nematoda: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 587 Td (C Grasslands. <i>Journal of Nematology</i> , 2017, 49, 42-66.	0.4	39
1710	A COI DNA barcoding survey of <i>Pratylenchus</i> species in the Great Plains Region of North America. <i>Journal of Nematology</i> , 2019, 51, 1-21.	0.4	11
1711	Cryptic species within the rotifer <i>Lecane bulla</i> (Rotifera: Monogononta: Lecanidae) from North America based on molecular species delimitation. <i>Revista Mexicana De Biodiversidad</i> , 2020, 91, 913116.	0.4	5
1712	Sequence-based species delimitation in the Balkan <i>Bythinella</i> (Gastropoda: Risssooidea) with general mixed yule coalescent model. <i>Folia Malacologica</i> , 2012, 20, 111-120.	0.1	9
1713	Spirorchiidiasis in marine turtles: the current state of knowledge. <i>Diseases of Aquatic Organisms</i> , 2019, 133, 217-245.	0.5	19
1714	Close congruence between Barcode Index Numbers (bins) and species boundaries in the Erebidae (Lepidoptera: Noctuoidea) of the Iberian Peninsula. <i>Biodiversity Data Journal</i> , 2017, 5, e19840.	0.4	21
1715	Phylogeny and species delimitations in the entomopathogenic genus <i>Beauveria</i> (Hypocreales, Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 342	0.8	37
1716	Cryptic diversity and mtDNA phylogeography of the invasive demon shrimp, <i>Dikerogammarus haemobaphes</i> (Eichwald, 1841), in Europe. <i>NeoBiota</i> , 0, 57, 53-86.	1.0	26
1717	Morphometrics and phylogeography of the cave-obligate land snail <i>Helicodiscus barri</i> (Gastropoda, Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50	5.0	10
1718	Species delimitation in northern European water scavenger beetles of the genus <i>Hydrobius</i> (Coleoptera, Hydrophilidae). <i>ZooKeys</i> , 2016, 564, 71-120.	0.5	24
1719	Phylogeography of a good Caribbean disperser: <i>Argiope argentata</i> (Araneae, Araneidae) and a new "cryptic" species from Cuba. <i>ZooKeys</i> , 2016, 625, 25-44.	0.5	37
1720	Species delimitation of the <i>Hyphydrus ovatus</i> complex in western Palaearctic with an update of species distributions (Coleoptera, Dytiscidae). <i>ZooKeys</i> , 2017, 678, 73-96.	0.5	3
1721	A revision of the new genus <i>Amiga</i> Nakahara, Willmott & Espeland, gen. n., described for <i>Papilio arnaca</i> Fabricius, 1776 (Lepidoptera, Nymphalidae, Satyrinae). <i>ZooKeys</i> , 2019, 821, 85-152.	0.5	11
1722	DNA barcoding of aphid-associated ants (Hymenoptera, Formicidae) in a subtropical area of southern China. <i>ZooKeys</i> , 2019, 879, 117-136.	0.5	9

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1723	Under an integrative taxonomic approach: the description of a new species of the genus <i>Loxosceles</i> (Araneae, Sicariidae) from Mexico City. <i>ZooKeys</i> , 2019, 892, 93-133.	0.5	12
1724	Twenty-six new species of <i>Hoploscopa</i> (Lepidoptera, Crambidae) from South-East Asia revealed by morphology and DNA barcoding. <i>ZooKeys</i> , 2020, 907, 1-99.	0.5	4
1725	Diversity and distribution of <i>Epeorus</i> (Caucasiron) (Ephemeroptera, Heptageniidae) in Iran, with descriptions of three new species. <i>ZooKeys</i> , 2020, 947, 71-102.	0.5	8
1726	Uncovering the shell game with barcodes: diversity of meiofaunal Caecidae snails (Truncatelloidea, Tj ETQq1 1 0.784314 rgBT /Overlo	0.5	2
1727	Species delimitation methods put into taxonomic practice: two new <i>Madascincus</i> species formerly allocated to historical species names (Squamata, Scincidae). <i>Zoosystematics and Evolution</i> , 2016, 92, 257-275.	0.4	5
1728	Molecular phylogeography and reproductive biology of the freshwater snail <i>Tarebia granifera</i> in Thailand and Timor (Cerithioidea, Thiaridae): morphological disparity versus genetic diversity. <i>Zoosystematics and Evolution</i> , 2018, 94, 461-493.	0.4	10
1729	Identification of <i>Sirex noctilio</i> and Native North American Woodwasp Larvae using DNA Barcode. <i>Journal of Entomology</i> , 2010, 7, 60-79.	0.2	13
1730	A pathogen of New Zealand <i>Pyropia plicata</i> (Bangiales, Rhodophyta), <i>Pythium porphyrae</i> (Oomycota). <i>Algae</i> , 2017, 32, 29-39.	0.9	18
1731	Molecular species discovery in the diatom <i>Sellaphora</i> and its congruence with mating trials.. <i>Fottea</i> , 2013, 13, 133-148.	0.4	33
1732	Difference without distinction? Gaps in cyanobacterial systematics; when more is just too much. <i>Fottea</i> , 2018, 18, 130-136.	0.4	13
1733	The Input of DNA Sequences to Animal Systematics: Rodents as Study Cases. , 0, , .		5
1734	HACSim: an R package to estimate intraspecific sample sizes for genetic diversity assessment using haplotype accumulation curves. <i>PeerJ Computer Science</i> , 2020, 6, e243.	2.7	18
1735	Molecular phylogeny and species delimitation of the freshwater prawn <i>Macrobrachium pilimanus</i> species group, with descriptions of three new species from Thailand. <i>PeerJ</i> , 2020, 8, e10137.	0.9	7
1736	Phylogenetic species delimitation for crayfishes of the genus <i>Pacifastacus</i> . <i>PeerJ</i> , 2016, 4, e1915.	0.9	29
1737	Bird migratory flyways influence the phylogeography of the invasive brine shrimp <i>Artemia franciscana</i> in its native American range. <i>PeerJ</i> , 2013, 1, e200.	0.9	44
1738	Mitochondrial DNA hyperdiversity and its potential causes in the marine periwinkle <i>Melarhaphe neritoides</i> (Mollusca: Gastropoda). <i>PeerJ</i> , 2016, 4, e2549.	0.9	15
1739	Genetic and morphological analyses indicate that the Australian endemic scorpion <i>Urodacus yaschenkoi</i> (Scorpiones: Urodacidae) is a species complex. <i>PeerJ</i> , 2017, 5, e2759.	0.9	6
1740	Neogene paleogeography provides context for understanding the origin and spatial distribution of cryptic diversity in a widespread Balkan freshwater amphipod. <i>PeerJ</i> , 2017, 5, e3016.	0.9	65

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1741	Phylogenetic surveys on the newt genus <i>Tylototriton sensu lato</i> (Salamandridae, Caudata) reveal cryptic diversity and novel diversification promoted by historical climatic shifts. PeerJ, 2018, 6, e4384.	0.9	17
1742	First endemic freshwater <i>Gammarus</i> from Crete and its evolutionary history—an integrative taxonomy approach. PeerJ, 2018, 6, e4457.	0.9	14
1743	A phylogenetic taxonomy of the <i>Cyrtodactylus peguensis</i> group (Reptilia: Squamata: Gekkonidae) with descriptions of two new species from Myanmar. PeerJ, 2018, 6, e5575.	0.9	16
1744	Uncovering hidden specific diversity of Andean glassfrogs of the <i>Centrolene buckleyi</i> species complex (Anura: Centrolenidae). PeerJ, 2018, 6, e5856.	0.9	8
1745	Multiple global radiations in tadpole shrimps challenge the concept of “living fossils”. PeerJ, 2013, 1, e62.	0.9	44
1746	The biogeography of <i>Elaphe sauromates</i> (Pallas, 1814), with a description of a new rat snake species. PeerJ, 2019, 7, e6944.	0.9	19
1747	Molecular phylogeny and intraspecific differentiation of the <i>Trapelus agilis</i> species complex in Iran (Squamata: Agamidae) inferred from mitochondrial DNA sequences. PeerJ, 2020, 8, e8295.	0.9	4
1748	Ancient genetic divergence in bumblebee catfish of the genus <i>Pseudopimelodus</i> (Pseudopimelodidae: Siluriformes) from northwestern South America. PeerJ, 2020, 8, e9028.	0.9	12
1749	A little frog leaps a long way: compounded colonizations of the Indian Subcontinent discovered in the tiny Oriental frog genus <i>Microhyla</i> (Amphibia: Microhylidae). PeerJ, 2020, 8, e9411.	0.9	29
1750	Resolving the <i>Tetrastigma loheri</i> s. l. Species Complex (Vitaceae) in the Philippines: No Evidence for Recognizing More than One Species. Systematic Botany, 2021, 46, 750-763.	0.2	3
1751	Marine Biodiscovery in a Changing World. Progress in the Chemistry of Organic Natural Products, 2021, 116, 1-36.	0.8	4
1752	species from south and south-eastern Australia (Ectocarpales, Phaeophyceae): a DNA barcoding approach. Australian Systematic Botany, 2021, 34, 587-594.	0.3	1
1754	Cryptic diversity and phylogeography of the <i>Rhabdophis nuchalis</i> group (Squamata: Colubridae). Molecular Phylogenetics and Evolution, 2022, 166, 107325.	1.2	4
1755	Evolution and phylogeny of the deep-sea isopod families Desmosomatidae Sars, 1897 and Nannoniscidae Hansen, 1916A (Isopoda: Asellota). Organisms Diversity and Evolution, 2021, , 1-27.	0.7	7
1756	Hidden Diversity Revealed in the Freshwater Snails, <i>Bythinella</i> and <i>Pseudamnicola</i> , in the Island of Crete. Integrative Zoology, 2021, , .	1.3	0
1757	Population genetics meets phylogenetics: new insights into the relationships among members of the genus <i>Euthynnus</i> (family Scombridae). Hydrobiologia, 2022, 849, 47-62.	1.0	5
1758	Molecular species delimitation of shrub frogs of the genus <i>Pseudophilautus</i> (Anura, Rhacophoridae). PLoS ONE, 2021, 16, e0258594.	1.1	6
1759	Improving the DNA Barcode Library of Mosquito Species With New Identifications and Discoveries in North-Central Argentina. Journal of Medical Entomology, 2022, 59, 173-183.	0.9	4

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1761	The origin of an extreme case of sisterâ€species sympatry in a palmâ€pollinator mutualistic system. <i>Journal of Biogeography</i> , 2021, 48, 3158-3169.	1.4	9
1762	Recent speciation and hybridization in Icelandic deepâ€sea isopods: An integrative approach using genomics and proteomics. <i>Molecular Ecology</i> , 2022, 31, 313-330.	2.0	15
1763	Efficacy and accuracy responses of DNA mini-barcodes in species identification under a supervised machine learning approach. , 2021, , .		1
1764	An Efficient Method for DNA-Based Species Assignment via Gene Tree and Species Tree Reconciliation. <i>Lecture Notes in Computer Science</i> , 2010, , 300-311.	1.0	6
1767	Phylogeny reconstruction: overview. , 2014, , 70-101.		0
1768	Comparison of phylogenetic methods and tests on trees. , 2014, , 153-181.		0
1769	Molecular clock and estimation of species divergence times. , 2014, , 361-389.		0
1770	Models of amino acid and codon substitution. , 2014, , 35-69.		0
1771	Maximum likelihood methods. , 2014, , 102-152.		0
1772	Bayesian theory. , 2014, , 182-213.		0
1773	Simulating molecular evolution. , 2014, , 418-441.		2
1774	Neutral and adaptive protein evolution. , 2014, , 390-417.		0
1776	Bayesian phylogenetics. , 2014, , 263-307.		0
1777	Coalescent theory and species trees. , 2014, , 308-360.		0
1778	Bayesian computation (MCMC). , 2014, , 214-262.		0
1779	Systematics and phylogeography of <i>Acanthodactylus schreiberi</i> and its relationships with <i>Acanthodactylus boskianus</i> (Reptilia: Squamata: Lacertidae). <i>Zoological Journal of the Linnean Society</i> , 2014, 172, 720-739.	1.0	8
1783	Organelle Genomes and Endosymbionts. , 2017, , 21-42.		0
1784	Genetic divergence across an oxygen minimum zone. <i>Marine Ecology - Progress Series</i> , 2017, 577, 79-91.	0.9	0

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1785	Isolation as a phylogeny-shaping factor: historical geology and cave habitats in the Mediterranean Truncatelloidea Gray, 1840 (Caenogastropoda). <i>Folia Malacologica</i> , 2017, 25, 231-229.	0.1	2
1786	DNA Barcoding of Ichthyoplankton and Juvenile Fishes of a Tropical River in Malaysia. , 2018, , 367-381.		0
1787	Combined effects of the microcystin MC-LR and temperature on the morphological features of <i>Brachionus calyciflorus</i> . <i>Hupo Kexue/Journal of Lake Sciences</i> , 2018, 30, 1027-1040.	0.3	0
1798	Genomic data show little geographical structure across the naturally fragmented range of the purple-gaped honeyeater. <i>Australian Journal of Zoology</i> , 2019, 67, 226.	0.6	1
1807	Molecular clocks, biogeography and species diversity in <i>Herichthys</i> with evaluation of the role of Punta del Morro as a vicariant brake along the Mexican Transition Zone in the context of local and global time frame of cichlid diversification. <i>PeerJ</i> , 2020, 8, e8818.	0.9	7
1808	A Revision of Philander (Marsupialia: Didelphidae), Part 2: Phylogenetic Relationships and Morphological Diagnosis of <i>P. nigratus</i> Thomas, 1923. <i>American Museum Novitates</i> , 2020, 2020, 1.	0.2	6
1809	Molecular Species Delimitation of the Genus <i>Reishia</i> (Mollusca: Gastropoda) along the Coasts of China and Korea. <i>Zoological Science</i> , 2020, 37, 382.	0.3	1
1812	Genetic Diversity of Ligidium Isopods in Hokkaido and Niigata, Northern Japan, Based on Mitochondrial DNA Analysis. <i>Zoological Science</i> , 2020, 37, 1.	0.3	4
1813	COI mtDNA barcoding and morphology for the description of a new species of ricinuleid of the genus <i>Pseudocellus</i> (Arachnida: Ricinulei: Ricinoididae) from El Triunfo Biosphere Reserve, Chiapas, Mexico. <i>European Journal of Taxonomy</i> , 0, 778, .	0.6	2
1817	Integrative taxonomy to assess the species richness of chalcidoid parasitoids (Hymenoptera) associated to Bruchinae (Coleoptera: Chrysomelidae) from Mexico. <i>Revista Mexicana De Biodiversidad</i> , 2020, 91, 913492.	0.4	4
1819	Systematic and diversity of ground beetles (Carabidae: Coleoptera) from district Haripur KPK, Pakistan. <i>Acta Entomology and Zoology</i> , 2020, 1, 05-07.	0.2	0
1820	A Molecular Approach to the Phylogeny of Theraphosidae and Their Kin. <i>Zoological Monographs</i> , 2020, , 25-75.	1.1	2
1821	Integrative systematics unveils the controversial identity of Engraulidae fishing stocks in a Neotropical estuary, northeast Brazil. <i>Neotropical Ichthyology</i> , 2020, 18, .	0.5	3
1822	<i>Gracilariopsis lemaneiformis</i> (Gracilariaceae, Rhodophyta) in the Mexican Coasts: A Case of Disjunct Distribution?. <i>American Journal of Plant Sciences</i> , 2020, 11, 111-124.	0.3	3
1823	Integrative taxonomy of the genus <i>Dudgeodes</i> Sartori, 2008 (Insecta, Ephemeroptera, Teloganodidae) from the Philippines with description of new species and supplementary descriptions of Southeast Asian species. <i>ZooKeys</i> , 2020, 910, 93-129.	0.5	11
1824	Cryptic diversity and gene introgression of Moinidae (Crustacea: Cladocera) in Nigeria. <i>Contributions To Zoology</i> , 2021, 90, 463-486.	0.2	2
1825	MtDNA species-level phylogeny and delimitation support significantly underestimated diversity and endemism in the largest Neotropical cichlid genus (Cichlidae: <i>Crenicichla</i>). <i>PeerJ</i> , 2021, 9, e12283.	0.9	6
1826	Genetic diversity in the <i>Diplosoma listerianum</i> complex (Ascidiacea: Didemnidae) from the Western Atlantic. <i>Systematics and Biodiversity</i> , 0, , 1-15.	0.5	1

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1827	Evolutionary history of <i>Massartella</i> (Ephemeroptera, Leptophlebiidae) suggests ancient vicariant event between biotas of the Pantepui and Atlantic Forest highlands. <i>Systematic Entomology</i> , 0, , .	1.7	1
1828	Fuzzy species borders of glacial survivalists in the Carpathian biodiversity hotspot revealed using a multimer approach. <i>Scientific Reports</i> , 2021, 11, 21629.	1.6	13
1829	Integrative systematics illuminates the relationships in two sponge-associated hydrozoan families (Capitata: Sphaerocorynidae and Zancleopsidae). <i>Contributions To Zoology</i> , 2021, 90, 487-525.	0.2	7
1830	A new species of <i>Epeorus</i> (Caucasiron) (Ephemeroptera, Heptageniidae) from Azerbaijan and Iran. <i>ZooKeys</i> , 2021, 1068, 13-26.	0.5	1
1834	Hidden or unnoticed? Multiple lines of evidence support the recognition of a new species of <i>Pseudocorynopoma</i> (Characidae: Corynopomini). <i>Journal of Fish Biology</i> , 2021, 98, 219-236.	0.7	6
1835	Molecular and phenotypic data reveal a new Amazonian species of pit vipers (Serpentes: Viperidae: Tj ETQq1 1 0.784314 rgBT /Overlo	0.2	5
1836	Addressing the Linnean shortfall in a cryptic species complex. <i>Zoological Journal of the Linnean Society</i> , 2021, 192, 277-305.	1.0	11
1837	The True Identity of the New World Iguanid Lizard <i>Müller and Hellmich 1932</i> (Iguania: Liolaemidae) and Description of a New Species in the Group. <i>Zoological Studies</i> , 2018, 57, e22.	0.3	7
1839	Cryptic diversity among populations of <i>Aegla</i> Leach, 1820 (Decapoda: Anomura: Aeglididae) from Tibagi River basin, Paraná state, Brazil, with descriptions of three new species. <i>Journal of Natural History</i> , 2021, 55, 2145-2171.	0.2	3
1840	Adding pieces to the puzzle: insights into diversity and distribution patterns of Cumacea (Crustacea: Tj ETQq1 1 0.784314 rgBT /Overlo	0.9	8
1841	Molecular phylogeny and phylogeography of the freshwater-fish genus <i>Pethia</i> (Teleostei: Cyprinidae) in Sri Lanka. <i>Bmc Ecology and Evolution</i> , 2021, 21, 203.	0.7	6
1842	Elevational Diversity Patterns of Green Lacewings (Neuroptera: Chrysopidae) Uncovered With DNA Barcoding in a Biodiversity Hotspot of Southwest China. <i>Frontiers in Ecology and Evolution</i> , 2021, 9, .	1.1	7
1843	Integrated analysis reveals a new species of <i>Corydoras</i> Lacépède, 1803 (Siluriformes: Callichthyidae) in the lower Iguassu River, Brazil. <i>Organisms Diversity and Evolution</i> , 0, , 1.	0.7	1
1844	The diversity of aphid parasitoids in East Africa and implications for biological control. <i>Pest Management Science</i> , 2022, 78, 1109-1116.	1.7	9
1845	DNA barcoding in Dorcadionini (Coleoptera, Cerambycidae) uncovers mitochondrial-morphological discordance and the hybridogenic origin of several subspecies. <i>Organisms Diversity and Evolution</i> , 2022, 22, 205-229.	0.7	7
1846	A new species of <i>Hoplias malabaricus</i> species complex (Characiformes: Tj ETQq1 1 0.784314 rgBT /Overlo <i>Journal of Fish Biology</i> , 2022, 100, 425-443.	0.7	2
1847	Validity of <i>Pampus liuorum</i> Liu & Li, 2013, Revealed by the DNA Barcoding of Pampus Fishes (Perciformes, Stromateidae). <i>Diversity</i> , 2021, 13, 618.	0.7	7
1848	Hidden biodiversity in microarthropods (Acari, Oribatida, Eremaeioidea, Caleremaeus). <i>Scientific Reports</i> , 2021, 11, 23123.	1.6	4

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1851	Molecular species delimitation of the genera <i>Anodus</i> , <i>Argonectes</i> , <i>Bivibranchia</i> and <i>Micromischodus</i> (Ostariophysi: Characiformes). <i>Neotropical Ichthyology</i> , 2021, 19, .	0.5	2
1852	The current approaches to the study of algae: DNA barcoding and DNA taxonomy. <i>Issues of Modern Algology</i> (Ð³Ð³/4Ð;Ñ€Ð³/4ÑÑ; ÑÐ³/4Ð²Ñ€ÐµÐ¹/4ÐµÐ¹/2Ð¹/2Ð³/4Ð¹ Ð°Ð»»Ñ€Ð³Ð³/4Ð»»Ð³/4Ð³Ð,Ð), 2021, , 124-130.	0.1	0
1853	Molecular analyses of flightless weevils <i>Chiloneus</i> from Sicily and adjoining islands revealed new synonymy (Coleoptera: Curculionidae). , 2021, 88, 948-958.		1
1854	Integrative taxonomy reveals a new species of Neotropical headstanding fish in genus <i>Schizodon</i> (Characiformes: Anostomidae). <i>Neotropical Ichthyology</i> , 2021, 19, .	0.5	9
1855	High-level taxonomic splitting in allopatric taxa causes confusion downstream: a revision of the nudibranch family <i>Ðjoryphellidae</i> . <i>Zoological Journal of the Linnean Society</i> , 2022, 196, 215-249.	1.0	6
1856	Extensive species diversification and marked geographic phylogenetic structure in the Mesoamerican genus <i>Stenopelmatus</i> (Orthoptera: Stenopelmatidae: Stenopelmatinae) revealed by mitochondrial and nuclear 3RAD data. <i>Invertebrate Systematics</i> , 2022, , .	0.5	0
1857	Does clustering of DNA barcodes agree with botanical classification directly at high taxonomic levels? Trees in French Guiana as a case study. <i>Molecular Ecology Resources</i> , 2022, 22, 1746-1761.	2.2	1
1858	Molecular assessment of the species of <i>Gracilariaceae</i> (Gracilariales, Rhodophyta) from the Yucatan Peninsula, Mexico, including two new records for the Mexican Atlantic. <i>Botanical Sciences</i> , 2022, 100, 493-505.	0.3	1
1859	A new cryptic species of <i>Pithecopus</i> (Anura, Phyllomedusidae) in north-eastern Brazil. <i>European Journal of Taxonomy</i> , 0, 723, 108-134.	0.6	9
1860	Shell features and anatomy of the springsnail genus <i>Radomaniola</i> (Caenogastropoda: Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 267 T of the Linnean Society, 2022, 196, 393-441.	1.0	5
1861	Characterization of the first <i>Wolbachia</i> from the genus <i>Scaptodrosophila</i> , a male-killer from the rainforest species <i>S. claytoni</i> . <i>Insect Science</i> , 2022, , .	1.5	0
1862	SPEDE sampler: an R Shiny application to assess how methodological choices and taxon sampling can affect Generalised Mixed Yule Coalescent (GMYC) output and interpretation. <i>Molecular Ecology Resources</i> , 2022, , .	2.2	1
1863	The Mitochondrial Genetic Diversity of the Olive Field Mouse <i>Abrothrix olivacea</i> (Cricetidae; Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 267 T Evolution, 2022, 29, 413-430.	1.0	2
1864	Assessing model adequacy leads to more robust phylogeographic inference. <i>Trends in Ecology and Evolution</i> , 2022, 37, 402-410.	4.2	7
1865	<i>Thainema</i> gen. nov. (Leptolyngbyaceae, Synechococcales): A new genus of simple trichal cyanobacteria isolated from a solar saltern environment in Thailand. <i>PLoS ONE</i> , 2022, 17, e0261682.	1.1	8
1866	Correct Species Identification and Its Implications for Conservation Using Haplonticidae (Crustacea, Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 267 T	1.2	13
1867	Using Species Groups to Approach the Large and Taxonomically Unresolved Freshwater Fish Family <i>Nemacheilidae</i> (Teleostei: Cypriniformes). <i>Biology</i> , 2022, 11, 175.	1.3	5
1869	Shedding light on species boundaries in small endogeic animals through an integrative approach: species delimitation in the centipede <i>Clinopodes carinthiacus</i> (Chilopoda: Geophilidae) in the south-eastern Alps. <i>Zoological Journal of the Linnean Society</i> , 2022, 196, 902-923.	1.0	5

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1870	Cryptic Species Diversity and Phylogenetic Relationship in the Rust Genus <i>Chrysomyxa</i> from China. <i>Journal of Fungi</i> (Basel, Switzerland), 2022, 8, 83.	1.5	6
1871	Phylogeny and phylogeography of <i>Diestramima</i> cave crickets (Orthoptera: Rhaphidophoridae): speciation driven by multiple dispersal and vicariance events. <i>Systematic Entomology</i> , 2022, 47, 179-201.	1.7	9
1872	A description of <i>Dendronotus shpataki</i> sp. nov. (Gastropoda: Nudibranchia) from the Sea of Japan: a contribution of citizen science to marine zoology. <i>Zoosystematica Rossica</i> , 2022, 31, 3-19.	0.2	0
1874	A description of <i>Dendronotus shpataki</i> sp. nov. (Gastropoda: Nudibranchia) from the Sea of Japan: a contribution of citizen science to marine zoology. <i>Zoosystematica Rossica</i> , 2022, 31, 3-19.	0.2	1
1875	A new PCR based molecular method for early and precise quantification of parasitization in the emerging olive pest <i>Dasineura oleae</i> . <i>Pest Management Science</i> , 2022, 78, 1842-1849.	1.7	6
1876	DNA Barcoding of Scavenging Amphipod Communities at Active and Inactive Hydrothermal Vents in the Indian Ocean. <i>Frontiers in Marine Science</i> , 2022, 8, .	1.2	5
1877	Phylogenetic relations and range history of jerboas of the Allactaginae subfamily (Dipodidae). <i>Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 502</i>	1.6	5
1878	Five New Species of the Lichen-Forming Fungal Genus <i>Peltula</i> from China. <i>Journal of Fungi</i> (Basel), Tj ETQq1 1 0.784314 rgBT /Overlock 1.5	1.5	3
1879	Hoplomertean larvae are planktonic predators that capture and devour active animal prey. <i>Invertebrate Biology</i> , 2022, 141, .	0.3	4
1880	Alternative prey mediate intraguild predation in the open field. <i>Pest Management Science</i> , 2022, 78, 3939-3946.	1.7	6
1881	Important alien and potential native invasive insect pests of key fruit trees in Sub-Saharan Africa: advances in sustainable pre- and post-harvest management approaches. <i>CABI Agriculture and Bioscience</i> , 2022, 3, .	1.1	5
1882	Molecular Analysis Reveals a High Diversity of Anopheline Mosquitoes in Yanomami Lands and the Pantanal Region of Brazil. <i>Genes</i> , 2021, 12, 1995.	1.0	2
1883	Pandora's Box in the Deep Sea "Intraspecific Diversity Patterns and Distribution of Two Congeneric Scavenging Amphipods. <i>Frontiers in Marine Science</i> , 2021, 8, .	1.2	11
1884	Global phylogeny and taxonomic reassessment of the lichen genus <i>Dendroscosticta</i> (Ascomycota: Peltigerales). <i>Taxon</i> , 2022, 71, 256-287.	0.4	3
1886	First molecular phylogeny and species delimitation of West Palaearctic <i>Pollenia</i> (Diptera). <i>Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 18</i>	1.0	5
1887	Population structuration and chromosomal features homogeneity in <i>Parodon nasus</i> (Characiformes). <i>Tj ETQq1 1 0.784314 rgBT /Overlock 0.5</i> <i>Ichthyology</i> , 2022, 20, .	0.5	2
1889	Global 16S rRNA diversity of provannid snail endosymbionts from Indo-Pacific deep-sea hydrothermal vents. <i>Environmental Microbiology Reports</i> , 2022, 14, 299-307.	1.0	8
1890	Sex pheromones and sex attractants of species within the genera <i>Idolus</i> Desbrochers des Loges and <i>Dalopius</i> Eschscholtz (Coleoptera: Elateridae) in the western United States. <i>Agricultural and Forest Entomology</i> , 2022, 24, 301-309.	0.7	2

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1891	The Phyllosymbiosis Pattern Between the Fig Wasps of the Same Genus and Their Associated Microbiota. <i>Frontiers in Microbiology</i> , 2021, 12, 800190.	1.5	1
1892	Phylogeny and Cryptic Diversity of <i>Diopatira</i> (Onuphidae, Annelida) in the East Atlantic. <i>Biology</i> , 2022, 11, 327.	1.3	5
1893	Multiple Recent Colonizations of the Australian Region by the <i>Chydorus sphaericus</i> Group (Crustacea: Tj ETQq0 0 0 rgBT /Overlock 10 T	1.2	8
1894	Pelagic larval polyclads that practice macrophagous carnivory. <i>Invertebrate Biology</i> , 2022, 141, .	0.3	2
1895	Does monitoring of saproxylic beetles benefit from inclusion of larvae?. <i>Insect Conservation and Diversity</i> , 2022, 15, 555-571.	1.4	3
1896	Revisiting taxonomic disparities in the genus <i>Naemorhedus</i> : new insights from Indian Himalayan Region. <i>Mammalia</i> , 2022, 86, 373-379.	0.3	4
1897	Ten new species of <i>Ulva</i> (Ulvophyceae, Chlorophyta) discovered in New Caledonia: genetic and morphological diversity, and bloom potential. <i>European Journal of Phycology</i> , 2022, 57, 458-478.	0.9	13
1898	The Multi-Ocean Distribution of the Hadal Amphipod, <i>Hirondellea dubia</i> Dahl, 1959 (Crustacea,) Tj ETQq1 1 0.784314 rgBT /Overlock 10	1.2	8
1899	Past climate cooling promoted global dispersal of amphipods from Tian Shan montane lakes to circumboreal lakes. <i>Global Change Biology</i> , 2022, 28, 3830-3845.	4.2	10
1900	DNA barcoding and phylogeography of the <i>Hoplias malabaricus</i> species complex. <i>Scientific Reports</i> , 2022, 12, 5288.	1.6	5
1901	Colonization rather than fragmentation explains the geographical distribution and diversification of treefrogs endemic to Brazilian shield sky islands. <i>Journal of Biogeography</i> , 2022, 49, 682-698.	1.4	5
1902	Phylogeny, species delimitation and ecological and morphological diversity of <i>Characithecium</i> (Monogenoidea: Dactylogyridae). <i>Parasitology</i> , 2022, , 1-17.	0.7	1
1903	A new obligate groundwater species of <i>Asellus</i> (Isopoda, Asellidae) from Iran. <i>Subterranean Biology</i> , 20, 42, 97-124.	5.0	4
1904	Evolutionary Genetics of Cacti: Research Biases, Advances and Prospects. <i>Genes</i> , 2022, 13, 452.	1.0	6
1905	Analysis of biodiversity data suggests that mammal species are hidden in predictable places. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2022, 119, e2103400119.	3.3	13
1906	Diversity investigation by application of DNA barcoding: A case study of lepidopteran insects in Xinjiang wild fruit forests, China. <i>Ecology and Evolution</i> , 2022, 12, e8678.	0.8	0
1907	Cryptic species of pondweeds (Potamogetonaceae) at an intercontinental scale revealed by molecular phylogenetic analyses. <i>Taxon</i> , 2022, 71, 531-551.	0.4	3
1908	Drivers of unique and asynchronous population dynamics in Malagasy herpetofauna. <i>Journal of Biogeography</i> , 2022, 49, 600-616.	1.4	7

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1909	A new Amazonian species of <i>Allobates</i> Zimmermann & Zimmermann, 1988 (Aromobatidae) with a trilled advertisement call. PeerJ, 2022, 10, e13026.	0.9	3
1911	Genome Skimming Contributes to Clarifying Species Limits in Paris Section Axiparis (Melanthiaceae). Frontiers in Plant Science, 2022, 13, 832034.	1.7	5
1912	First record of <i>Pheretima vungtauensis</i> (Clitellata: Megascolecidae) in India and its phylogenetic relationship with <i>Metaphire houlleti</i> . , 0, , 1.		0
1913	An Annotated Checklist of Recent Opossums (Mammalia: Didelphidae). Bulletin of the American Museum of Natural History, 2022, 455, .	1.2	10
1914	Geographic distance, sedimentation, and substrate shape cryptic crustose coralline algal assemblages in the world's largest subtropical intertidal algal reef. Molecular Ecology, 2022, 31, 3056-3071.	2.0	6
1915	Population genomics supports multiple hybrid zone origins of socially hybridogenetic lineages of <i>Pogonomyrmex</i> harvester ants. Evolution; International Journal of Organic Evolution, 2022, 76, 1016-1032.	1.1	4
1916	Diversity, biogeography, and reproductive evolution in the genus <i>Pipa</i> (Amphibia: Anura: Pipidae). Molecular Phylogenetics and Evolution, 2022, 170, 107442.	1.2	11
1917	Diversity of <i>Brachionus plicatilis</i> species complex (Rotifera) in inland saline waters from China: Presence of a new mitochondrial clade on the Tibetan Plateau. Molecular Phylogenetics and Evolution, 2022, 171, 107457.	1.2	4
1918	A taxonomically complex catfish group from an underrepresented geographic area: Systematics and species limits in <i>Hypostomus</i> Lacépède, 1803 (Siluriformes, Loricariidae) from Eastern South America. Journal of Zoological Systematics and Evolutionary Research, 2021, 59, 1994-2009.	0.6	2
1919	A molecular and chromosomic meta-analysis approach and its implications for the taxonomy of the genus <i>Makalata</i> Husson, 1978 (Rodentia, Echimyidae) including an amended diagnosis for <i>M. macrura</i> (Wagner, 1842). Journal of Zoological Systematics and Evolutionary Research, 2021, 59, 2387-2409.	0.6	0
1920	Complete Genome Sequence, Molecular Characterization and Phylogenetic Relationships of a Novel Tern Atadenovirus. Microorganisms, 2022, 10, 31.	1.6	3
1921	Phylogenetic Relationships, Speciation, and Origin of <i>Armillaria</i> in the Northern Hemisphere: A Lesson Based on rRNA and Elongation Factor 1-Alpha. Journal of Fungi (Basel, Switzerland), 2021, 7, 1088.	1.5	8
1922	Mersin Üniversitesi Su Ürünleri Fakültesi Uygulama Birimleri'nde Rejeneratif Tıp Araştırmaları'nda Model Organizma Olarak Yetiştirilen <i>Ambystoma mexicanum</i> 'un DNA Barkodlaması ve Filogenisi. Commagene Journal of Biology, 0, , 161-176.	0.1	0
1923	Species delimitation of rice seed bugs complex: Insights from mitochondrial genomes and ddRAD-seq data. Zoologica Scripta, 2022, 51, 185-198.	0.7	5
1924	Postulating the Modality of Integrative Taxonomy in Describing the Cryptic Congener <i>Pampus griseus</i> (Cuvier) and Systematics of the Genus <i>Pampus</i> (Perciformes: Stromateidae). Frontiers in Marine Science, 2021, 8, .	1.2	4
1925	Uncovering Hidden Diversity: Three New Species of the <i>Keratella</i> Genus (Rotifera, Monogononta.) Tj ETQq1 1 0.784314 rgBT /Overl	0.7	4
1926	Molecular evidence for extensive discontinuity between peracarid (Crustacea) fauna of Macaronesian islands and nearby continental coasts: over fifty candidate endemic species. Marine Biology, 2022, 169, 1.	0.7	5
1927	<sc>DNA</sc>-based assessment of environmental degradation in an unknown fauna: The freshwater macroinvertebrates of the <sc>Indo-Burmese</sc> hotspot. Journal of Applied Ecology, 2022, 59, 1644-1658.	1.9	2

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1928	Stripes Matter: Integrative Systematics of <i>Coryphellina rubrolineata</i> Species Complex (Gastropoda: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5	0.7	5
1929	Reassessing fish diversity of Penang Island's freshwaters (northwest Peninsular Malaysia) through a molecular approach raises questions on its conservation status. <i>Biodiversity and Conservation</i> , 0, , 1.	1.2	0
1930	Lack of Statistical Rigor in DNA Barcoding Likely Invalidates the Presence of a True Species' Barcode Gap. <i>Frontiers in Ecology and Evolution</i> , 2022, 10, .	1.1	10
1931	Identifying cryptic fern gametophytes using DNA barcoding: A review. <i>Applications in Plant Sciences</i> , 2022, 10, e11465.	0.8	8
1966	Analysing diversification dynamics using barcoding data: The case of an obligate mycorrhizal symbiont. <i>Molecular Ecology</i> , 2022, 31, 3496-3512.	2.0	6
1967	Examination of the generic concept and species boundaries of the genus <i>Erioscyphella</i> (Lachnaceae,) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 5 materials. <i>MycoKeys</i> , 2022, 87, 1-52.	0.8	0
1968	Molecular data reveal multiple lineages of <i>Scinax nebulosus</i> (Spix, 1824) (Anura: Hylidae) with Plio-Pleistocene diversification in different Brazilian regions. <i>Anais Da Academia Brasileira De Ciencias</i> , 2022, 94, e20200733.	0.3	1
1969	Systematics and phylogeography of western Mediterranean tarantulas (Araneae: Theraphosidae). <i>Zoological Journal of the Linnean Society</i> , 2022, 196, 845-884.	1.0	8
1970	Species limits and recent diversification of <i>Cerradomys</i> (Sigmodontinae: Oryzomyini) during the Pleistocene. <i>PeerJ</i> , 2022, 10, e13011.	0.9	3
1971	<i>Microglena antarctica</i> sp. nov. a New Antarctic Green Alga from Inexpressible Island (Terra Nova Bay,) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 5	0.7	5
1972	Phylogenetic placement and species delimitation of the crab spider genus <i>Phrynarachne</i> (Araneae:) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5	1.2	2
1973	Genetic differentiation and phylogeography of rotifer <i>Polyarthra dolichoptera</i> and <i>P. vulgaris</i> populations between Southeastern China and eastern North America: High intercontinental differences. <i>Ecology and Evolution</i> , 2022, 12, e8912.	0.8	4
1974	Four in One: Cryptic Diversity in Geoffroy's Side-Necked Turtle <i>Phrynops geoffroanus</i> (Schweigger) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5	0.7	2
1975	DNA Barcoding of Cold-Water Coral-Associated Ophiuroid Fauna from the North Atlantic. <i>Diversity</i> , 2022, 14, 358.	0.7	2
1976	Phylogeography of the freshwater rotifer <i>Brachionus calyciflorus</i> species complex in China. <i>Hydrobiologia</i> , 2022, 849, 2813-2829.	1.0	4
1977	Coalescent-based species delimitation in North American pinyon pines using low-copy nuclear genes and plastomes. <i>American Journal of Botany</i> , 2022, 109, 706-726.	0.8	3
1978	A revision of the red algal genus <i>Ahnfeltia</i> on the Russian coast of the North Pacific. <i>Phycologia</i> , 0, , 1-7.	0.6	1
1979	Phylogenetics in space: How continuous spatial structure impacts tree inference. <i>Molecular Phylogenetics and Evolution</i> , 2022, 173, 107505.	1.2	0

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1980	Uncovering overlooked diversity using molecular phylogenetic approach: A case of Japanese sphaeriid clams (<i>Bivalvia</i> : <i>Sphaeriidae</i>). <i>Molecular Phylogenetics and Evolution</i> , 2022, 173, 107508.	1.2	5
1981	Mitochondrial discordance and patterns of reproductive isolation in a complex of simultaneously hermaphroditic species, the <i>Alloobophora chlorotica</i> case study. <i>Journal of Evolutionary Biology</i> , 2022, 35, 831-843.	0.8	1
1982	Disentangling phylogenetic relations and biogeographic history within the <i>Cucujus haematodes</i> species group (Coleoptera: Cucujidae). <i>Molecular Phylogenetics and Evolution</i> , 2022, 173, 107527.	1.2	1
1983	Recurrent founder-event speciation across the Mediterranean likely shaped the species diversity and geographic distribution of the freshwater snail genus <i>Mercuria</i> Boeters, 1971 (Caenogastropoda). <i>Trends in Ecology and Evolution</i> , 2022, 33, 107-114.	0.784314	14
1984	DNA barcoding and metabarcoding of highly diverse aquatic mites (Acarina) can improve their use in routine biological monitoring. <i>Marine and Freshwater Research</i> , 2022, 73, 900-914.	0.7	4
1985	Evolution and Biogeography of Freshwater Snails of the Genus <i>Bulinus</i> (Gastropoda) in Afrotropical Extreme Environments. <i>Frontiers in Environmental Science</i> , 2022, 10, .	1.5	6
1986	Life without blood: Molecular and functional analysis of hirudins and hirudin-like factors of the Asian non-hematophagous leech <i>Whitmania pigra</i> . <i>Journal of Thrombosis and Haemostasis</i> , 2022, 20, 1808-1817.	1.9	8
1987	Unveiling the <i>Mycodrosophila projectans</i> (Diptera, Drosophilidae) species complex: Insights into the evolution of three Neotropical cryptic and syntopic species. <i>PLoS ONE</i> , 2022, 17, e0268657.	1.1	1
1988	The taxonomy and phylogeny of the <i>Cyrtodactylus brevipalmatus</i> group (Squamata: Gekkonidae) with emphasis on <i>C. interdigitalis</i> and <i>C. ngati</i> . <i>Vertebrate Zoology</i> , 0, 72, 245-269.	2.0	5
1989	Is what you see what you get? The relationship between field observed and laboratory observed aphid parasitism rates in canola fields. <i>Pest Management Science</i> , 2022, 78, 3596-3607.	1.7	3
1990	DNA barcoding and species delimitation of the genus <i>Oxynoemacheilus</i> (Teleostei). <i>Trends in Ecology and Evolution</i> , 2022, 33, 107-114.	0.7	5
1991	DNA barcoding reveals deep divergent molecular units in <i>Pomatomus saltatrix</i> (Perciformes). <i>Trends in Ecology and Evolution</i> , 2022, 33, 107-114.	0.4	2
1992	Integrative taxonomy on the rare sky-island <i>Ligidium</i> species from southwest China (Isopoda). <i>Trends in Ecology and Evolution</i> , 2022, 33, 107-114.	0.3	4
1993	Ancient Tethyan Vicariance and Long-Distance Dispersal Drive Global Diversification and Cryptic Speciation in the Red Seaweed <i>Pterocladia</i> . <i>Frontiers in Plant Science</i> , 2022, 13, .	1.7	7
1995	Species diversity of the freshwater red algal genus <i>Kumanoa</i> in Taiwan with the description of two new species: <i>Kumanoa taiwanensis</i> sp. nov. and <i>Kumanoa yuanyangensis</i> sp. nov. <i>Phycologia</i> , 0, , 1-14.	0.6	0
1996	Unravelling the species diversity, phylogeny and biogeography of the mycoheterotrophic <i>Voyriaceae</i> (Gentianaceae) and the description of a new species. <i>Taxon</i> , 0, , .	0.4	0
1998	Life history strategies of <i>Cotylurus</i> spp. Szidat, 1928 (Trematoda, Strigeidae) in the molecular era – Evolutionary consequences and implications for taxonomy. <i>International Journal for Parasitology: Parasites and Wildlife</i> , 2022, 18, 201-211.	0.6	0
1999	New insights into the genetic diversity of the Balkan bush-cricket of the <i>Poecilimon ornatus</i> group (Orthoptera: Tettigoniidae). <i>Arthropod Systematics and Phylogeny</i> , 0, 80, 243-259.	5.5	0

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2000	Impact of mitotype diversity on metabarcoding biodiversity estimations in Insecta and Arachnida using different sample preparation strategies. <i>Molecular Ecology Resources</i> , 0, , .	2.2	1
2001	Resolved and Redeemed: A New Fleck to the Evolutionary Divergence in the Genus <i>Scomberomorus</i> Lacepède, 1801 (Scombridae) With Cryptic Speciation. <i>Frontiers in Marine Science</i> , 0, 9, .	1.2	5
2002	Genetic diversity of sympatric <i>Schizymenia dubyi</i> and <i>S. apoda</i> (Schizymeniaceae). <i>Journal of Phycology</i> , 38, 10, 150-160.	0.6	0
2003	Abyssal vent field habitats along plate margins in the Central Indian Ocean yield new species in the genus <i>Anatoma</i> (Vetigastropoda: Anatomidae). <i>European Journal of Taxonomy</i> , 0, 826, 135-162.	0.6	0
2004	Systematic Review of the Genus <i>Nalepa</i> Reitter, 1887 (Coleoptera, Tenebrionidae, Blaptinae, Blaptini) from the Tibetan Plateau, with Description of Six New Species and Two Larvae. <i>Insects</i> , 2022, 13, 598.	1.0	5
2005	Seven snail species hidden in one: Biogeographic diversity in an apparently widespread periwinkle in the Southern Ocean. <i>Journal of Biogeography</i> , 2022, 49, 1521-1534.	1.4	9
2006	Delimiting the cryptic diversity and host preferences of <i>Sycophila</i> parasitoid wasps associated with oak galls using phylogenomic data. <i>Molecular Ecology</i> , 2022, 31, 4417-4433.	2.0	11
2007	Secondary reversion to sexual monomorphism associated with tissue-specific loss of <i>doublesex</i> expression. <i>Evolution; International Journal of Organic Evolution</i> , 2022, 76, 2089-2104.	1.1	3
2008	A DNA barcode reference library for endemic Ponto-Caspian amphipods. <i>Scientific Reports</i> , 2022, 12, .	1.6	16
2009	A new species of <i>Astronotus</i> (Teleostei, Cichlidae) from the Orinoco River and Gulf of Paria basins, northern South America. <i>ZooKeys</i> , 0, 1113, 111-152.	0.5	1
2010	The Biodiversity of Calcaxonian Octocorals from the Irish Continental Slope Inferred from Multilocus Mitochondrial Barcoding. <i>Diversity</i> , 2022, 14, 576.	0.7	4
2011	Phylogeny and Integrative Taxonomy of the Genera <i>Gymnaetoides</i> and <i>Pseudotachycines</i> (Orthoptera: Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 6	1.0	1
2012	An Integrative Taxonomic Revision of Aneuraceae H.Klinggr. (Marchantiophyta) from Guadeloupe and Martinique, French West Indies. <i>Cryptogamie, Bryologie</i> , 2022, 43, .	0.1	0
2013	Molecular systematics and biogeographic insights of the <i>Calomys callosus</i> complex (Rodentia). <i>Journal of Biogeography</i> , 2022, 49, 1521-1534.	0.7	2
2014	Out of southern Africa: Origins and cryptic speciation in <i>Chamaeleo</i> , the most widespread chameleon genus. <i>Molecular Phylogenetics and Evolution</i> , 2022, 175, 107578.	1.2	4
2015	<i>Ceriodaphnia</i> (Cladocera: Daphniidae) in China: Lineage diversity, phylogeography and possible interspecific hybridization. <i>Molecular Phylogenetics and Evolution</i> , 2022, 175, 107586.	1.2	5
2016	Phylogeography of Chinese cereal cyst nematodes sheds lights on their origin and dispersal. <i>Evolutionary Applications</i> , 2022, 15, 1236-1248.	1.5	3
2017	Species diversity and systematics of the <i>Leptodactylus melanonotus</i> group (Anura). <i>Systematics and Biodiversity</i> , 2022, 20, 1-31.	0.5	3

#	ARTICLE	IF	CITATIONS
2018	Distinguishing Long-Discussed Cryptic Species of the Epibiotic Goose-Neck Barnacle of the Genus <i>Conchoderma</i> (Thoracalcaea: Lepadidae) with Integrative Taxonomy. <i>Diversity</i> , 2022, 14, 593.	0.7	0
2019	The Application of DNA Barcoding in Crustacean Larvae Identification from the Zhongsha Islands, South China. <i>Frontiers in Marine Science</i> , 0, 9, .	1.2	2
2020	DNA Barcoding and Species Delimitation for Dogfish Sharks Belonging to the <i>Squalus</i> Genus (Squaliformes: Squalidae). <i>Diversity</i> , 2022, 14, 544.	0.7	5
2024	Resolving the phylogenetic relationship between <i>Parmotrema crinitum</i> and <i>Parmotrema perlatum</i> populations. <i>Lichenologist</i> , 2022, 54, 183-194.	0.5	1
2025	Annotated checklist and genetic data for parasitic helminths infecting New Zealand marine invertebrates. <i>Invertebrate Biology</i> , 2022, 141, .	0.3	5
2026	Predicting Species Boundaries and Assessing Undescribed Diversity in <i>Pneumocystis</i> , an Obligate Lung Symbiont. <i>Journal of Fungi (Basel, Switzerland)</i> , 2022, 8, 799.	1.5	1
2027	Assessing diversity of King Crab <i>Lithodes</i> spp. in the south-eastern pacific using phylogeny and molecular species delimitation methods. <i>Ecology and Evolution</i> , 2022, 12, .	0.8	1
2028	Multi-gene phylogeny of North American clear-winged moths (Lepidoptera: Sesiidae): a foundation for future evolutionary study of a speciose mimicry complex. <i>Cladistics</i> , 2023, 39, 1-17.	1.5	2
2029	Narrowly defined taxa on a global scale: The phylogeny and taxonomy of the genera <i>Catriona</i> and <i>Tenellia</i> (Nudibranchia, Trinchetiidae) favours fine-scale taxonomic differentiation and dissolution of the 'lumpers & splitters' dilemma. <i>Evolutionary Applications</i> , 2023, 16, 428-460.	1.5	4
2030	Late Cenozoic environmental changes drove the diversification of a weevil genus endemic to the Cape Floristic Region. <i>Zoologica Scripta</i> , 2022, 51, 724-740.	0.7	5
2031	Integrative systematics of the genus <i>Dondice</i> Marcus, 1958 (Gastropoda, Nudibranchia, Myrrhinidae) in the Western Atlantic. <i>Marine Biodiversity</i> , 2022, 52, .	0.3	2
2032	Cryptic subterranean diversity: regional phylogeography of the sand termite <i>Psammotermes allocerus</i> Silvestri, 1908 in the wider Namib region. <i>Organisms Diversity and Evolution</i> , 0, , .	0.7	0
2033	A new nurse frog of the <i>Allobates tapajo</i> species complex (Anura: Aromobatidae) from the upper Madeira River, Brazilian Amazonia. <i>PeerJ</i> , 0, 10, e13751.	0.9	5
2034	Cladocera (Crustacea: Branchiopoda) of Man-Made Lakes at the Northeast Part of the United Arab Emirates with a Hypothesis on Their Origin. <i>Diversity</i> , 2022, 14, 688.	0.7	1
2035	A tale of two bellies: systematics of the oval frogs (Anura: Microhylidae: <i>Elachistocleis</i>). <i>Zoological Journal of the Linnean Society</i> , 2023, 197, 545-568.	1.0	0
2037	Molecular techniques for the taxonomy of <i>Aedes</i> Meigen, 1818 (Culicidae: Aedini): A review of studies from 2010 to 2021. <i>Acta Tropica</i> , 2022, 236, 106694.	0.9	3
2038	Integrative taxonomy reveals a new species of <i>Cyphocharax</i> (Characiformes: Curimatidae) from the Upper Paraíba do Sul River basin, Brazil. <i>Neotropical Ichthyology</i> , 2022, 20, .	0.5	1
2039	New insights into the genetic variability of <i>Fasciola hepatica</i> (Trematoda) in Algeria and relationships with other geographic regions revealed by mitochondrial DNA. <i>Helminthologia</i> , 2022, 59, 152-164.	0.3	1

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2040	Molecular Characterization of <i>Anopheles algeriensis</i> Theobald, 1903 (Diptera: Culicidae) Populations from Europe. <i>Pathogens</i> , 2022, 11, 990.	1.2	1
2041	Sympatric morphotypes of the restricted-range Tashan Cave Garra: distinct species or a case of phenotypic plasticity?. <i>Environmental Biology of Fishes</i> , 2022, 105, 1251-1260.	0.4	5
2042	<scp>DNA</scp> metabarcoding of gut contents reveals key habitat and seasonal drivers of trophic networks involving generalist predators in agricultural landscapes. <i>Pest Management Science</i> , 2022, 78, 5390-5401.	1.7	4
2043	Coexisting Cryptic Species as a Model System in Integrative Taxonomy. , 2022, , 169-196.		3
2044	Disentangling the identity of <i>Lebertia porosa</i> Thor, 1900 using integrative taxonomy (Acari: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 582 T	0.6	3
2045	Long-read Sequencing Data Reveals Dynamic Evolution of Mitochondrial Genome Size and the Phylogenetic Utility of Mitochondrial DNA in Hercules Beetles (<i>Dynastes</i>; Scarabaeidae). <i>Genome Biology and Evolution</i> , 2022, 14, .	1.1	5
2046	Multiples lines of evidence unveil cryptic diversity in the<i>LophostomaÂbrasiliense</i>(Chiroptera:) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5	0.5	3
2047	Molecular Studies of Relationships and Identifications Among Insects of the Subfamily Panchaetothripinae (Thysanoptera, Thripidae). <i>Journal of Insect Science</i> , 2022, 22, .	0.6	1
2048	Diversification in Caucasian <i>Epeorus</i> (<i>Caucasiron</i>) mayflies (Ephemeroptera:) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 427 T <i>Entomology</i> , 2022, 47, 603-617.	1.7	1
2049	Rapid and sensitive onâ€site genetic diagnostics of pest fruit flies using <scp>CRISPRâ€Cas12a</scp>. <i>Pest Management Science</i> , 2023, 79, 68-75.	1.7	7
2050	Remarks on phylogeny and molecular variations of criconematid species (Nematoda: Criconematidae) with case studies from Vietnam. <i>Scientific Reports</i> , 2022, 12, .	1.6	5
2051	ï»¿Pinctada phuketensis sp. nov. (Bivalvia, Ostreida, Margaritidae), a new pearl oyster species from Phuket, western coast of Thailand. <i>ZooKeys</i> , 0, 1119, 181-195.	0.5	2
2052	New Record of <i>Anisopteromalus calandrae</i> 1 at Veracruz, Mexico: A Parasitoid of <i>Sitophilus zeamais</i> 2. <i>Southwestern Entomologist</i> , 2022, 47, .	0.1	0
2053	ï»¿A DNA barcode library for katydids, cave crickets, and leaf-rolling crickets (Tettigoniidae,) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 5	0.5	1
2054	Phylogenetics, integrative taxonomy and systematics of the <i>Sesamia cretica</i> species group (Lepidoptera: Noctuidae: Apameini: Sesamiina), with the description of 21 new species from the Afrotropical region. <i>Annales De La Societe Entomologique De France</i> , 2022, 58, 387-454.	0.4	0
2055	Specific and Intraspecific Diversity of Symphypleona and Neelipleona (Hexapoda: Collembola) in Southern High Appalachia (USA). <i>Diversity</i> , 2022, 14, 847.	0.7	4
2056	Multilocus species delimitation analyses show junior synonyms and deep-sea unknown species of genus <i>Gaidropsarus</i> (Teleostei: Gadiformes) in the North Atlantic/Mediterranean Sea area. <i>Marine Biology</i> , 2022, 169, .	0.7	2
2058	Unrecognized species diversity and endemism in the cichlid genus <i>Bujurquina</i> (Teleostei: Cichlidae) together with a molecular phylogeny document large-scale transformation of the western Amazonian river network and reveal complex paleogeography of the Ecuadorian Amazon. <i>Hydrobiologia</i> , 2023, 850, 2199-2229.	1.0	2

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2059	Hidden diversity of the genus <i>Trinomys</i> (Rodentia: Echimyidae): phylogenetic and populational structure analyses uncover putative new lineages. <i>Zoological Journal of the Linnean Society</i> , 0, , .	1.0	0
2060	Taxonomic revision of the Australian stick insect genus <i>Candovia</i> (Phasmida: Necrosciinae): insight from molecular systematics and species-delimitation approaches. <i>Zoological Journal of the Linnean Society</i> , 0, , .	1.0	1
2061	DNA barcoding echinoderms from the East Coast of South Africa. The challenge to maintain DNA data connected with taxonomy. <i>PLoS ONE</i> , 2022, 17, e0270321.	1.1	2
2062	Species delimitation in the genus <i>Ochraethes</i> Chevrolat, 1860 (Coleoptera: Cerambycidae), with description of two new species. <i>European Journal of Taxonomy</i> , 0, 845, .	0.6	0
2063	OTU Delimitation with Earthworm DNA Barcodes: A Comparison of Methods. <i>Diversity</i> , 2022, 14, 866.	0.7	9
2064	Extremely divergent <i>COI</i> sequences within an amphipod species complex: A possible role for endosymbionts?. <i>Ecology and Evolution</i> , 2022, 12, .	0.8	2
2065	Historical DNA of rare yellow-eared bats <i>Vampyressa</i> Thomas, 1900 (Chiroptera, Phyllostomidae) clarifies phylogeny and species boundaries within the genus. <i>Systematics and Biodiversity</i> , 2022, 20, 1-13.	0.5	1
2067	Diversification of the <i>Pristimantis conspicillatus</i> group (Anura: Craugastoridae) within distinct neotropical areas throughout the Neogene. <i>Systematics and Biodiversity</i> , 2022, 20, 1-16.	0.5	7
2068	Molecular and morphological characterization of <i>Digenea</i> (Rhodomelaceae, Rhodophyta) in the Mexican Atlantic. <i>Botanica Marina</i> , 2022, .	0.6	0
2069	Cryptic diversity of <i>Oxythyrea</i> flower chafers and its implication for conservation of non-forest biotopes in the Balkans. <i>Insect Conservation and Diversity</i> , 0, , .	1.4	1
2070	Phylogeography and evolutionary lineage diversity in the small-eared greater galago, <i>Otolemur garnettii</i> (Primates: Galagidae). <i>Zoological Journal of the Linnean Society</i> , 0, , .	1.0	1
2071	Disentangling a Neotropical pest species complex: Genetic diversity and population structure of the native rice stink bug <i>Oebalus poecilus</i> and the invasive <i>O. ypsilon-griseus</i> . <i>Pest Management Science</i> , 0, , .	1.7	0
2072	New Botrylloides, Botryllus, and Symplegma (Ascidiacea: Styelidae) in Coral Reefs of the Southern Gulf of Mexico and Mexican Caribbean Sea. <i>Diversity</i> , 2022, 14, 977.	0.7	6
2073	Historical biogeography highlights the role of Miocene landscape changes on the diversification of a clade of Amazonian tree frogs. <i>Organisms Diversity and Evolution</i> , 2023, 23, 395-414.	0.7	7
2074	Towards a genetic theory of island biogeography: Inferring processes from multidimensional community-scale data. <i>Global Ecology and Biogeography</i> , 2023, 32, 4-23.	2.7	2
2075	Genome-Based Taxa Delimitation (GBTD): A New Approach. <i>Diversity</i> , 2022, 14, 948.	0.7	1
2076	Integrative taxonomy of the stalk-eyed bug genus <i>Chauliops</i> (Heteroptera: Malcidae): Tj ETQq0 0 0 rBT /Overlock 10 Tf 50 107 932-947.	1.6	2
2077	Ecological speciation of Japanese hedgehog mushroom: <i>Hydnum subalpinum</i> sp. nov. is distinguished from its sister species <i>H. repando-orientale</i> by means of integrative taxonomy. <i>Mycological Progress</i> , 2022, 21, .	0.5	2

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2078	Diversity of <i>Siphonaria</i> Sowerby I, 1823 (Gastropoda, Siphonariidae) in the Seychelles Bank and beyond. <i>Zoologica Scripta</i> , 0, , .	0.7	0
2079	Uncertainties and risks in delimiting species of <i>Cambeva</i> (Siluriformes: Trichomycteridae) with single-locus methods and geographically restricted data. <i>Neotropical Ichthyology</i> , 2022, 20, .	0.5	3
2080	Two new species of <i>Centroptilum</i> Eaton, 1869 from North Africa (Ephemeroptera, Baetidae). <i>ZooKeys</i> , 0, 1131, 71-97.	0.5	8
2081	Unweaving a hard taxonomic knot in coral reef dwellers: integrative systematics reveals two parallel cryptic species complexes in marbled shrimps of the genus <i>Saron</i> Thallwitz 1891 (Caridea). <i>TJ ETQq1 1 0.764814 rgBT /Overlock</i>	1.0	14
2082	Standardized nuclear markers improve and homogenize species delimitation in Metazoa. <i>Methods in Ecology and Evolution</i> , 2023, 14, 543-555.	2.2	13
2083	Multilocus phylogeny and species delimitation suggest synonymies of two <i>Lucanus</i> Scopoli, 1763 (Coleoptera, Lucanidae) species names. <i>ZooKeys</i> , 0, 1135, 139-155.	0.5	1
2084	To be or not to be? Integrative taxonomy and species delimitation in the daddy long-legs spiders of the genus <i>Physocyclus</i> (Araneae, Pholcidae) using DNA barcoding and morphology. <i>ZooKeys</i> , 0, 1135, 93-118.	0.5	7
2085	Species diversity and distribution of genus <i>Pampus</i> (Pelagiaria: Stromateidae) based on global mitochondrial data. <i>Frontiers in Marine Science</i> , 0, 9, .	1.2	2
2086	<i>Cucullanus pinnai pinnai</i> and <i>C. pinnai pterodorasi</i> (Nematoda Cucullanidae): what does the integrative taxonomy tell us about these species and subspecies classification?. <i>Parasitology Research</i> , 2023, 122, 557-569.	0.6	1
2087	Comparing the Efficiency of Single-Locus Species Delimitation Methods within Trochoidea (Gastropoda: Vetigastropoda). <i>Genes</i> , 2022, 13, 2273.	1.0	6
2088	A new species of <i>Odontophrynus</i> (Anura, Odontophrynidae) from the southern portion of the Mantiqueira mountains. <i>European Journal of Taxonomy</i> , 0, 847, .	0.6	1
2089	Evaluating species richness using proteomic fingerprinting and DNA barcoding—a case study on meiobenthic copepods from the Clarion Clipperton Fracture Zone. <i>Marine Biodiversity</i> , 2022, 52, .	0.3	1
2090	DNA barcoding unveils a high diversity of caddisflies (Trichoptera) in the Mount Halimun Salak National Park (West Java; Indonesia). <i>PeerJ</i> , 0, 10, e14182.	0.9	2
2091	Quality control of fighting fish nucleotide sequences in public repositories reveals a dark matter of systematic taxonomic implication. <i>Genes and Genomics</i> , 2023, 45, 169-181.	0.5	2
2092	Diversity of interstitial nemertean of the genus <i>Ototyphlonemertes</i> (Nemertea: Monostilifera): <i>TJ ETQq0 0 0 rgBT /Overlock 10 Tf 50 18</i> genus. <i>Frontiers in Marine Science</i> , 0, 9, .	1.2	0
2093	Historical dispersal and host-switching formed the evolutionary history of a globally distributed multi-host parasite—The <i>Ligula intestinalis</i> species complex. <i>Molecular Phylogenetics and Evolution</i> , 2023, 180, 107677.	1.2	6
2094	To explore strange new worlds—The diversification in <i>Tremella caloplacae</i> was linked to the adaptive radiation of the Teloschistaceae. <i>Molecular Phylogenetics and Evolution</i> , 2023, 180, 107680.	1.2	2
2095	DNA Barcoding of Lepidoptera Species from the Maltese Islands: New and Additional Records, with an Insight into Endemic Diversity. <i>Diversity</i> , 2022, 14, 1090.	0.7	1

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2096	Diversity of the Piscicola Species (Hirudinea, Piscicolidae) in the Eastern Palaeartic with a Description of Three New Species and Notes on Their Biogeography. Diversity, 2023, 15, 98.	0.7	3
2097	Insights into the divergent evolution of the oceanic squid <i>Sthenoteuthis oualaniensis</i> (Cephalopoda: Ommastrephidae) from the Indian Ocean. Integrative Zoology, 2023, 18, 924-948.	1.3	4
2098	Diversification in the mountains: Evolutionary history and molecular phylogeny of Anatolian rock lizards. Molecular Phylogenetics and Evolution, 2023, 180, 107675.	1.2	1
2101	Big diversity in a small hotspot: two new species of Leptophlebiidae (Insecta, Ephemeroptera) from New Caledonia. ZooKeys, 0, 1143, 71-88.	0.5	0
2102	Molecular Weevil Identification Project: A thoroughly curated barcode release of 1300 Western Palearctic weevil species (Coleoptera, Curculionoidea). Biodiversity Data Journal, 0, 11, .	0.4	2
2103	Phylogenetic assessment of endangered and look-alike Pigtoe species in a freshwater mussel diversity hotspot. Ecology and Evolution, 2023, 13, .	0.8	3
2104	COI-Barcoding and Species Delimitation Assessment of Toad-Headed Agamas of the Genus Phrynocephalus (Agamidae, Squamata) Reveal Unrecognized Diversity in Central Eurasia. Diversity, 2023, 15, 149.	0.7	3
2105	Genetic evidence for widespread population size expansion in North American boreal birds prior to the Last Glacial Maximum. Proceedings of the Royal Society B: Biological Sciences, 2023, 290, .	1.2	6
2106	Mechanisms of reproductive isolation among cryptic species in monogonont rotifers. Hydrobiologia, 2023, 850, 4705-4718.	1.0	3
2107	Integrative Taxonomy Approach Reveals Cryptic Diversity within the Phoretic Pseudoscorpion Genus Lamprochernes (Pseudoscorpiones: Chernetidae). Insects, 2023, 14, 122.	1.0	6
2108	New species of Rhyacoglanis (Siluriformes: Pseudopimelodidae) from the upper rio Tocantins basin. Neotropical Ichthyology, 2023, 21, .	0.5	0
2109	Deep mtDNA Sequence Divergences and Possible Species Radiation of Whip Spiders (Arachnida, Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 182 Td (2023, 3, 133-147.	0.4	3
2110	Cryptic carnivores: Intercontinental sampling reveals extensive novel diversity in a genus of freshwater annelids. Molecular Phylogenetics and Evolution, 2023, 182, 107748.	1.2	0
2111	Multi-gene phylogeny reveals a new genus and species of Hapalidiales (Rhodophyta) from Antarctica: <i>Thalassolithon adeliense</i> gen. & sp. nov. Phycologia, 2023, 62, 83-98.	0.6	1
2112	Morpho-Molecular Discordance? Re-Approaching Systematics of <i>Cambeva</i> (Siluriformes: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 182 Td (2023, 3, 133-147.	0.7	2
2113	Still Many to Be Named: An Integrative Analysis of the Genus <i>Dendronotus</i> (Gastropoda: Nudibranchia) in the North Pacific Revealed Seven New Species. Diversity, 2023, 15, 162.	0.7	0
2114	Allopatric mosaics in the Indo-West Pacific crab subfamily Chlorodiellinae reveal correlated patterns of sympatry, genetic divergence, and genitalic disparity. Molecular Phylogenetics and Evolution, 2023, 181, 107710.	1.2	1
2115	The identity of <i>Argyria lacteella</i> (Fabricius, 1794) (Lepidoptera, Pyraloidea, Crambinae), synonyms, and related species revealed by morphology and DNA capture in type specimens. ZooKeys, 0, 1146, 1-42.	0.5	0

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2116	Khargia gen. nov., a new genus of simple trichal Cyanobacteria from the Persian Gulf. <i>Fottea</i> , 2023, 23, 49-61.	0.4	1
2117	A new lineage of fresh- and brackish-water mussels (Bivalvia, Mytilidae) from Southeast Asia. <i>Zoologica Scripta</i> , 2023, 52, 298-313.	0.7	0
2118	Small organelle-enriched metagenomics: An improved method for environmental DNA-based identification of marine plankton. <i>Limnology and Oceanography: Methods</i> , 2023, 21, 178-191.	1.0	1
2119	Species delimitation, molecular phylogeny and historical biogeography of the sweetlips fish (Perciformes, Haemulidae). <i>Zoosystematics and Evolution</i> , 2023, 99, 135-147.	0.4	1
2120	Diversity of the genus <i>Avrainvillea</i> (Dichotomosiphonaceae, Chlorophyta): new insights and eight new species. <i>European Journal of Phycology</i> , 0, , 1-28.	0.9	1
2121	Ophiotholia (Echinodermata: Ophiuroidea): A little-known deep-sea genus present in polymetallic nodule fields with the description of a new species. <i>Frontiers in Marine Science</i> , 0, 10, .	1.2	1
2122	DNA barcoding of the genus <i>Alburnoides</i> Jettles, 1861 (Actinopterygii, Cyprinidae) from Anatolia, Turkey. <i>Zoosystematics and Evolution</i> , 2023, 99, 185-194.	0.4	1
2123	Fly <i>DNA</i> suggests strict reliance of the causative agent of sylvatic anthrax on rainforest ecosystems. <i>Environmental DNA</i> , 2024, 6, .	3.1	2
2124	Resistance in pea (<i>Pisum sativum</i>) genetic resources to the pea aphid, <i>Acyrtosiphon pisum</i> . <i>Entomologia Experimentalis Et Applicata</i> , 2023, 171, 435-448.	0.7	1
2125	Molecular and morphological analyses disclose the existence of three species of <i>Dolichopoda</i> (Orthoptera: Rhaphidophoridae) in the Calabria region (Italy). <i>Journal of Natural History</i> , 2023, 57, 372-394.	0.2	0
2126	Similar Ones Are Not Related and Vice Versa—New Dendronotus Taxa (Nudibranchia: Dendronotidae) from the North Atlantic Ocean Provide a Platform for Discussion of Global Marine Biodiversity Patterns. <i>Diversity</i> , 2023, 15, 504.	0.7	0
2127	A new species of <i>Parauchenoglanis</i> (<i>Auchenoglanididae</i> : Siluriformes) from the Upper Lualaba River (Upper Congo) , with further evidence of hidden species diversity within the genus. <i>Journal of Fish Biology</i> , 0, , .	0.7	0
2128	A new lizard species of the <i>Liolaemus kingii</i> group (Squamata: Liolaemidae) from northwestern Chubut province (Argentina). <i>Zootaxa</i> , 2023, 5264, 235-255.	0.2	2
2129	Diversity, distribution and composition of abyssal benthic Isopoda in a region proposed for deep-seafloor mining of polymetallic nodules: a synthesis. <i>Marine Biodiversity</i> , 2023, 53, .	0.3	0
2130	A widespread Ponto-Caspian invader with a mistaken identity: integrative taxonomy elucidates the confusing taxonomy of <i>Trichogammarus trichiatus</i> (= <i>Echinogammarus</i>) (Crustacea: Tj ETQq0 0 0 rgBT, Overlock 10 Tf 50		
2131	Systematics and evolutionary history of the genus <i>Micromys</i> (Mammalia: Rodentia: Muridae). <i>Mammalian Biology</i> , 0, , .	0.8	1
2132	The phylogenetic structure and coalescent species delimitation of an endemic trapdoor spider genus, <i>Stasimopus</i> (Araneae, Mygalomorphae, Stasimopidae) in the Karoo region of South Africa. <i>Molecular Phylogenetics and Evolution</i> , 2023, 184, 107798.	1.2	2
2175	DNA Barcoding for Assessing Biodiversity. , 2023, , 21-45.		0

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