

# An Empirical Test of Bootstrapping as a Method for Ass Analysis

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Citation Report

#	ARTICLE	IF	CITATIONS
2	Thecal designs in isorophinid edrioasteroids. <i>Lethaia</i> , 1993, 26, 289-302.	0.6	38
3	Is There Something Wrong with the Bootstrap on Phylogenies? A Reply to Hillis and Bull. <i>Systematic Biology</i> , 1993, 42, 193.	2.7	25
4	An Empirical Test of Bootstrapping as a Method for Assessing Confidence in Phylogenetic Analysis. <i>Systematic Biology</i> , 1993, 42, 182.	2.7	2,316
5	On Islands of Trees and the Efficacy of Different Methods of Branch Swapping in Finding Most-Parsimonious Trees. <i>Systematic Biology</i> , 1993, 42, 200.	2.7	0
6	Experimental Molecular Evolution of Bacteriophage T7. <i>Evolution; International Journal of Organic Evolution</i> , 1993, 47, 993.	1.1	17
7	On Islands of Trees and the Efficacy of Different Methods of Branch Swapping in Finding Most-Parsimonious Trees. <i>Systematic Biology</i> , 1993, 42, 200-210.	2.7	39
8	[34] Analysis of DNA sequence data: Phylogenetic inference. <i>Methods in Enzymology</i> , 1993, 224, 456-487.	0.4	113
9	EXPERIMENTAL MOLECULAR EVOLUTION OF BACTERIOPHAGE T7. <i>Evolution; International Journal of Organic Evolution</i> , 1993, 47, 993-1007.	1.1	61
10	Animals and fungi are each other's closest relatives: congruent evidence from multiple proteins.. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1993, 90, 11558-11562.	3.3	526
11	Is there Something Wrong with the Bootstrap on Phylogenies? A Reply to Hillis and Bull. <i>Systematic Biology</i> , 1993, 42, 193-200.	2.7	270
12	Chloroplast DNA systematics: a review of methods and data analysis. <i>American Journal of Botany</i> , 1994, 81, 1205-1224.	0.8	480
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16	Intraspecific Cladogram Estimation: Accuracy at Higher Levels of Divergence. <i>Systematic Biology</i> , 1994, 43, 222-235.	2.7	118
17	What is the Bootstrap Technique?. <i>Systematic Biology</i> , 1994, 43, 424-430.	2.7	26
18	Evolution of a Plant Homeotic Multigene Family: Toward Connecting Molecular Systematics Andmolecular Developmental Genetics. <i>Systematic Biology</i> , 1994, 43, 307-328.	2.7	62
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21	Phylogeny of <i>Drosophila</i> and related genera inferred from the nucleotide sequence of the Cu,Zn Sod gene. <i>Journal of Molecular Evolution</i> , 1994, 38, 443-454.	0.8	93
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32	Evolutionary History of the Symbiosis Between Fungus-Growing Ants and Their Fungi. <i>Science</i> , 1994, 266, 1691-1694.	6.0	376
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34	Mitochondrial DNA Sequence Divergence and Phylogenetic Relationships among Eight Chromosome Races of the <i>Sceloporus Grammicus</i> Complex (Phrynosomatidae) in Central Mexico. <i>Systematic Biology</i> , 1994, 43, 387-418.	2.7	687
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37	Phylogeny and Systematics of <i>Sphenostylis</i> and <i>Nesphostylis</i> (Leguminosae: Phaseoleae) Based on Morphological and Chloroplast DNA Data. <i>Systematic Botany</i> , 1994, 19, 389.	0.2	14

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53	Objections to Bootstrapping Phylogenies: A Critique. <i>Systematic Biology</i> , 1995, 44, 299.	2.7	26
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94	A Combined Morphological and Molecular Approach to the Phylogeny of Asteroids (Asteroidea): Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 7	2.7	54
95	Molecular phylogeny of the grass genus <i>Brachypodium</i> P. Beauv. based on RFLP and RAPD analysis. <i>Botanical Journal of the Linnean Society</i> , 1995, 117, 263-280.	0.8	45
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1411	Phylogenetic Inference Using Molecular Data. <i>Crustacean Issues</i> , 2009, , 67-88.	0.9	3
1412	<i>Cymadothea trifolii</i> , an obligate biotrophic leaf parasite of <i>Trifolium</i> , belongs to <i>Mycosphaerellaceae</i> as shown by nuclear ribosomal DNA analyses. <i>Persoonia: Molecular Phylogeny and Evolution of Fungi</i> , 2009, 22, 49-55.	1.6	34
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1414	Phylogenetic Relationships Between and within <i>Phacelia</i> Sections <i>Whitlavia</i> and <i>Gymnobythus</i> (Boraginaceae). <i>Systematic Botany</i> , 2009, 34, 737-746.	0.2	9
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1416	Polyphasic taxonomy of <i>Aspergillus</i> section <i>Sparsi</i> . <i>IMA Fungus</i> , 2010, 1, 187-195.	1.7	12
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1428	Bacterial community diversity in paper mills processing recycled paper. <i>Journal of Industrial Microbiology and Biotechnology</i> , 2010, 37, 1061-1069.	1.4	16
1429	Bacterial communities associated with the rhizosphere of pioneer plants ( <i>Bahia xylopoda</i> and <i>Viguiera</i> ) <a href="#">Tj ETQq1 1 0.784314 rgBT /Over</a>	0.7	7
1430	A molecular phylogenetic study of <i>Hemsleya</i> (Cucurbitaceae) based on ITS, rpl16, trnH-psbA, and trnL DNA sequences. <i>Plant Systematics and Evolution</i> , 2010, 285, 23-32.	0.3	9
1431	Phylogeny of <i>Salsola</i> s.l. (Chenopodiaceae) based on DNA sequence data from ITS, psbA-psbH, and rbcL, with emphasis on taxa of northwestern China. <i>Plant Systematics and Evolution</i> , 2010, 288, 25-42.	0.3	41
1432	Molecular phylogeny of <i>Phalaenopsis</i> Blume (Orchidaceae) on the basis of plastid and nuclear DNA. <i>Plant Systematics and Evolution</i> , 2010, 288, 77-98.	0.3	33
1433	Narrow species concepts in the <i>Frullania dilatata</i> "appalachiana" "eboracensis" complex (Porellales). <a href="#">Tj ETQq1 1 0.784314 rgBT /Over</a> <i>Evolution</i> , 2010, 290, 151-158.	0.3	14
1434	Phylogenetics and Biogeography of the <i>Phalaenopsis violacea</i> (Orchidaceae) Species Complex Based on Nuclear and Plastid DNA. <i>Journal of Plant Biology</i> , 2010, 53, 453-460.	0.9	7
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1441	Applying phylogenetic analysis to viral livestock diseases: Moving beyond molecular typing. <i>Veterinary Journal</i> , 2010, 184, 130-137.	0.6	13
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1446	Epidemiology of vampire bat-transmitted rabies virus in Goiás, central Brazil: re-evaluation based on G-L intergenic region. <i>BMC Research Notes</i> , 2010, 3, 288.	0.6	4
1447	Fumonisin contamination and fumonisin producing black <i>Aspergilli</i> in dried vine fruits of different origin. <i>International Journal of Food Microbiology</i> , 2010, 143, 143-149.	2.1	82
1448	Characterization of a diversity of tetraphyllidean and rhinebothriidean cestode larval types, with comments on host associations and life-cycles. <i>International Journal for Parasitology</i> , 2010, 40, 889-910.	1.3	106
1449	Molecular diversity of peanut- $\epsilon$ nodulating rhizobia in soils of Argentina. <i>Journal of Basic Microbiology</i> , 2010, 50, 274-279.	1.8	13
1450	Species authentication of octopus, cuttlefish, bobtail and bottle squids (families Octopodidae,) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 18	4.2	33
1451	Molecular identification and pathology of <i>Anisakis pegreffii</i> (Nematoda: Anisakidae) infection in the Mediterranean loggerhead sea turtle ( <i>Caretta caretta</i> ). <i>Veterinary Parasitology</i> , 2010, 174, 65-71.	0.7	20
1452	Phylogenetic relationships of <i>Ansonia</i> from Southeast Asia inferred from mitochondrial DNA sequences: Systematic and biogeographic implications (Anura: Bufonidae). <i>Molecular Phylogenetics and Evolution</i> , 2010, 54, 561-570.	1.2	49
1453	The evolutionary diversification of parrots supports a taxon pulse model with multiple trans-oceanic dispersal events and local radiations. <i>Molecular Phylogenetics and Evolution</i> , 2010, 54, 984-994.	1.2	66

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1455	Complex patterns of continental speciation: Molecular phylogenetics and biogeography of sub-Saharan puddle frogs ( <i>Phrynobatrachus</i> ). <i>Molecular Phylogenetics and Evolution</i> , 2010, 55, 883-900.	1.2	49
1456	Effect of geological vicariance on mitochondrial DNA differentiation in Common Pheasant populations of the Loess Plateau and eastern China. <i>Molecular Phylogenetics and Evolution</i> , 2010, 55, 409-417.	1.2	25
1457	An expanded phylogeny of treefrogs (Hylidae) based on nuclear and mitochondrial sequence data. <i>Molecular Phylogenetics and Evolution</i> , 2010, 55, 871-882.	1.2	138
1458	Large multi-locus plastid phylogeny of the tribe Arundinarieae (Poaceae: Bambusoideae) reveals ten major lineages and low rate of molecular divergence. <i>Molecular Phylogenetics and Evolution</i> , 2010, 56, 821-839.	1.2	83
1459	One species or at least eight? Delimitation and distribution of <i>Frullania tamarisci</i> (L.) Dumort. s. l. (Jungermanniopsida, Porellales) inferred from nuclear and chloroplast DNA markers. <i>Molecular Phylogenetics and Evolution</i> , 2010, 56, 1105-1114.	1.2	99
1460	Polyphyly of the genus <i>Axinella</i> and of the family Axinellidae (Porifera: Demospongiaep). <i>Molecular Phylogenetics and Evolution</i> , 2010, 57, 35-47.	1.2	51
1461	Phylogeography and historical demography of <i>Polypedates leucomystax</i> in the islands of Indonesia and the Philippines: Evidence for recent human-mediated range expansion?. <i>Molecular Phylogenetics and Evolution</i> , 2010, 57, 598-619.	1.2	64
1462	Late Miocene diversification and phylogenetic relationships of the huge toads in the <i>Rhinella marina</i> (Linnaeus, 1758) species group (Anura: Bufonidae). <i>Molecular Phylogenetics and Evolution</i> , 2010, 57, 787-797.	1.2	66
1463	Phylogeny of the leafy liverwort <i>Ptilidium</i> : Cryptic speciation and shared haplotypes between the Northern and Southern Hemispheres. <i>Molecular Phylogenetics and Evolution</i> , 2010, 57, 1260-1267.	1.2	55
1464	DNA-based characterisation and classification of forensically important flesh flies (Diptera: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 342 Td	1.3	53
1465	Including RNA secondary structures improves accuracy and robustness in reconstruction of phylogenetic trees. <i>Biology Direct</i> , 2010, 5, 4.	1.9	154
1466	Biogeographic regions of the Iberian Peninsula: butterflies as biogeographical indicators. <i>Journal of Zoology</i> , 2010, 282, 180-190.	0.8	15
1467	A neglected lineage of North American turtles fills a major gap in the fossil record. <i>Palaeontology</i> , 2010, 53, 241-248.	1.0	41
1468	Parallel habitat specialization within the wolf spider genus <i>Hogna</i> from the Galápagos. <i>Molecular Ecology</i> , 2010, 19, 4029-4045.	2.0	28
1469	A phylogeny of Adelanthaceae (Jungermanniales, Marchantiophyta) based on nuclear and chloroplast DNA markers, with comments on classification, cryptic speciation and biogeography. <i>Molecular Phylogenetics and Evolution</i> , 2010, 55, 293-304.	1.2	58
1470	Diversification in species complexes: Tests of species origin and delimitation in the <i>Bursera simaruba</i> clade of tropical trees (Burseraceae). <i>Molecular Phylogenetics and Evolution</i> , 2010, 57, 798-811.	1.2	33
1471	Discordant mitochondrial and nuclear gene phylogenies in emydid turtles: implications for speciation and conservation. <i>Biological Journal of the Linnean Society</i> , 0, 99, 445-461.	0.7	117

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1473	Distinctions between optimal and expected support. <i>Cladistics</i> , 2010, 26, 657-663.	1.5	7
1474	Recovering phylogenetic signal from frog mating calls. <i>Zoologica Scripta</i> , 2010, 39, 141-154.	0.7	56
1475	Systematics of Andean gladiator frogs of the <i>Hypsiboas pulchellus</i> species group (Anura, Hylidae). <i>Zoologica Scripta</i> , 2010, 39, 572-590.	0.7	23
1476	Major clades and phylogenetic relationships between lichenized and non-lichenized lineages in <i>Ostropales</i> (Ascomycota: Lecanoromycetes). <i>Taxon</i> , 2010, 59, 1483-1494.	0.4	74
1477	A new species of the genus <i>Tropidophorus</i> Duméril & Bibron, 1839 (Squamata: Sauria): <i>Tropidophorus</i> sp. nov. <i>Zootaxa</i> , 2010, 2391, 33.	0.2	8
1478	Taxonomic re-assessment of the Australian and New Guinean green-eyed treefrogs <i>Litoria eucnemis</i> , <i>L. genimaculata</i> and <i>L. serrata</i> (Anura: Hylidae). <i>Zootaxa</i> , 2010, 2391, 33.	0.2	8
1479	Systematic reassessments of fanged frogs from China and adjacent regions (Anura: Dicoglossidae). <i>Zootaxa</i> , 2010, 2345, 33.	0.2	18
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1482	Genetic relationships among <i>Heliconia</i> (Heliconiaceae) species based on RAPD markers. <i>Genetics and Molecular Research</i> , 2010, 9, 1377-1387.	0.3	18
1483	Phylogenetic incongruence arising from fragmented speciation in enteric bacteria. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 11453-11458.	3.3	62
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1485	cTBP: A Successful Intron Length Polymorphism (ILP)-Based Genotyping Method Targeted to Well Defined Experimental Needs. <i>Diversity</i> , 2010, 2, 572-585.	0.7	38
1486	Phylogeny and morphology of four new species of <i>Lasiodiplodia</i> from Iran. <i>Persoonia: Molecular Phylogeny and Evolution of Fungi</i> , 2010, 25, 1-10.	1.6	135
1487	<i>Neoleurodiscus fujii</i> , a new genus and new species found at the timberline in Japan. <i>Mycologia</i> , 2010, 102, 217-223.	0.8	12
1488	An Ancient Origin for the Enigmatic Flat-Headed Frogs (Bombinatoridae: <i>Barbourula</i> ) from the Islands of Southeast Asia. <i>PLoS ONE</i> , 2010, 5, e12090.	1.1	71
1489	Short Communication: Biological and Genetic Characterization of HIV Type 1 Subtype B and Nonsubtype B Transmitted Viruses: Usefulness for Vaccine Candidate Assessment. <i>AIDS Research and Human Retroviruses</i> , 2010, 26, 1019-1025.	0.5	23

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1491	Light at the end of the tunnel: insights into the molecular systematics of East African puddle frogs (Anura: Phrynobatrachidae). <i>Systematics and Biodiversity</i> , 2010, 8, 39-47.	0.5	16
1492	DNA barcode data confirm new species and reveal cryptic diversity in Chilean <i>Smicridea</i> ( <i>Smicridea</i> ) (Trichoptera:Hydropsychidae). <i>Journal of the North American Benthological Society</i> , 2010, 29, 1058-1074.	3.0	72
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1497	Two Independent Epidemics of HIV in Maryland. <i>Journal of Acquired Immune Deficiency Syndromes</i> (1999), 2010, 54, 297-303.	0.9	19
1498	<i>Remototrachyna</i> , a newly recognized tropical lineage of lichens in the Hypotrachyna clade (Parmeliaceae, Ascomycota), originated in the Indian subcontinent. <i>American Journal of Botany</i> , 2010, 97, 579-590.	0.8	61
1499	Biogeographic Patterns of Diversification and the Origins of <i>Cleome</i> in the Neotropics (Cleomaceae). <i>Systematic Botany</i> , 2010, 35, 811-826.	0.2	77
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1501	Multigene phylogeny and mating tests reveal three cryptic species related to <i>Calonectria pauciramosa</i> . <i>Studies in Mycology</i> , 2010, 66, 15-30.	4.5	63
1502	Phylogeny and systematics of the genus <i>Calonectria</i> . <i>Studies in Mycology</i> , 2010, 66, 31-69.	4.5	119
1503	Rapid Identification of Seaweeds in Food Products by PCR Combined with ALF-RFLP and FINS Methodologies. <i>Journal of Agricultural and Food Chemistry</i> , 2010, 58, 11586-11592.	2.4	8
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1509	Phylogenetic Relationships in <i>Eleocharis</i> (Cyperaceae): C <sub>4</sub> Photosynthesis Origins and Patterns of Diversification in the Spikerushes. <i>Systematic Botany</i> , 2010, 35, 257-271.	0.2	39
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1513	Correlation, hierarchies, and networks in financial markets. <i>Journal of Economic Behavior and Organization</i> , 2010, 75, 40-58.	1.0	287
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1517	Isolation, biochemical characterization, and molecular modeling of American lobster digestive cathepsin D1. <i>Comparative Biochemistry and Physiology - B Biochemistry and Molecular Biology</i> , 2010, 157, 394-400.	0.7	16
1518	Molecular epidemiology of rabies virus isolates in Uganda. <i>Virus Research</i> , 2010, 147, 135-138.	1.1	7
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1520	A New Squeaker Frog (Arthroleptidae: Arthroleptis) from the Mountains of Cameroon and Nigeria. <i>Herpetologica</i> , 2010, 66, 335-348.	0.2	18
1521	Phylogenetic status of four new species of <i>Acanthobothrium</i> (Cestoda:Tetracanthocephala) parasitic on the wedgefish <i>Rhynchobatus laevis</i> (Elasmobranchii:Rhynchobatidae): implications for interpreting host associations. <i>Invertebrate Systematics</i> , 2010, 24, 419.	0.5	25
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1524	The phylogenetic utility of the nuclear protein-coding gene EF-1 $\alpha$ for resolving recent divergences in Opiliones, emphasizing intron evolution. <i>Journal of Arachnology</i> , 2010, 38, 9-20.	0.3	20
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1527	Inference of phylogenetic relationships among the subfamilies of grasses (Poaceae: Poales) using meso-scale exemplar-based sampling of the plastid genome. Botany, 2010, 88, 65-84.	0.5	27
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1534	Studies on microbiologically influenced corrosion of SS304 by a novel manganese oxidizer, <i>Bacillus flexus</i> . Biofouling, 2011, 27, 675-683.	0.8	19
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1542	Insight into the Short-Finned Squid <i>Illex coindetii</i> (Cephalopoda: Ommastrephidae) Feeding Ecology: Is There a Link Between Helminth Parasites and Food Composition?. Journal of Parasitology, 2011, 97, 55-62.	0.3	28
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1546	<i>Cladophialophora psammophila</i> , a novel species of Chaetothyriales with a potential use in the bioremediation of volatile aromatic hydrocarbons. <i>Fungal Biology</i> , 2011, 115, 1019-1029.	1.1	73
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1557	A New Puddle Frog (Phrynobatrachidae: Phrynobatrachus) from the Obudu Plateau In Eastern Nigeria. <i>Herpetologica</i> , 2011, 67, 271-287.	0.2	11
1558	Species identification of Tanzanian antelopes using DNA barcoding. <i>Molecular Ecology Resources</i> , 2011, 11, 442-449.	2.2	30
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1560	An Extremely Hyperapolytic <i>Acanthobothrium</i> Species (Cestoda: Tetraphyllidea) from the Japanese Wobbegong, <i>Orectolobus japonicus</i> (Elasmobranchii: Orectolobiformes) in Taiwan. <i>Comparative Parasitology</i> , 2011, 78, 4-14.	0.0	23
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1563	Phylogeny, morphology, and biogeography of <i>Haplophyllum</i> (Rutaceae), a species-rich genus of the Irano-Turanian floristic region. <i>Taxon</i> , 2011, 60, 513-527.	0.4	24

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1565	A new species of Cycloramphus Tschudi (Anura: Cycloramphidae) from the Parque Nacional da Serra dos Açúços, Southeastern Brazil. Zootaxa, 2011, 2737, 19.	0.2	12
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1581	Phylogeography of Middle American gophersnakes: mixed responses to biogeographical barriers across the Mexican Transition Zone. Journal of Biogeography, 2011, 38, 1570-1584.	1.4	101



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1583	Macroevolutionary patterns in the diversification of parrots: effects of climate change, geological events and key innovations. <i>Journal of Biogeography</i> , 2011, 38, 2176-2194.	1.4	60
1584	Biogeography and morphological evolution in a Pacific island ant radiation. <i>Molecular Ecology</i> , 2011, 20, 114-130.	2.0	34
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1586	A new genus and species of nothrotheriid sloth ( <i>Xenarthra</i> , Tardigrada, Nothrotheriidae) from the Late Miocene (Huayquerian) of Peru. <i>Palaeontology</i> , 2011, 54, 171-205.	1.0	51
1587	MOLECULAR PHYLOGENETIC RELATIONSHIPS IN THE FRESHWATER FAMILY HYDRODICTYACEAE (SPHAEROPLEALES, CHLOROPHYCEAE), WITH AN EMPHASIS ON <i>PEDIASTRUM DUPLEX</i> . <i>Journal of Phycology</i> , 2011, 47, 152-163.	1.0	66
1588	Molecular phylogeny and diversity of the Corsican red-legged partridge: hybridization and management issues. <i>Journal of Zoology</i> , 2011, 285, no-no.	0.8	15
1589	Intraclass Evolution and Classification of the Colpodea (Ciliophora). <i>Journal of Eukaryotic Microbiology</i> , 2011, 58, 397-415.	0.8	32
1590	Speciation at the Mogollon Rim in the Arizona Mountain Kingsnake ( <i>Lampropeltis pyromelana</i> ). <i>Molecular Phylogenetics and Evolution</i> , 2011, 60, 445-454.	1.2	51
1591	Congeneric phylogeographical sampling reveals polyphyly and novel biodiversity within black basses (Centrarchidae: <i>Micropterus</i> ). <i>Biological Journal of the Linnean Society</i> , 2011, 104, 346-363.	0.7	22
1592	A phylogenetic analysis of morphological and molecular characters of <i>Lithospermum</i> L. (Boraginaceae) and related taxa: evolutionary relationships and character evolution. <i>Cladistics</i> , 2011, 27, 559-580.	1.5	31
1593	A molecular phylogeny of Equatorial African Lacertidae, with the description of a new genus and species from eastern Democratic Republic of the Congo. <i>Zoological Journal of the Linnean Society</i> , 2011, 163, 913-942.	1.0	25
1594	Systematics and spicule evolution in dictyonal sponges (Hexactinellida: Sceptrulophora) with description of two new species. <i>Zoological Journal of the Linnean Society</i> , 2011, 163, 1003-1025.	1.0	22
1595	Typing of nitrogen-fixing <i>Frankia</i> strains by matrix-assisted laser desorption ionization-time-of-flight (MALDI-TOF) mass spectrometry. <i>Systematic and Applied Microbiology</i> , 2011, 34, 63-68.	1.2	26
1596	A methodological investigation of hominoid craniodental morphology and phylogenetics. <i>Journal of Human Evolution</i> , 2011, 60, 47-57.	1.3	19
1597	Multi-locus sequence analysis reveals host specific association between <i>Bartonella washoensis</i> and squirrels. <i>Veterinary Microbiology</i> , 2011, 148, 60-65.	0.8	22
1598	Evolutionary dynamics of rabies viruses highlights the importance of China rabies transmission in Asia. <i>Virology</i> , 2011, 410, 403-409.	1.1	31
1599	A new xenoma-forming microsporidium infecting intestinal tract of Atlantic bluefin tuna ( <i>Thunnus</i> ). <i>Journal of Eukaryotic Microbiology</i> , 2011, 58, 397-415.	0.4	3

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1601	New insights on systematics and phylogenetics of Mediterranean <i>Blaps</i> species (Coleoptera: Tenebrionidae: Blaptini), assessed through morphology and dense taxon sampling. <i>Systematic Entomology</i> , 2011, 36, 340-361.	1.7	22
1602	Molecular and chemical characters to evaluate species status of two cuckoo bumblebees: <i>Bombus barbutellus</i> and <i>Bombus maxillosus</i> (Hymenoptera, Apidae, Bombini). <i>Systematic Entomology</i> , 2011, 36, 453-469.	1.7	34
1603	Molecular characterization of <i>Fasciola</i> spp. from the endemic area of northern Iran based on nuclear ribosomal DNA sequences. <i>Experimental Parasitology</i> , 2011, 128, 196-204.	0.5	49
1604	Molecular characterization of <i>Fasciola gigantica</i> from Mauritania based on mitochondrial and nuclear ribosomal DNA sequences. <i>Experimental Parasitology</i> , 2011, 129, 127-136.	0.5	25
1605	Genetic divergence of <i>Octopus vulgaris</i> species in the eastern Mediterranean. <i>Biochemical Systematics and Ecology</i> , 2011, 39, 277-282.	0.6	14
1606	Freshwater ascomycetes: two new species of <i>Lindgomyces</i> (Lindgomycetaceae, Pleosporales). <i>Journal of Eukaryotic Microbiology</i> , 2011, 57, 50-52.	0.8	23
1607	Two new aflatoxin producing species, and an overview of <i>Aspergillus</i> section <i>Flavi</i> . <i>Studies in Mycology</i> , 2011, 69, 57-80.	4.5	274
1608	<i>Parmelia sulcata</i> (Ascomycota: Parmeliaceae), a sympatric monophyletic species complex. <i>Lichenologist</i> , 2011, 43, 585-601.	0.5	49
1609	Conundrums in species concepts: the discovery of a new cryptic species segregated from <i>Parmelina tiliacea</i> (Ascomycota: Parmeliaceae). <i>Lichenologist</i> , 2011, 43, 603-616.	0.5	33
1610	Molecular phylogeny of the <i>Sinocyclocheilus</i> (Cypriniformes: Cyprinidae) fishes in northwest part of Guangxi, China. <i>Environmental Biology of Fishes</i> , 2011, 92, 371-379.	0.4	6
1611	Characterization of Expressed Sequence Tags from Flower Buds of Alpine <i>Lilium formosanum</i> using a Subtractive cDNA Library. <i>Plant Molecular Biology Reporter</i> , 2011, 29, 88-97.	1.0	16
1612	h-TBP: an approach based on intron-length polymorphism for the rapid isolation and characterization of the multiple members of the $\beta$ -tubulin gene family in <i>Camelina sativa</i> (L.) Crantz. <i>Molecular Breeding</i> , 2011, 28, 635-645.	1.0	25
1613	A large Finnish echovirus 30 outbreak was preceded by silent circulation of the same genotype. <i>Virus Genes</i> , 2011, 42, 28-36.	0.7	41
1614	Molecular characterization of <i>Hysterothylacium aduncum</i> (Nematoda: Raphidascaridae) from different fish caught off the Tunisian coast based on nuclear ribosomal DNA sequences. <i>Parasitology Research</i> , 2011, 109, 1429-1437.	0.6	20
1615	Nucleotide diversity of a genomic sequence similar to SHATTERPROOF (PvSHP1) in domesticated and wild common bean ( <i>Phaseolus vulgaris</i> L.). <i>Theoretical and Applied Genetics</i> , 2011, 123, 1341-1357.	1.8	44
1616	FINS methodology to identification of sardines and related species in canned products and detection of mixture by means of SNP analysis systems. <i>European Food Research and Technology</i> , 2011, 232, 1077-1086.	1.6	17
1617	Authentication of swordfish ( <i>Xiphias gladius</i> ) by RT-PCR and FINS methodologies. <i>European Food Research and Technology</i> , 2011, 233, 195-202.	1.6	20

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1618	Identification of <i>Tropilaelaps</i> mites (Acari, Laelapidae) infesting <i>Apis mellifera</i> in China. <i>Apidologie</i> , 2011, 42, 485-498.	0.9	10
1619	Isolation and Partial Characterization of Halotolerant Lactic Acid Bacteria from Two Mexican Cheeses. <i>Applied Biochemistry and Biotechnology</i> , 2011, 164, 889-905.	1.4	65
1620	Bioaccumulation of cerium and neodymium by <i>Bacillus cereus</i> isolated from rare earth environments of Chavara and Manavalakurichi, India. <i>Indian Journal of Microbiology</i> , 2011, 51, 488-495.	1.5	30
1621	A new kinosternoid from the Late Cretaceous Hell Creek Formation of North Dakota and Montana and the origin of the <i>Dermatemys mawii</i> lineage. <i>Palaontologische Zeitschrift</i> , 2011, 85, 125-142.	0.8	26
1622	Another example of cryptic diversity in lichen-forming fungi: the new species <i>Parmelia mayi</i> (Ascomycota: Parmeliaceae). <i>Organisms Diversity and Evolution</i> , 2011, 11, 331-342.	0.7	65
1623	<i>Scleroramularia</i> gen. nov. associated with sooty blotch and flyspeck of apple and pawpaw from the Northern Hemisphere. <i>Fungal Diversity</i> , 2011, 46, 53-66.	4.7	26
1624	<i>Astrosphaeriella</i> is polyphyletic, with species in <i>Fissuroma</i> gen. nov., and <i>Neoastrosphaeriella</i> gen. nov.. <i>Fungal Diversity</i> , 2011, 51, 135-154.	4.7	81
1625	First molecular identification of the zoonotic parasite <i>Anisakis pegreffii</i> (Nematoda: Anisakidae) in a paraffin-embedded granuloma taken from a case of human intestinal anisakiasis in Italy. <i>BMC Infectious Diseases</i> , 2011, 11, 82.	1.3	130
1626	Rebooting the human mitochondrial phylogeny: an automated and scalable methodology with expert knowledge. <i>BMC Bioinformatics</i> , 2011, 12, 174.	1.2	15
1627	Plio-Pleistocene sea level and temperature fluctuations in the northwestern Pacific promoted speciation in the globally-distributed flathead mullet <i>Mugil cephalus</i> . <i>BMC Evolutionary Biology</i> , 2011, 11, 83.	3.2	146
1628	Lizards from the end of the world: Phylogenetic relationships of the <i>Liolaemus lineomaculatus</i> section (Squamata: Iguania: Liolaemini). <i>Molecular Phylogenetics and Evolution</i> , 2011, 59, 364-376.	1.2	75
1629	Local-scale environmental variation generates highly divergent lineages associated with stream drainages in a terrestrial salamander, <i>Plethodon caddoensis</i> . <i>Molecular Phylogenetics and Evolution</i> , 2011, 59, 399-411.	1.2	28
1630	Phylogeny and character evolution in the jelly fungi (Tremellomycetes, Basidiomycota, Fungi). <i>Molecular Phylogenetics and Evolution</i> , 2011, 61, 12-28.	1.2	114
1631	Spurious 99% bootstrap and jackknife support for unsupported clades. <i>Molecular Phylogenetics and Evolution</i> , 2011, 61, 177-191.	1.2	76
1632	The Interconversion of UDP-Arabinopyranose and UDP-Arabinofuranose Is Indispensable for Plant Development in <i>Arabidopsis</i> . <i>Plant Cell</i> , 2011, 23, 1373-1390.	3.1	134
1633	Molecular Epidemiology of Rift Valley Fever Virus. <i>Emerging Infectious Diseases</i> , 2011, 17, 2270-2276.	2.0	128
1634	Are Melanistic Populations of the Karoo Girdled Lizard, <i>Karusasaurus polyzonus</i> , Relics or Ecotypes? A Molecular Investigation. <i>African Zoology</i> , 2011, 46, 146-155.	0.2	4
1635	<i>Alternaria hungarica</i> sp. nov., a minor foliar pathogen of wheat in Hungary. <i>Mycologia</i> , 2011, 103, 94-100.	0.8	15

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1637	Genotypic and Phenotypic Variation in Pseudomonas aeruginosa Reveals Signatures of Secondary Infection and Mutator Activity in Certain Cystic Fibrosis Patients with Chronic Lung Infections. Infection and Immunity, 2011, 79, 4802-4818.	1.0	31
1638	Mitochondrial DNA sequence variation and phylogeography of the scarlet kingsnake (Lampropeltis tigris) in the Florida Keys. Systematic Biology, 2011, 60, 1075-1085.	0.6	10
1639	Vahliellaceae, a new family of cyanobacterial lichens (Peltigerales, Ascomycetes). Lichenologist, 2011, 43, 67-72.	0.5	22
1641	Independent Origin of Radial Floral Symmetry in the Gloxinieae (Gesnerioideae: Gesneriaceae) is Supported by the Rediscovery of <i>Phinaea pulchella</i> in Cuba. Systematic Botany, 2011, 36, 757-767.	0.2	7
1642	Survey of Branch Support Methods Demonstrates Accuracy, Power, and Robustness of Fast Likelihood-based Approximation Schemes. Systematic Biology, 2011, 60, 685-699.	2.7	912
1643	Formalizing morphologically cryptic biological entities: New insights from DNA taxonomy, hybridization, and biogeography in the leafy liverwort <i>Porella platyphylla</i> (Jungermanniopsida). Systematic Botany, 2011, 36, 757-767.	0.8	10
1644	A new species of Andean toad (Bufonidae, Osornophryne) discovered using molecular and morphological data, with a taxonomic key for the genus. ZooKeys, 2011, 108, 73-97.	0.5	14
1645	Amplified fragment length polymorphism markers for DNA fingerprinting in the genus Salvia. Plant Biosystems, 2011, 145, 274-277.	0.8	13
1646	The Devil in the Details: Interactions between the Branch-Length Prior and Likelihood Model Affect Node Support and Branch Lengths in the Phylogeny of the Psoraceae. Systematic Biology, 2011, 60, 541-561.	2.7	44
1647	Performance, Accuracy, and Web Server for Evolutionary Placement of Short Sequence Reads under Maximum Likelihood. Systematic Biology, 2011, 60, 291-302.	2.7	476
1648	Additions to the Mycosphaerella complex. IMA Fungus, 2011, 2, 49-64.	1.7	35
1649	Detection and Classification of SPLCV Isolates in the U.S. Sweetpotato Germplasm Collection via a Real-Time PCR Assay and Phylogenetic Analysis. Plant Disease, 2011, 95, 1385-1391.	0.7	5
1650	Testing the phylogenetic utility of MCM7 in the Ascomycota. MycoKeys, 2011, 1, 63-94.	0.8	58
1651	Morphological, molecular characterisation and phylogenetic position of Longidorus mindanaoensis n. sp. (Nematoda: Longidoridae) from a Philippine Avicennia mangrove habitat. Nematology, 2012, 14, 285-307.	0.2	11
1652	A new species of Orobdella (Hirudinida, Arhynchobdellida, Gastrostomobdellidae) and redescription of O. kawakatsuorum from Hokkaido, Japan with the phylogenetic position of the new species. ZooKeys, 2012, 169, 9-30.	0.5	38
1653	DACTAL: divide-and-conquer trees (almost) without alignments. Bioinformatics, 2012, 28, i274-i282.	1.8	41
1654	A new sexannulate species of Orobdella (Hirudinida, Arhynchobdellida, Orobdellidae) from Yakushima Island, Japan. ZooKeys, 2012, 181, 79-93.	0.5	24

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1655	High Diversity of Rabies Viruses Associated with Insectivorous Bats in Argentina: Presence of Several Independent Enzootics. <i>PLoS Neglected Tropical Diseases</i> , 2012, 6, e1635.	1.3	22
1656	A new species of <i>Orobdella</i> (Hirudinida, Arhynchobdellida, Orobdellidae) from Taipei, Taiwan. <i>ZooKeys</i> , 2012, 207, 49-63.	0.5	15
1657	First report of the exotic blue land planarian, <i>Caenoplana coerulea</i> (Platyhelminthes, Geoplanidae), on Menorca (Balearic Islands, Spain). <i>ZooKeys</i> , 2012, 199, 91-105.	0.5	14
1658	Efficiency of Nuclear and Mitochondrial Markers Recovering and Supporting Known Amniote Groups. <i>Evolutionary Bioinformatics</i> , 2012, 8, EBO.S9656.	0.6	5
1659	Taxon Influence Index: Assessing Taxon-Induced Incongruities in Phylogenetic Inference. <i>Systematic Biology</i> , 2012, 61, 337-345.	2.7	10
1660	The multiple fuzzy origins of woodiness within Balsaminaceae using an integrated approach. Where do we draw the line?. <i>Annals of Botany</i> , 2012, 109, 783-799.	1.4	34
1661	Resolving the <i>Diplodia</i> complex on apple and other <i>Rosaceae</i> hosts. <i>Persoonia: Molecular Phylogeny and Evolution of Fungi</i> , 2012, 29, 29-38.	1.6	70
1662	Molecular Characteristics of Subterranean Termites of the genus <i>Reticulitermes</i> (Isoptera: Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50	1.3	10
1663	A New Emerald Ash Borer (Coleoptera: Buprestidae) Parasitoid Species of <i>Spathius</i> Nees (Hymenoptera: Braconidae: Doryctinae) from the Russian Far East and South Korea. <i>Annals of the Entomological Society of America</i> , 2012, 105, 165-178.	1.3	52
1664	<i>Parmelina yalungana</i> resurrected and reported from Alaska, China and Russia. <i>Bryologist</i> , 2012, 115, 557-565.	0.1	5
1665	Genetic analysis of a novel nidovirus from fathead minnows. <i>Journal of General Virology</i> , 2012, 93, 1247-1252.	1.3	32
1666	<i>Arabidopsis&lt;i&gt;Deficient in Cutin Ferulate&lt;/i&gt;Encodes a Transferase Required for Feruloylation of 1%-Hydroxy Fatty Acids in Cutin Polyester</i> <i>Plant Physiology</i> , 2012, 158, 654-665.	2.3	86
1667	Genetic structuring of European anchovy (<i>Engraulis encrasicolus</i>) populations through mitochondrial DNA sequences. <i>Mitochondrial DNA</i> , 2012, 23, 62-69.	0.6	9
1668	<i>Frullania knightbridgei</i> , a new liverwort (Frullaniaceae, Marchantiophyta) species from the deep south of Aotearoa-New Zealand based on an integrated evidence-based approach. <i>PhytoKeys</i> , 2012, 8, 13.	0.4	19
1669	Ochratoxigenic Black Species of <i>Aspergilli</i> in Grape Fruits of Northern Italy Identified by an Improved PCR-RFLP Procedure. <i>Toxins</i> , 2012, 4, 42-54.	1.5	23
1670	Fungal trunk pathogens associated with wood decay of almond trees on Mallorca (Spain). <i>Persoonia: Molecular Phylogeny and Evolution of Fungi</i> , 2012, 28, 1-13.	1.6	156
1671	Epitypification, morphology, and phylogeny of <i>Tothia fuscella</i>. <i>Mycotaxon</i> , 2012, 118, 203-211.	0.1	11
1672	Phylogeny and taxonomy of the genus <i>Gliocladiopsis</i>. <i>Persoonia: Molecular Phylogeny and Evolution of Fungi</i> , 2012, 28, 25-33.	1.6	18

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1673	Black Aspergilli and fumonisin contamination in onions purchased in Hungary. <i>Acta Alimentaria</i> , 2012, 41, 414-423.	0.3	22
1674	A 150 year-old mystery solved: Transfer of the rheophytic endemic liverwort <i>Myriocolea irrorata</i> to <i>Colura</i> . <i>Phytotaxa</i> , 2012, 66, 55.	0.1	29
1675	Phylogeny and proposed circumscription of <i>Breynia</i> , <i>Sauropus</i> and <i>Synostemon</i> (Phyllanthaceae), based on chloroplast and nuclear DNA sequences. <i>Australian Systematic Botany</i> , 2012, 25, 313.	0.3	30
1676	Integrative taxonomy justifies a new genus, <i>Nodastrella</i> gen. nov., for North Atlantic "Rossella" species (Porifera: Hexactinellida: Rossellidae). <i>Zootaxa</i> , 2012, 3383, .	0.2	20
1677	Diversification of the newly recognized lichen-forming fungal lineage <i>Montanelia</i> (Parmeliaceae, Ascomycota) and its relation to key geological and climatic events. <i>American Journal of Botany</i> , 2012, 99, 2014-2026.	0.8	51
1678	Molecular evolution and diversification of the moss family Daltoniaceae (Hookeriales, Bryophyta) with emphasis on the unravelling of the phylogeny of <i>Distichophyllum</i> and its allies. <i>Botanical Journal of the Linnean Society</i> , 2012, 170, 157-175.	0.8	11
1679	Phylogenetic placement of pelican spiders (Archaeidae, Araneae), with insight into evolution of the "neck" and predatory behaviours of the superfamily Palpimanoidea. <i>Cladistics</i> , 2012, 28, 598-626.	1.5	53
1680	Cryptic species in <i>Iphisa elegans</i> Gray, 1851 (Squamata: Gymnophthalmidae) revealed by hemipenial morphology and molecular data. <i>Zoological Journal of the Linnean Society</i> , 2012, 166, 361-376.	1.0	70
1681	The Parvidrilidae - a diversified groundwater family: description of six new species from southern Europe, and clues for its phylogenetic position within Clitellata (Annelida). <i>Zoological Journal of the Linnean Society</i> , 2012, 166, 530-558.	1.0	17
1682	Disentangling dispersal, vicariance and adaptive radiation patterns: A case study using armyworms in the pest genus <i>Spodoptera</i> (Lepidoptera: Noctuidae). <i>Molecular Phylogenetics and Evolution</i> , 2012, 65, 855-870.	1.2	82
1683	Sequence variation of the mitochondrial 12S rRNA gene among <i>Unionicola</i> ( <i>Wolcottatax</i> ) <i>arcuata</i> (Acari: Unionicolidae) from freshwater mussels in China. <i>International Journal of Acarology</i> , 2012, 38, 394-401.	0.3	5
1684	Species delimitation in annual killifishes from the Brazilian Caatinga, the <i>Hypsolebias flavicaudatus</i> complex (Cyprinodontiformes: Rivulidae): implications for taxonomy and conservation. <i>Systematics and Biodiversity</i> , 2012, 10, 71-91.	0.5	27
1685	Phylogeny-Based Species Delimitation In Philippine Slender Skinks (Reptilia: Squamata: Scincidae) III: Taxonomic Revision of the <i>Brachymeles Gracilis</i> Complex, With Descriptions of Three New Species. <i>Herpetological Monographs</i> , 2012, 26, 135-172.	1.1	18
1686	Authentication of the most important species of rockfish by means of fins. <i>European Food Research and Technology</i> , 2012, 235, 929-937.	1.6	5
1687	Freshwater ascomycetes: <i>Coniochaeta gigantospora</i> sp. nov. based on morphological and molecular data. <i>Mycoscience</i> , 2012, 53, 373-380.	0.3	14
1688	Molecular characterization of microalgae used in aquaculture with biotechnology potential. <i>Aquaculture International</i> , 2012, 20, 847-857.	1.1	15
1689	Basal and 3-methylcholanthrene-induced expression of cytochrome P450 1A, 1B and 1C genes in the Brazilian guppy, <i>Poecilia vivipara</i> . <i>Aquatic Toxicology</i> , 2012, 124-125, 106-113.	1.9	19
1690	<i>Lasiodiplodia</i> species associated with dieback disease of mango ( <i>Mangifera indica</i> ) in Egypt. <i>Australasian Plant Pathology</i> , 2012, 41, 649-660.	0.5	94

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1691	Molecular phylogeny of <i>Gyrodactylus</i> (Monogenea) parasitizing fishes in fresh water, estuarine, and marine habitats in Canada. <i>Canadian Journal of Zoology</i> , 2012, 90, 776-786.	0.4	37
1692	Overcoming Deep Roots, Fast Rates, and Short Internodes to Resolve the Ancient Rapid Radiation of Eupolypod II Ferns. <i>Systematic Biology</i> , 2012, 61, 490.	2.7	127
1693	The <i>Colletotrichum boninense</i> species complex. <i>Studies in Mycology</i> , 2012, 73, 1-36.	4.5	306
1694	The <i>Colletotrichum acutatum</i> species complex. <i>Studies in Mycology</i> , 2012, 73, 37-113.	4.5	656
1695	Can computational biology improve the phylogenetic analysis of insulin?. <i>Computer Methods and Programs in Biomedicine</i> , 2012, 108, 860-872.	2.6	8
1696	Hybridization between native and introduced species of deer in Eastern Europe. <i>Journal of Mammalogy</i> , 2012, 93, 1331-1341.	0.6	48
1697	Characterization of Human Endogenous Retroviral Elements in the Blood of HIV-1-Infected Individuals. <i>Journal of Virology</i> , 2012, 86, 262-276.	1.5	89
1698	Strain Characterization of <i>Potato virus S</i> Isolates from Tasmania, Australia. <i>Plant Disease</i> , 2012, 96, 813-819.	0.7	12
1699	Diversification in the Mexican horned lizard <i>Phrynosoma orbiculare</i> across a dynamic landscape. <i>Molecular Phylogenetics and Evolution</i> , 2012, 62, 87-96.	1.2	50
1700	Molecular systematics of sclerosomatid harvestmen (Opiliones, Phalangioidea, Sclerosomatidae): Geography is better than taxonomy in predicting phylogeny. <i>Molecular Phylogenetics and Evolution</i> , 2012, 62, 224-236.	1.2	56
1701	Relative roles of Neogene vicariance and Quaternary climate change on the historical diversification of bunchgrass lizards ( <i>Sceloporus scalaris</i> group) in Mexico. <i>Molecular Phylogenetics and Evolution</i> , 2012, 62, 447-457.	1.2	64
1702	Speciation in an avian complex endemic to the mountains of Middle America ( <i>Ergaticus</i> , Aves): Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 18	1.2	46
1703	Molecular phylogeny and morphometric analyses reveal deep divergence between Amazonia and Atlantic Forest species of <i>Dendrophryniscus</i> . <i>Molecular Phylogenetics and Evolution</i> , 2012, 62, 826-838.	1.2	79
1704	A phylogeny of the northern temperate leafy liverwort genus <i>Scapania</i> (Scapaniaceae,): Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 18	1.2	36
1705	A different perspective on the phylogenetic relationships of the Moxostomatini (Cypriniformes): Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 18	1.2	20
1706	Molecular phylogeny of Aphyocharacinae (Characiformes, Characidae) with morphological diagnoses for the subfamily and recognized genera. <i>Molecular Phylogenetics and Evolution</i> , 2012, 64, 297-307.	1.2	26
1707	Rapid range expansion in the Great Plains narrow-mouthed toad ( <i>Gastrophryne olivacea</i> ) and a revised taxonomy for North American microhylids. <i>Molecular Phylogenetics and Evolution</i> , 2012, 64, 645-653.	1.2	18
1708	The genus <i>Cladosporium</i> . <i>Studies in Mycology</i> , 2012, 72, 1-401.	4.5	521

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1709	SATÁ©-II: Very Fast and Accurate Simultaneous Estimation of Multiple Sequence Alignments and Phylogenetic Trees. <i>Systematic Biology</i> , 2012, 61, 90.	2.7	310
1710	From Amazonia to the Atlantic forest: Molecular phylogeny of <i>Phyzelaphryninae</i> frogs reveals unexpected diversity and a striking biogeographic pattern emphasizing conservation challenges. <i>Molecular Phylogenetics and Evolution</i> , 2012, 65, 547-561.	1.2	124
1711	The biogeographic legacy of an imperilled taxon provides a foundation for assessing lineage diversification, demography and conservation genetics. <i>Diversity and Distributions</i> , 2012, 18, 689-703.	1.9	12
1712	Experimental Design in Phylogenetics: Testing Predictions from Expected Information. <i>Systematic Biology</i> , 2012, 61, 661-674.	2.7	14
1713	Accuracy and Precision of Species Trees: Effects of Locus, Individual, and Base Pair Sampling on Inference of Species Trees in Lizards of the <i>Liolaemus darwini</i> Group (Squamata, Liolaemidae). <i>Systematic Biology</i> , 2012, 61, 272.	2.7	86
1714	Cathepsin B from the white shrimp <i>Litopenaeus vannamei</i> : cDNA sequence analysis, tissues-specific expression and biological activity. <i>Comparative Biochemistry and Physiology - B Biochemistry and Molecular Biology</i> , 2012, 161, 32-40.	0.7	41
1715	Vascular Streak Dieback of cacao in Southeast Asia and Melanesia: in planta detection of the pathogen and a new taxonomy. <i>Fungal Biology</i> , 2012, 116, 11-23.	1.1	48
1716	Species limits in the Andean toad genus <i>Osornophryne</i> (Bufonidae). <i>Molecular Phylogenetics and Evolution</i> , 2012, 65, 805-822.	1.2	17
1717	Morphological characterization and molecular fingerprinting of <i>Nostoc</i> strains by multiplex RAPD. <i>Microbiology</i> , 2012, 81, 710-720.	0.5	4
1718	Quartet decomposition server: a platform for analyzing phylogenetic trees. <i>BMC Bioinformatics</i> , 2012, 13, 123.	1.2	6
1719	Evolutionarily consistent families in SCOP: sequence, structure and function. <i>BMC Structural Biology</i> , 2012, 12, 27.	2.3	14
1720	Phylogeny of Chaetothyriaceae in northern Thailand including three new species. <i>Mycologia</i> , 2012, 104, 382-395.	0.8	44
1721	Molecular phylogeny of <i>Sydowiellaceae</i> resolving the position of <i>Cainiella</i> . <i>Mycologia</i> , 2012, 104, 419-426.	0.8	9
1722	Phylogenetic and mixed Yule-coalescent analyses reveal cryptic lineages within two South American marine snails of the genus <i>Crepidatella</i> (Gastropoda: Calyptraeidae). <i>Invertebrate Biology</i> , 2012, 131, 301-311.	0.3	10
1723	First molecular phylogenetic insights into the evolution of free-tailed bats in the subfamily Molossinae (Molossidae, Chiroptera). <i>Journal of Mammalogy</i> , 2012, 93, 12-28.	0.6	55
1724	Phylogeography of the Teiid Lizard <i>Kentropyx calcarata</i> and the Sphaerodactylid <i>Gonatodes humeralis</i> (Reptilia: Squamata): Testing A Geological Scenario for the Lower Amazon-Tocantins Basins, Amazonia, Brazil. <i>Herpetologica</i> , 2012, 68, 272.	0.2	11
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1726	Molecular phylogenetics, species diversity, and biogeography of the Andean lizards of the genus <i>Proctoporus</i> (Squamata: Gymnophthalmidae). <i>Molecular Phylogenetics and Evolution</i> , 2012, 65, 953-964.	1.2	28



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1728	The biological roles of glutaredoxins. <i>Biochemical Journal</i> , 2012, 446, 333-348.	1.7	100
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1731	Congruence and indifference between two molecular markers for understanding oral evolution in the Marynidae sensu lato (Ciliophora, Colpodea). <i>European Journal of Protistology</i> , 2012, 48, 297-304.	0.5	13
1732	Multiple origins of sequestrate basidiomes within <i>Entoloma</i> inferred from molecular phylogenetic analyses. <i>Fungal Biology</i> , 2012, 116, 1250-1262.	1.1	18
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1734	The Essentials of Computational Molecular Evolution. <i>Methods in Molecular Biology</i> , 2012, 855, 111-152.	0.4	28
1736	Lindane biodegradation by the <i>Fusarium verticillioides</i> AT-100 strain, isolated from <i>Agave tequilana</i> leaves: Kinetic study and identification of metabolites. <i>International Biodeterioration and Biodegradation</i> , 2012, 74, 36-47.	1.9	51
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1738	Genetic and Morphologic Variation in the Davis Mountains Cottontail ( <i>Sylvilagus robustus</i> ). <i>Southwestern Naturalist</i> , 2012, 57, 1-7.	0.1	5
1739	Sequence Data Reveals Phylogenetic Affinities of <i>Acrocallymma aquaticasp. nov.</i> , <i>Aquasubmersa mircensisgen. et sp. nov.</i> and <i>Clohesyomyces aquaticus</i> (Freshwater Coelomycetes). <i>Cryptogamie, Mycologie</i> , 2012, 33, 333-346.	0.2	37
1740	<i>Bambusicola</i> , a New Genus from Bamboo with Asexual and Sexual Morphs. <i>Cryptogamie, Mycologie</i> , 2012, 33, 363-379.	0.2	45
1741	Estimating divergence dates and evaluating dating methods using phylogenomic and mitochondrial data in squamate reptiles. <i>Molecular Phylogenetics and Evolution</i> , 2012, 65, 974-991.	1.2	141
1742	Molecular phylogeny of microhylid frogs (Anura: Microhylidae) with emphasis on relationships among New World genera. <i>BMC Evolutionary Biology</i> , 2012, 12, 241.	3.2	72
1743	Towards a natural classification of Botryosphaeraiales. <i>Fungal Diversity</i> , 2012, 57, 149-210.	4.7	198
1744	Trichomeriaceae, a new sooty mould family of Chaetothyriales. <i>Fungal Diversity</i> , 2012, 56, 63-76.	4.7	58
1745	Phyllosticta species associated with freckle disease of banana. <i>Fungal Diversity</i> , 2012, 56, 173-187.	4.7	52

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1747	Phylogeography of the Endangered Otago Skink, <i>Oligosoma ottagense</i> : Population Structure, Hybridisation and Genetic Diversity in Captive Populations. PLoS ONE, 2012, 7, e34599.	1.1	16
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1755	Multiple Quaternary Refugia in the Eastern Guiana Shield Revealed by Comparative Phylogeography of 12 Frog Species. Systematic Biology, 2012, 61, 461.	2.7	113
1756	<i>Tremella diploschistina</i> ( <i>Tremellales</i> , Basidiomycota, Fungi), a new lichenicolous species growing on <i>Diploschistes</i> . Lichenologist, 2012, 44, 321-332.	0.5	24
1757	Molecular phylogeny of glass sponges (Porifera, Hexactinellida): increased taxon sampling and inclusion of the mitochondrial protein-coding gene, cytochrome oxidase subunit I. Hydrobiologia, 2012, 687, 11-20.	1.0	24
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1759	Molecular Identification and Antifungal Susceptibilities of Black Aspergillus Isolates from Otomycosis Cases in Hungary. Mycopathologia, 2012, 174, 143-147.	1.3	48
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1765	Genetic diversity and insular colonization of <i>Liolaemus pictus</i> (Squamata, Liolaeminae) in northwestern Patagonia. <i>Austral Ecology</i> , 2012, 37, 67-77.	0.7	14
1766	Evaluation of High-resolution Melting Curve Analysis as a New Tool for Root-knot Nematode Diagnostics. <i>Journal of Phytopathology</i> , 2012, 160, 59-66.	0.5	34
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1772	Phylogenetic analyses and toxigenic profiles of <i>Fusarium equiseti</i> and <i>Fusarium acuminatum</i> isolated from cereals from Southern Europe. <i>Food Microbiology</i> , 2012, 31, 229-237.	2.1	72
1773	Molecular identification and population dynamic of <i>Anisakis pegreffii</i> (Nematoda: Anisakidae Dujardin). <i>Journal of Food Microbiology</i> , 2012, 157, 224-229.	2.1	48
1774	Evolution of the bomolochiform superfamily complex (Copepoda: Cyclopoida): New insights from ssrDNA and morphology, and origin of umazuracolids from polychaete-infesting ancestors rejected. <i>International Journal for Parasitology</i> , 2012, 42, 71-92.	1.3	40
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1780	Phylogeography of the catfish <i>Hatcheria macraei</i> reveals a negligible role of drainage divides in structuring populations. <i>Molecular Ecology</i> , 2012, 21, 942-959.	2.0	32
1781	Coalescence patterns of endemic Tibetan species of stream salamanders (Hynobiidae). <i>Journal of Biogeography</i> , 2012, 39, 1707-1719.	2.0	35

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1783	Phylogeny and biogeography of the parrot genus <i>Prioniturus</i> (Aves: Psittaciformes). <i>Journal of Zoological Systematics and Evolutionary Research</i> , 2012, 50, 145-156.	0.6	12
1784	Species assignment and antifungal susceptibilities of black aspergilli recovered from otomycosis cases in Iran. <i>Mycoses</i> , 2012, 55, 333-338.	1.8	49
1785	Phylogenetic position of gastrostomobdellid leeches (Hirudinida, Arhynchobdellida,) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Jf 50 62 23	0.7	23
1786	Coâ€œcirculation of two Puumala hantavirus lineages in Latvia: A russian lineage described previously and a novel Latvian lineage. <i>Journal of Medical Virology</i> , 2012, 84, 314-318.	2.5	22
1787	<i>Trichoderma stromaticum</i> and its overseas relatives. <i>Mycological Progress</i> , 2012, 11, 215-254.	0.5	27
1788	Phylogeny of the industrial relevant, thermophilic genera <i>Myceliophthora</i> and <i>Corynascus</i> . <i>Fungal Diversity</i> , 2012, 52, 197-207.	4.7	59
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1790	A study of genetic variations, population size, and population dynamics of the catadromous Japanese eel <i>Anguilla japonica</i> (Pisces) in northern Taiwan. <i>Hydrobiologia</i> , 2012, 683, 203-216.	1.0	7
1791	Molecular characterization of peste des petits ruminants virus from the Karamoja region of Uganda (2007-2008). <i>Archives of Virology</i> , 2012, 157, 29-35.	0.9	27
1792	Fineâ€œscale biogeographical and temporal diversification processes of peacock swallowtails ( <i>Papilio</i> subgenus <i>Achillides</i> ) in the Indoâ€œAustralian Archipelago. <i>Cladistics</i> , 2013, 29, 88-111.	1.5	43
1793	How to contend with paraphyly in the taxonomy of the delphinine cetaceans?. <i>Marine Mammal Science</i> , 2013, 29, 567-588.	0.9	27
1794	<i>Botryosphaeria</i> , <i>Neofusicoccum</i> , <i>Neoscytalidium</i> and <i>Pseudofusicoccum</i> species associated with mango in Brazil. <i>Fungal Diversity</i> , 2013, 61, 195-208.	4.7	62
1795	The <i>Colletotrichum orbiculare</i> species complex: Important pathogens of field crops and weeds. <i>Fungal Diversity</i> , 2013, 61, 29-59.	4.7	90
1796	<i>Ilyonectria</i> black foot rot associated with Proteaceae. <i>Australasian Plant Pathology</i> , 2013, 42, 337-349.	0.5	23
1797	Does Computational Biology Help us to Understand the Molecular Phylogenetics and Evolution of Cluster of Differentiation (CD) Proteins?. <i>Protein Journal</i> , 2013, 32, 143-154.	0.7	2
1798	A phylogenetic analysis of Bovine Viral Diarrhoea Virus (BVDV) isolates from six different regions of the UK and links to animal movement data. <i>Veterinary Research</i> , 2013, 44, 43.	1.1	49
1799	Molecular systematics of bonneted bats (Molossidae:Eumops) based on mitochondrial and nuclear DNA sequences. <i>Journal of Mammalogy</i> , 2013, 94, 867-880.	0.6	15

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1801	Quantification and relative severity of inflated branch-support values generated by alternative methods: An empirical example. <i>Molecular Phylogenetics and Evolution</i> , 2013, 67, 277-296.	1.2	27
1802	First Record of the Genus <i>Oreolalax</i> (Anura: Megophryidae) from Vietnam with Description of a New Species. <i>Copeia</i> , 2013, 2013, 213-222.	1.4	14
1803	Phylogenetic relationships of the enigmatic longtailed rattlesnakes ( <i>Crotalus ericsmithi</i> , <i>C. lannomi</i> ). <i>Tj ETQq1 1 0.784314 rgBT /Over</i>	1.2	36
1804	Metal corrosion by aerobic bacteria isolated from stimulated corrosion systems: Effects of additional nitrate sources. <i>International Biodeterioration and Biodegradation</i> , 2013, 83, 158-165.	1.9	33
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1806	Testing robustness of relative complexity measure method constructing robust phylogenetic trees for <i>Galanthus L.</i> Using the relative complexity measure. <i>BMC Bioinformatics</i> , 2013, 14, 20.	1.2	5
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1808	Freshwater Ascomycetes: <i>Minutisphaera</i> (Dothideomycetes) revisited, including one new species from Japan. <i>Mycologia</i> , 2013, 105, 959-976.	0.8	28
1809	Identification of <i>Anisakis</i> species (Nematoda: Anisakidae) in marine fish hosts from Papua New Guinea. <i>Veterinary Parasitology</i> , 2013, 193, 126-133.	0.7	27
1810	Morphological analysis of phylogenetic relationships among extant rhynchonellide brachiopods. <i>Journal of Paleontology</i> , 2013, 87, 550-569.	0.5	15
1811	Phylogenetic analysis, fumonisin production and pathogenicity of <i>Fusarium fujikuroi</i> strains isolated from rice in the Philippines. <i>Journal of the Science of Food and Agriculture</i> , 2013, 93, 3032-3039.	1.7	29
1812	Quantification and in situ localisation of <i>abcb1</i> and <i>abcc9</i> genes in toxicant-exposed sea urchin embryos. <i>Environmental Science and Pollution Research</i> , 2013, 20, 8600-8611.	2.7	7
1813	<i>Diplodia quercivora</i> sp. nov.: a new species of <i>Diplodia</i> found on declining <i>Quercus canariensis</i> trees in Tunisia. <i>Mycologia</i> , 2013, 105, 1266-1274.	0.8	48
1814	A new genus of megalonychid sloth (Mammalia, Xenarthra) from the late Pleistocene (Lujanian) of Sierra de Perija, Zulia State, Venezuela. <i>Journal of Vertebrate Paleontology</i> , 2013, 33, 1226-1238.	0.4	32
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1816	Sky Islands of the Cameroon Volcanic Line: a diversification hot spot for puddle frogs ( <i>Phrynobatrachidae</i> : <i>Phrynobatrachus</i> ). <i>Zoologica Scripta</i> , 2013, 42, 591-611.	0.7	15
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1819	Phylogeny of the non-monophyletic <i>Cayratia</i> Juss. (Vitaceae) and implications for character evolution and biogeography. <i>Molecular Phylogenetics and Evolution</i> , 2013, 68, 502-515.	1.2	54
1820	Phylogeography of <i>Coreoperca whiteheadi</i> (Perciformes: Coreoperca) in China based on mitochondrial and nuclear gene sequences. <i>Biochemical Systematics and Ecology</i> , 2013, 50, 223-231.	0.6	12
1821	A molecular survey of a captive wallaby population for periodontopathogens and the co-incidence of <i>Fusobacterium necrophorum</i> subspecies <i>necrophorum</i> with periodontal diseases. <i>Veterinary Microbiology</i> , 2013, 163, 335-343.	0.8	17
1822	Molecular characterization of <i>Fasciola hepatica</i> from Sardinia based on sequence analysis of genomic and mitochondrial gene markers. <i>Experimental Parasitology</i> , 2013, 135, 471-478.	0.5	17
1823	A Molecular Phylogeny of the Species-Rich Neotropical Genus <i>Anthurium</i> (Araceae) based on Combined Chloroplast and Nuclear DNA. <i>Systematic Botany</i> , 2013, 38, 576-588.	0.2	41
1824	Increased phylogenetic diversity of bovine viral diarrhoea virus type 1 isolates in England and Wales since 2001. <i>Veterinary Microbiology</i> , 2013, 162, 315-320.	0.8	26
1825	Planistromellaceae (Botryosphaerales). <i>Cryptogamie, Mycologie</i> , 2013, 34, 45.	0.2	13
1826	Phenotypic, Molecular, and Pathological Characterization of <i>Colletotrichum acutatum</i> Associated with Andean Lupine and Tamarillo in the Ecuadorian Andes. <i>Plant Disease</i> , 2013, 97, 819-827.	0.7	20
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1829	Families of Dothideomycetes. <i>Fungal Diversity</i> , 2013, 63, 1-313.	4.7	509
1830	Reevaluation of the anatomy of the Cenomanian (Upper Cretaceous) hind-limbed marine fossil snakes <i>Pachyrhachis</i> , <i>Haasiophis</i> , and <i>Eupodophis</i> . <i>Journal of Vertebrate Paleontology</i> , 2013, 33, 1328-1342.	0.4	19
1831	Diversity of <i>Botryosphaeriaceae</i> species associated with conifers in Portugal. <i>European Journal of Plant Pathology</i> , 2013, 135, 791-804.	0.8	29
1832	Characterisation and pathogenicity of <i>Pestalotiopsis uvicola</i> and <i>Pestalotiopsis clavispora</i> causing grey leaf spot of mango ( <i>Mangifera indica</i> L.) in Italy. <i>European Journal of Plant Pathology</i> , 2013, 135, 619-625.	0.8	39
1833	The evolutionary history of the order Antipatharia (Cnidaria: Anthozoa: Hexacorallia) as inferred from mitochondrial and nuclear DNA: implications for black coral taxonomy and systematics. <i>Zoological Journal of the Linnean Society</i> , 2013, 169, 312-361.	1.0	62
1834	Genetic population structure in <i>Siniperca scherzeri</i> (Perciformes: Siniperca) in China inferred from mitochondrial DNA sequences and microsatellite loci. <i>Biochemical Systematics and Ecology</i> , 2013, 51, 160-170.	0.6	4
1835	Human rhinoviruses in INDISâ€”study materialâ€”evidence for recovery of viable rhinovirus from fecal specimens. <i>Journal of Medical Virology</i> , 2013, 85, 1466-1472.	2.5	11

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1836	Phylogeny of the <i>Oenanthe lugens</i> complex (Aves, Muscicapidae: Saxicolinae): Paraphyly of a morphologically cohesive group within a recent radiation of open-habitat chats. <i>Molecular Phylogenetics and Evolution</i> , 2013, 69, 450-461.	1.2	20
1837	Phylogeny and Morphology of <i>Leptosphaerulina saccharicola</i> sp. nov. and <i>Pleosphaerulina oryzae</i> and Relationships with <i>Pithomyces</i> . <i>Cryptogamie, Mycologie</i> , 2013, 34, 303-319.	0.2	18
1838	Multi-Gene Analyses Reveal Taxonomic Placement of <i>Scolicosporium minkeviciusii</i> in Phaeosphaeriaceae (Pleosporales). <i>Cryptogamie, Mycologie</i> , 2013, 34, 357-366.	0.2	11
1839	DNA barcoding common non-native freshwater fish species in Turkey: Low genetic diversity but high population structuring. <i>Mitochondrial DNA</i> , 2013, 24, 276-287.	0.6	23
1840	The relationships of <i>Odontotrema</i> (Odontotremataceae) and the resurrected <i>Sphaeropezia</i> (Stictidaceae) – new combinations and three new <i>Sphaeropezia</i> species. <i>Mycologia</i> , 2013, 105, 384-397.	0.8	26
1841	Emended diagnosis and phylogenetic relationships of the Upper Cretaceous fossil snake <i>Najash rionegrina</i> Apesteguía and Zaher, 2006. <i>Journal of Vertebrate Paleontology</i> , 2013, 33, 131-140.	0.4	25
1842	Genus-level phylogeny of snakes reveals the origins of species richness in Sri Lanka. <i>Molecular Phylogenetics and Evolution</i> , 2013, 66, 969-978.	1.2	86
1843	Multilocus phylogeny and MALDI-TOF analysis of the plant pathogenic species <i>Alternaria dauci</i> and relatives. <i>Fungal Biology</i> , 2013, 117, 32-40.	1.1	40
1844	Building megaphylogenies for macroecology: taking up the challenge. <i>Ecography</i> , 2013, 36, 13-26.	2.1	79
1845	Mazaedium evolution in the Ascomycota (Fungi) and the classification of mazaediate groups of formerly unclear relationship. <i>Cladistics</i> , 2013, 29, 296-308.	1.5	65
1846	Out of the Bassian province: historical biogeography of the Australasian platycercine parrots (Aves). <i>Trends in Ecology and Evolution</i> , 2013, 28, 19-27.	8.7	19
1847	Phylogenetic and taxonomic relationships of the <i>Polypedates leucomystax</i> complex (Amphibia). <i>Zoologica Scripta</i> , 2013, 42, 54-70.	0.7	35
1848	Authentication of gadooids from highly processed products susceptible to include species mixtures by means of DNA sequencing methods. <i>European Food Research and Technology</i> , 2013, 236, 171-180.	1.6	7
1849	Multi-gene region phylogenetic analyses suggest reticulate evolution and a clade of Australian origin among paleotropical woody bamboos (Poaceae: Bambusoideae: Bambuseae). <i>Plant Systematics and Evolution</i> , 2013, 299, 239-257.	0.3	33
1850	Incidence, severity and causal fungal species of <i>Mycosphaerella</i> and <i>Teratosphaeria</i> diseases in Eucalyptus stands in Galicia (NW Spain). <i>Forest Ecology and Management</i> , 2013, 302, 379-389.	1.4	7
1851	Automated subtyping of HIV-1 genetic sequences for clinical and surveillance purposes: Performance evaluation of the new REGA version 3 and seven other tools. <i>Infection, Genetics and Evolution</i> , 2013, 19, 337-348.	1.0	313
1852	<i>Lindgomyces angustiascus</i> , (Lindgomycetaceae, Pleosporales, Dothideomycetes), a new lignicolous species from freshwater habitats in the USA. <i>Mycoscience</i> , 2013, 54, 353-361.	0.3	25
1853	Molecular characterization of major and minor rDNA repeats and genetic variability assessment in different species of mahseer found in North India. <i>Gene</i> , 2013, 527, 248-258.	1.0	10

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1854	Active site characterization and molecular cloning of <i>Tenebrio molitor</i> midgut trehalase and comments on their insect homologs. <i>Insect Biochemistry and Molecular Biology</i> , 2013, 43, 768-780.	1.2	20
1855	Species concepts in <i>Cercospora</i> : spotting the weeds among the roses. <i>Studies in Mycology</i> , 2013, 75, 115-170.	4.5	290
1856	Phylogenetic lineages in <i>Pseudocercospora</i> . <i>Studies in Mycology</i> , 2013, 75, 37-114.	4.5	175
1857	A phylogenetic re-evaluation of <i>Phyllosticta</i> (Botryosphaerales). <i>Studies in Mycology</i> , 2013, 76, 1-29.	4.5	104
1858	The complete mitochondrial genome of <i>Biston panterinaria</i> (Lepidoptera: Geometridae), with phylogenetic utility of mitochondrial genome in the Lepidoptera. <i>Gene</i> , 2013, 515, 349-358.	1.0	61
1859	Identification of novel and zoonotic <i>Cryptosporidium</i> species in fish from Papua New Guinea. <i>Veterinary Parasitology</i> , 2013, 198, 1-9.	0.7	45
1860	A wide geographical survey of mitochondrial DNA variation in the great spotted woodpecker complex, <i>Dendrocopos major</i> (Aves: Picidae). <i>Biological Journal of the Linnean Society</i> , 2013, 108, 173-188.	0.7	23
1861	Molecular identification and larval morphological description of <i>Contraecaecum pelagicum</i> (Nematoda: Anisakidae) from the anchovy <i>Engraulis anchoita</i> (Engraulidae) and fish-eating birds from the Argentine North Patagonian Sea. <i>Parasitology International</i> , 2013, 62, 309-319.	0.6	23
1862	An extraordinary tail - integrative review of the agamid genus <i>Xenagama</i> . <i>Journal of Zoological Systematics and Evolutionary Research</i> , 2013, 51, 144-164.	0.6	9
1863	Sequence-based molecular phylogenetics and phylogeography of the American box turtles ( <i>Terrapene</i> ) Tj ETQq1 1 0.784314 rrgBT /Ov	1.2	39
1864	Genome-wide <i>scp&gt;RAD&lt;/scp&gt;</i> sequence data provide unprecedented resolution of species boundaries and relationships in the <i>scp&gt;L&lt;/scp&gt;</i> ake <i>scp&gt;V&lt;/scp&gt;</i> ictoria cichlid adaptive radiation. <i>Molecular Ecology</i> , 2013, 22, 787-798.	2.0	415
1865	Molecular phylogeny of <i>Trissolcus</i> species (Hymenoptera: Scelionidae). <i>Biochemical Systematics and Ecology</i> , 2013, 48, 85-91.	0.6	12
1866	Phylogeny and taxonomy of <i>Botryosphaeria</i> and <i>Neofusicoccum</i> species in Iran, with description of <i>Botryosphaeria scharifii</i> sp. nov.. <i>Mycologia</i> , 2013, 105, 210-220.	0.8	50
1867	TESTING THE MUSEUM VERSUS CRADLE TROPICAL BIOLOGICAL DIVERSITY HYPOTHESIS: PHYLOGENY, DIVERSIFICATION, AND ANCESTRAL BIOGEOGRAPHIC RANGE EVOLUTION OF THE ANTS. <i>Evolution; International Journal of Organic Evolution</i> , 2013, 67, 2240-2257.	1.1	290
1868	Phylogenetic analysis of <i>Lappula</i> Moench (Boraginaceae) based on molecular and morphological data. <i>Plant Systematics and Evolution</i> , 2013, 299, 913-926.	0.3	9
1869	Structural variation and dynamics of the nuclear ribosomal intergenic spacer region in key members of the <i>Gibberella fujikuroi</i> species complex. <i>Genome</i> , 2013, 56, 205-213.	0.9	10
1870	Species of <i>Lasiodiplodia</i> associated with mango in Brazil. <i>Fungal Diversity</i> , 2013, 61, 181-193.	4.7	96
1871	<i>Stachybotrys</i> from soil in China, identified by morphology and molecular phylogeny. <i>Mycological Progress</i> , 2013, 12, 693-698.	0.5	8



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1872	Phylogeny and fruit evolution in Menispermaceae. American Journal of Botany, 2013, 100, 883-905.	0.8	33
1873	Diversification patterns and processes of wingless endemic insects in the Mediterranean Basin: historical biogeography of the genus <i>Blaps</i> (Coleoptera: Tenebrionidae). Journal of Biogeography, 2013, 40, 1899-1913.	1.4	25
1874	A new species of <i>Gracixalus</i> (Amphibia: Anura: Rhacophoridae) from northern Vietnam. Organisms Diversity and Evolution, 2013, 13, 203-214.	0.7	15
1875	Identifying localized biases in large datasets: A case study using the avian tree of life. Molecular Phylogenetics and Evolution, 2013, 69, 1021-1032.	1.2	78
1876	Combined Morphological and Molecular Phylogeny of the Clusioid Clade (Malpighiales) and the Placement of the Ancient Rosid Macrofossil <i>Paleoclusia</i> . International Journal of Plant Sciences, 2013, 174, 910-936.	0.6	39
1877	Biodiversity in wood-decay macro-fungi associated with declining arid zone trees of India as revealed by nuclear rDNA analysis. European Journal of Plant Pathology, 2013, 136, 373-382.	0.8	13
1878	Assessing statistical reliability of phylogenetic trees via a speedy double bootstrap method. Molecular Phylogenetics and Evolution, 2013, 67, 429-435.	1.2	5
1879	Phylogeny as a Proxy for Ecology in Seagrass Amphipods: Which Traits Are Most Conserved?. PLoS ONE, 2013, 8, e57550.	1.1	37
1880	High phylogeographic structure in sylvatic vectors of Chagas disease of the genus <i>Mepraia</i> (Hemiptera: Reduviidae). Infection, Genetics and Evolution, 2013, 19, 280-286.	1.0	22
1881	A new genus and subfamily of mosasaurs from the Upper Cretaceous of northern Italy. Journal of Vertebrate Paleontology, 2013, 33, 599-612.	0.4	49
1882	Resolving the phylogenetic history of the short-necked turtles, genera <i>Elseya</i> and <i>Myuchelys</i> (Testudines: Chelidae) from Australia and New Guinea. Molecular Phylogenetics and Evolution, 2013, 68, 251-258.	1.2	16
1883	Molecular phylogenetics and phylogeographic structure of <i>Sumichrasti</i> 's harvest mouse ( <i>Reithrodontomys sumichrasti</i> : Cricetidae) based on mitochondrial and nuclear DNA sequences. Molecular Phylogenetics and Evolution, 2013, 68, 282-292.	1.2	22
1884	The enigmatic truffle <i>Fevansia aurantiaca</i> is an ectomycorrhizal member of the <i>Albatrellus</i> lineage. Mycorrhiza, 2013, 23, 663-668.	1.3	8
1885	Life-history effects of arsenic toxicity in clades of the earthworm <i>Lumbricus rubellus</i> . Environmental Pollution, 2013, 172, 200-207.	3.7	24
1886	Molecular Phylogeny of Mulletts (Teleostei: Mugilidae) in Iran Based on Mitochondrial DNA. Biochemical Genetics, 2013, 51, 334-340.	0.8	3
1887	Structure and enzymatic mechanism of a moonlighting dUTPase. Acta Crystallographica Section D: Biological Crystallography, 2013, 69, 2298-2308.	2.5	21
1889	Molecular phylogeny of the Iranian Plateau five-toed jerboa, <i>Allactaga</i> (Dipodidea: Rodentia), inferred from mtDNA. Mammalia, 2013, 77, .	0.3	11
1890	A new <i>Myrmecridium</i> species from Guizhou, China. Mycotaxon, 2013, 124, 1-8.	0.1	8

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1891	Phylogenetic Analysis Reveals a High Prevalence of <i>Sporothrix brasiliensis</i> in Feline Sporotrichosis Outbreaks. <i>PLoS Neglected Tropical Diseases</i> , 2013, 7, e2281.	1.3	223
1892	Incorporating trnH-psbA to the core DNA barcodes improves significantly species discrimination within southern African Combretaceae. <i>ZooKeys</i> , 2013, 365, 129-147.	0.5	34
1893	DNA barcodes and phylogenetic affinities of the terrestrial slugs <i>Arion gilvus</i> and <i>A. ponsi</i> (Gastropoda, Pulmonata, Arionidae). <i>ZooKeys</i> , 2013, 365, 83-104.	0.5	3
1894	Diversity of <i>Ptychadena</i> in Rwanda and taxonomic status of <i>P. chrysogaster</i> Laurent, 1954 (Amphibia.) <i>TJ ETQq1 1 0.784314 rgBT /Ov</i>	0.5	4
1895	Efficacy of the core DNA barcodes in identifying processed and poorly conserved plant materials commonly used in South African traditional medicine. <i>ZooKeys</i> , 2013, 365, 215-233.	0.5	16
1896	&lt;l&gt;Diaporthe&lt;/l&gt;; a genus of endophytic, saprobic and plant pathogenic fungi. <i>Persoonia: Molecular Phylogeny and Evolution of Fungi</i> , 2013, 31, 1-41.	1.6	468
1897	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
1898	Massive Mitochondrial Gene Transfer in a Parasitic Flowering Plant Clade. <i>PLoS Genetics</i> , 2013, 9, e1003265.	1.5	115
1899	Molecular and Morphological Data Support the Existence of a Sexual Cycle in Species of the Genus <i>Paracoccidioides</i> . <i>Eukaryotic Cell</i> , 2013, 12, 380-389.	3.4	38
1900	Molecular Phylogeny of the Leafy Liverwort <i>Lejeunea</i> (Porellales): Evidence for a Neotropical Origin, Uneven Distribution of Sexual Systems and Insufficient Taxonomy. <i>PLoS ONE</i> , 2013, 8, e82547.	1.1	53
1901	The role of numeracy in moderating the influence of statistics in climate change messages. <i>Public Understanding of Science</i> , 2013, 22, 785-798.	1.6	37
1902	<i>Calonectria metrosideri</i>, a highly aggressive pathogen causing leaf blight, root rot, and wilt of <i>Metrosideros</i> spp. in <i>Brazil</i>. <i>Forest Pathology</i> , 2013, 43, 257-265.	0.5	20
1903	Tracking the impact of Pliocene/Pleistocene sea level and climatic oscillations on the cladogenesis of the Cape legless skink, <i>Acontias meleagris</i> species complex, in South Africa. <i>Journal of Biogeography</i> , 2013, 40, 492-506.	1.4	23
1904	Evaluating molecular support for and against the monophyly of the Peritrichia and phylogenetic relationships within the Mobilida (Ciliophora, Oligohymenophorea). <i>Zoologica Scripta</i> , 2013, 42, 213-226.	0.7	29
1905	Morphological and Genetic Identification of Spontaneously Spawned Larvae of Captive Bluefin Tuna in the Adriatic Sea. <i>Fisheries</i> , 2013, 38, 410-417.	0.6	8
1906	Molecular and Genetic Evidence for a Tetrapolar Mating System in the Basidiomycetous Yeast <i>Kwoniella mangrovensis</i> and Two Novel Sibling Species. <i>Eukaryotic Cell</i> , 2013, 12, 746-760.	3.4	25
1907	Uranium (U)-Tolerant Bacterial Diversity from U Ore Deposit of Domiasiat in North-East India and Its Prospective Utilisation in Bioremediation. <i>Microbes and Environments</i> , 2013, 28, 33-41.	0.7	61
1908	Full Issue PDF Volume 38, Issue 9. <i>Fisheries</i> , 2013, 38, 389-428.	0.6	1

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1909	<i>Colletotrichum fructicola</i> , first record of bitter rot of apple in China. Mycotaxon, 2014, 126, 23-30.	0.1	9
1910	Size doesn't matter—recircumscription of <i>Microlejeunea</i> (Lejeuneaceae, Porellales) based on molecular and morphological evidence. Phytotaxa, 2013, 85, 41.	0.1	44
1911	A new species of <i>Calonectria</i> ; causing leaf disease of water lily in China. Mycotaxon, 2013, 122, 177-185.	0.1	14
1912	Generic circumscriptions in <i>Geoglossomycetes</i> . Persoonia: Molecular Phylogeny and Evolution of Fungi, 2013, 31, 101-111.	1.6	23
1913	Exploring Species Level Taxonomy and Species Delimitation Methods in the Facultatively Self-Fertilizing Land Snail Genus <i>Rumina</i> (Gastropoda: Pulmonata). PLoS ONE, 2013, 8, e60736.	1.1	68
1914	Multiple Glacial Refugia of the Low-Dispersal Ground Beetle <i>Carabus irregularis</i> : Molecular Data Support Predictions of Species Distribution Models. PLoS ONE, 2013, 8, e61185.	1.1	51
1915	Patterns of Genetic and Reproductive Traits Differentiation in Mainland vs. Corsican Populations of Bumblebees. PLoS ONE, 2013, 8, e65642.	1.1	72
1916	Molecular Phylogeny and Biogeography of <i>Petaurista</i> Inferred from the Cytochrome b Gene, with Implications for the Taxonomic Status of <i>P. caniceps</i> , <i>P. marica</i> and <i>P. sybilla</i> . PLoS ONE, 2013, 8, e70461.	1.1	13
1917	Support Measures to Estimate the Reliability of Evolutionary Events Predicted by Reconciliation Methods. PLoS ONE, 2013, 8, e73667.	1.1	22
1918	The Evolution and Expression of the Moth Visual Opsin Family. PLoS ONE, 2013, 8, e78140.	1.1	20
1919	Phylogeny of the <i>Cyrtodactylus irregularis</i> species complex (Squamata: Gekkonidae) from Vietnam with the description of two new species. Zootaxa, 2013, 3737, 399.	0.2	29
1920	To name or not to name: Criteria to promote economy of change in Linnaean classification schemes. Zootaxa, 2013, 3636, 201-44.	0.2	170
1921	A new species of the <i>Gekko japonicus</i> group (Squamata: Sauria: Gekkonidae) from the border region between China and Vietnam. Zootaxa, 2013, 3652, 501.	0.2	13
1922	A new species of <i>Calonectria</i> causing leaf blight and cutting rot of three forest tree species in Brazil. Tropical Plant Pathology, 2013, 38, 513-521.	0.8	22
1923	Transfer of <i>Lejeunea huctumalcensis</i> to <i>Physantholejeunea</i> (Lejeuneaceae, Porellales). Australian Systematic Botany, 2013, 26, 386.	0.3	15
1924	The Fatal Case of Lyssavirus Encephalitis in the Russian Far East. , 2013, , .		2
1925	Systematics and biogeography of <i>Sternarchellini</i> (Gymnotiformes: Aptereronotidae): Diversification of electric fishes in large Amazonian rivers. Neotropical Ichthyology, 2014, 12, 565-584.	0.5	19
1926	HIV-1 pol Diversity among Female Bar and Hotel Workers in Northern Tanzania. PLoS ONE, 2014, 9, e102258.	1.1	5

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1927	Promiscuous Speciation with Gene Flow in Silverside Fish Genus <i>Odontesthes</i> (Atheriniformes). <i>Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 74</i>	1.1	18
1928	Comparative Analysis of the Mitochondrial Genomes of Callitettixini Spittlebugs (Hemiptera: Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 74) Presence of Short Conservative Elements at the Tribal Level. <i>PLoS ONE</i> , 2014, 9, e109140.	1.1	25
1929	Molecular and Morphological Analysis Reveals Five New Species of <i>Zygophiala</i> Associated with Flyspeck Signs on Plant Hosts from China. <i>PLoS ONE</i> , 2014, 9, e110717.	1.1	5
1930	<p><strong>Taxonomic assessment of Alligator Snapping Turtles </strong><strong>(Chelydridae: <em>Macrochelys</em>), with the description of </strong><strong>two new species from the southeastern United States</strong></p></strong>. <i>Zootaxa</i> , 2014, 3786, 141.	0.2	41
1931	DNA barcoding of Vietnamese bent-toed geckos (Squamata: Gekkonidae: <i>Cyrtodactylus</i> ) and the description of a new species. <i>Zootaxa</i> , 2014, 3784, 48-66.	0.2	35
1932	<strong>Systematics of the blindsnakes (Serpentes: Scolecophidia: Typhlopoidea) based on molecular and morphological evidence</strong>. <i>Zootaxa</i> , 2014, 3829, 1.	0.2	55
1933	A taxonomic revision of the Asian keelback snakes, genus <i>Amphiesma</i> (Serpentes: Colubridae: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 74)	0.2	42
1934	Antigenic characterisation of lyssaviruses in South Africa. <i>Onderstepoort Journal of Veterinary Research</i> , 2014, 81, .	0.6	8
1935	Molecular systematics of the world's most polytypic bird: the <em>Pachycephala pectoralis</em><em>melanura</em> (Aves: Pachycephalidae) species complex. <i>Zoological Journal of the Linnean Society</i> , 0, , .	1.0	0
1938	DNA barcoding for identification of the endangered species <i>Aquilaria sinensis</i>: comparison of data from heated or aged wood samples. <i>Holzforschung</i> , 2014, 68, 487-494.	0.9	62
1939	Evolutionary history of <i>Chaetognatha</i> inferred from molecular and morphological data: a case study for body plan simplification. <i>Frontiers in Zoology</i> , 2014, 11, 84.	0.9	20
1940	Systematics of the genus <i>Geastrum</i> (Fungi: Basidiomycota) revisited. <i>Taxon</i> , 2014, 63, 477-497.	0.4	34
1942	A new quadrannulate species of <i>Orobdella</i> (Hirudinida, Arhynchobdellida, Orobdellidae) from central Honshu, Japan. <i>ZooKeys</i> , 2014, 445, 57-76.	0.5	10
1943	Characterization of Corrosive Bacterial Consortia Isolated from Water in a Cooling Tower. <i>ISRN Corrosion</i> , 2014, 2014, 1-11.	0.3	22
1944	Evaluation of criteria for species delimitation of bagworm moths (Lepidoptera: Psychidae). <i>European Journal of Entomology</i> , 2014, 111, 121-136.	1.2	7
1945	Population Genetics of Two Key Mosquito Vectors of Rift Valley Fever Virus Reveals New Insights into the Changing Disease Outbreak Patterns in Kenya. <i>PLoS Neglected Tropical Diseases</i> , 2014, 8, e3364.	1.3	31
1946	Integrative taxonomy of New Caledonian beetles: species delimitation and definition of the <i>Uloma isoceroides</i> species group (Coleoptera, Tenebrionidae, Ulomini), with the description of four new species. <i>ZooKeys</i> , 2014, 415, 133-167.	0.5	9
1947	Phylogenetic Analyses. , 2014, , 93-110.		0

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1948	Comparative Dynamics and Distribution of Influenza Drug Resistance Acquisition to Protein M2 and Neuraminidase Inhibitors. <i>Molecular Biology and Evolution</i> , 2014, 31, 355-363.	3.5	31
1949	Phylogeny, biogeography and description of <i>Ameripassalus</i> , a new Mesoamerican genus of Passalidae (Coleoptera). <i>Invertebrate Systematics</i> , 2014, 28, 124.	0.5	17
1950	Multilocus Adaptation Associated with Heat Resistance in Reef-Building Corals. <i>Current Biology</i> , 2014, 24, 2952-2956.	1.8	216
1951	Mitochondrial DNA variability to explore the relationship complexity of Schizothoracine (Teleostei): Tj ETQq1 1 0.784314 rgBTj /Overlock 10 Tf 50 347 Td (<i>Schizothorax</i>). <i>Journal of Molecular Evolution</i> , 2014, 78, 1-14.	0.5	10
1952	Phylogenetic relationships of species of genus <i>Arachis</i> based on genic sequences. <i>Genome</i> , 2014, 57, 327-334.	0.9	5
1953	Naming and outline of Dothideomycetes 2014 including proposals for the protection or suppression of generic names. <i>Fungal Diversity</i> , 2014, 69, 1-55.	4.7	216
1954	<i>Fodinomyces uranophilus</i> gen. nov. sp. nov. and <i>Coniochaeta fodinicola</i> sp. nov., two uranium mine-inhabiting Ascomycota fungi from northern Australia. <i>Mycologia</i> , 2014, 106, 1073-1089.	0.8	43
1955	Inferring influenza global transmission networks without complete phylogenetic information. <i>Evolutionary Applications</i> , 2014, 7, 403-412.	1.5	4
1956	Predominance of a single phylogenetic species in colonization events among a sextet of decollate land snail, <i>Rumina decollata</i> (Mollusca: Pulmonata: Subulinidae), species. <i>Genome</i> , 2014, 57, 161-167.	0.9	7
1957	Metabolite of tryptophan promoting changes in EEG signal and the oxidative status of the brain. <i>Cell Biochemistry and Function</i> , 2014, 32, 496-501.	1.4	3
1958	Genetic persistence of an initially introduced brown trout ( <i>Salmo trutta</i> ) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 347 Td (<i>Salmo trutta</i>). <i>Journal of Molecular Evolution</i> , 2014, 78, 485-497.	0.7	10
1959	Removal of bisphenol A (BPA) in a nitrifying system with immobilized biomass. <i>Bioresource Technology</i> , 2014, 171, 305-313.	4.8	52
1960	A review of taxonomy and phylogenetic relationships in the genus <i>Costasiella</i> (Heterobranchia): Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 347 Td (<i>Costasiella</i>). <i>Journal of Molecular Evolution</i> , 2014, 78, 1-14.	0.4	11
1961	Towards a monophyletic classification of Lejeuneaceae II: subtribes Pycnolejeuneinae and Xylolejeuneinae subtr. nov., transfer of <i>Otolejeunea</i> to <i>Lepidolejeuninae</i> , and generic refinements. <i>Phytotaxa</i> , 2014, 163, 61.	0.1	24
1962	The <i>Colletotrichum destructivum</i> species complex - hemibiotrophic pathogens of forage and field crops. <i>Studies in Mycology</i> , 2014, 79, 49-84.	4.5	156
1963	<i>Brianaria</i> (Psoraceae), a new genus to accommodate the <i>Micareia sylvicola</i> group. <i>Lichenologist</i> , 2014, 46, 285-294.	0.5	17
1964	Electrochemistry of calcium precipitating bacteria in orthodontic wire. <i>Oral Science International</i> , 2014, 11, 22-29.	0.3	9
1965	New SSU-rDNA sequences for eleven colpodeans (Ciliophora, Colpodea) and description of <i>Apocrytolophosis</i> nov. gen. <i>European Journal of Protistology</i> , 2014, 50, 40-46.	0.5	14

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1966	Environmental siblings of black agents of human chromoblastomycosis. <i>Fungal Diversity</i> , 2014, 65, 47-63.	4.7	56
1967	Phylogenetic perspectives on biome shifts in <i>Lecanocoryne</i> ( <i>Ascomycota</i> ) in relation to climatic niche evolution in western South America. <i>Journal of Biogeography</i> , 2014, 41, 328-338.	1.4	33
1968	Diversification across the Palaearctic desert belt throughout the Pleistocene: phylogeographic history of the Houbara-Macqueen's bustard complex ( <i>Otididae</i> : <i>Chlamydotis</i> ) as revealed by mitochondrial DNA. <i>Journal of Zoological Systematics and Evolutionary Research</i> , 2014, 52, 65-74.	0.6	10
1969	Molecular phylogeny, biogeography, and host plant shifts in the bee genus <i>Melitta</i> (Hymenoptera: Tj ETQq1 1 0.784314 rgBT /Overlock	1.2	34
1970	Species limits and phylogenetic relationships of red-finned cryptic species of the seasonal killifish genus <i>Hypsolebias</i> from the Brazilian semi-arid Caatinga (Teleostei: Cyprinodontiformes: Tj ETQq0 0 0 rgBT /Overlock	1.0	15
1971	Phylogeny, character evolution, and biogeography of <i>Cuscuta</i> (dodders; Convolvulaceae) inferred from coding plastid and nuclear sequences. <i>American Journal of Botany</i> , 2014, 101, 670-690.	0.8	77
1972	Primate phylogenetic relationships and divergence dates inferred from complete mitochondrial genomes. <i>Molecular Phylogenetics and Evolution</i> , 2014, 75, 165-183.	1.2	260
1973	Reassessment of the systematics of the widespread Neotropical genus <i>Cercomacra</i> (Aves: Tj ETQq1 1 0.784314 rgBT /Overlock	1.0	19
1974	Divergent maximum-likelihood-branch-support values for polytomies. <i>Molecular Phylogenetics and Evolution</i> , 2014, 73, 87-96.	1.2	49
1975	Effects of an adaptive zone shift on morphological and ecological diversification in terapontid fishes. <i>Evolutionary Ecology</i> , 2014, 28, 205-227.	0.5	7
1976	The complex of <i>Diplodia</i> species associated with <i>Fraxinus</i> and some other woody hosts in Italy and Portugal. <i>Fungal Diversity</i> , 2014, 67, 143-156.	4.7	55
1977	Discovery of the Roosevelt's Barking Deer ( <i>Muntiacus rooseveltorum</i> ) in Vietnam. <i>Conservation Genetics</i> , 2014, 15, 993-999.	0.8	4
1978	Phylogenetic analysis of canine parvovirus partial VP2 gene in India. <i>Virus Genes</i> , 2014, 48, 89-95.	0.7	32
1979	Species of <i>Lasiodiplodia</i> associated with papaya stem-end rot in Brazil. <i>Fungal Diversity</i> , 2014, 67, 127-141.	4.7	86
1980	Evolution within the nuthatches ( <i>Sittidae</i> : Aves, Passeriformes): molecular phylogeny, biogeography, and ecological perspectives. <i>Journal of Ornithology</i> , 2014, 155, 755-765.	0.5	31
1981	Molecular data do not provide unambiguous support for the monophyly of flatfishes ( <i>Pleuronectiformes</i> ): A reply to Betancur-R and OrtÅ. <i>Molecular Phylogenetics and Evolution</i> , 2014, 75, 149-153.	1.2	25
1982	Molecular phylogeny of the New World gecko genus <i>Homonota</i> ( <i>Squamata</i> : <i>Phyllodactylidae</i> ). <i>Zoologica Scripta</i> , 2014, 43, 249-260.	0.7	24
1983	Genetic diversity assessed by SSR markers and chemotyping of <i>Fusarium culmorum</i> causal agent of foot and root rot of wheat collected from two different fields in Tunisia. <i>European Journal of Plant Pathology</i> , 2014, 139, 481-495.	0.8	10

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1984	Botryosphaeriaceae associated with diseases of mango ( <i>Mangifera indica</i> ). <i>Australasian Plant Pathology</i> , 2014, 43, 425.	0.5	18
1985	Molecular characterization of <i>Diplodia seriata</i> , a new pathogen of <i>Prunus laurocerasus</i> in Italy. <i>Phytoparasitica</i> , 2014, 42, 189-197.	0.6	11
1986	Phylogeography and biogeography of the lower Cenozoic North America: diversification between two continents and between two seas. <i>Biological Reviews</i> , 2014, 89, 767-790.	4.7	89
1987	A New Species of <i>Siphlophis</i> (Serpentes: Dipsadidae: Xenodontinae) from the Eastern Andean Slopes of Ecuador. <i>South American Journal of Herpetology</i> , 2014, 9, 30-45.	0.5	5
1988	A phylogeographic evaluation of the <i>Amolops mantzorum</i> species group: Cryptic species and plateau uplift. <i>Molecular Phylogenetics and Evolution</i> , 2014, 73, 40-52.	1.2	35
1989	Integrative taxonomy and conservation of seasonal killifishes, <i>Xenurolebias</i> (Teleostei: Tj ETQq1 1 0.784314 rgBT / Overlock 10 ff	0.5	10
1990	A phylogeny of softshell turtles (Testudines: Trionychidae) with reference to the taxonomic status of the critically endangered, giant softshell turtle, <i>Rafetus swinhoei</i> . <i>Organisms Diversity and Evolution</i> , 2014, 14, 279-293.	0.7	27
1991	<i>Geoglossum simile</i> of North America and Europe: distribution of a widespread earth tongue species and designation of an epitype. <i>Mycological Progress</i> , 2014, 13, 857-866.	0.5	5
1992	Molecular identification of <i>Anisakis</i> species (Nematoda: Anisakidae) from marine fishes collected in Turkish waters. <i>Veterinary Parasitology</i> , 2014, 201, 82-94.	0.7	53
1993	First molecular phylogeny of the subfamily Polycerinae (Mollusca, Nudibranchia, Polyceridae). <i>Helgoland Marine Research</i> , 2014, 68, 143-153.	1.3	16
1994	A molecular phylogeny of Pacific honeyeaters (Aves: Meliphagidae) reveals extensive paraphyly and an isolated Polynesian radiation. <i>Molecular Phylogenetics and Evolution</i> , 2014, 71, 308-315.	1.2	30
1995	Molecular systematics of the world's most polytypic bird: the <i>Pachycephala pectoralis</i> / <i>melanura</i> (Aves: Pachycephalidae) species complex. <i>Zoological Journal of the Linnean Society</i> , 2014, 170, 566-588.	1.0	48
1996	TBP-assisted species and hybrid identification in the genus <i>Passiflora</i> . <i>Molecular Breeding</i> , 2014, 33, 209-219.	1.0	12
1997	Multilocus phylogeny of talpine moles (Talpini, Talpidae, Eulipotyphla) and its implications for systematics. <i>Molecular Phylogenetics and Evolution</i> , 2014, 70, 513-521.	1.2	33
1998	Burst speciation processes and genomic expansion in the neotropical annual killifish genus <i>Austrolebias</i> (Cyprinodontiformes, Rivulidae). <i>Genetica</i> , 2014, 142, 87-98.	0.5	22
1999	Species Delimitation Using Bayes Factors: Simulations and Application to the <i>Sceloporus scalaris</i> Species Group (Squamata: Phrynosomatidae). <i>Systematic Biology</i> , 2014, 63, 119-133.	2.7	247
2000	Going further on an intricate and challenging group of nudibranchs: description of five novel species and a more complete molecular phylogeny of the subfamily <i>Nembrothinae</i> (Polyceridae). <i>Cladistics</i> , 2014, 30, 607-634.	1.5	14
2001	New generic proposal for the European Neogene large testudinids (Cryptodira) and the first phylogenetic hypothesis for the medium and large representatives of the European Cenozoic record. <i>Zoological Journal of the Linnean Society</i> , 2014, 172, 653-719.	1.0	21

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2002	Placing Environmental Next-Generation Sequencing Amplicons from Microbial Eukaryotes into a Phylogenetic Context. <i>Molecular Biology and Evolution</i> , 2014, 31, 993-1009.	3.5	97
2003	Introducing the Novel Species, <i>Dothiorella symphoricarposicola</i> , from Snowberry in Italy. <i>Cryptogamie, Mycologie</i> , 2014, 35, 257-270.	0.2	12
2004	Genetic and Morphological Approaches Distinguish the Three Sibling Species of the <i>Anisakis simplex</i> Species Complex, with a Species Designation as <i>Anisakis berlandi</i> n. sp. for <i>A. simplex</i> sp. C (Nematoda: Anisakidae). <i>Journal of Parasitology</i> , 2014, 100, 199-214.	0.3	167
2005	A New Host Record for <i>Tritetrabdella taiwana</i> (Hirudinida: Arhynchobdellida: Haemadipsidae) from the Asian Painted Frog <i>Kaloula pulchra</i> (Anura: Microhylidae) in Hong Kong, China, with a Taxonomic Note on <i>T. taiwana</i> . <i>Comparative Parasitology</i> , 2014, 81, 125-129.	0.0	5
2006	A new <i>Alternaria</i> species from grapevine in China. <i>Mycological Progress</i> , 2014, 13, 1119.	0.5	10
2008	Higher level molecular phylogeny of darkling beetles (Coleoptera: Tenebrionidae). <i>Journal of Molecular Evolution</i> , 2014, 78, 1-10.	1.7	542
2009	<i>Microbacterium endophyticum</i> sp. nov. and <i>Microbacterium halimionae</i> sp. nov., endophytes isolated from the salt-marsh plant <i>Halimione portulacoides</i> and emended description of the genus <i>Microbacterium</i> . <i>Systematic and Applied Microbiology</i> , 2014, 37, 474-479.	1.2	46
2010	Taxonomic review of <i>Scinax fuscomarginatus</i> (Lutz, 1925) and related species (Anura: Hylidae). <i>Zoological Journal of the Linnean Society</i> , 2014, 171, 783-821.	1.0	37
2011	Identification of bacteria associated with underground parts of <i>Crocus sativus</i> by 16S rRNA gene targeted metagenomic approach. <i>World Journal of Microbiology and Biotechnology</i> , 2014, 30, 2701-2709.	1.7	18
2012	Greensporones: Resorcylic Acid Lactones from an Aquatic <i>Halenospora</i> sp.. <i>Journal of Natural Products</i> , 2014, 77, 2088-2098.	1.5	69
2013	A multilocus phylogeny reveals deep lineages within African galagids (Primates: Galagidae). <i>BMC Evolutionary Biology</i> , 2014, 14, 72.	3.2	80
2014	Increase in transmitted resistance to non-nucleoside reverse transcriptase inhibitors among newly diagnosed HIV-1 infections in Europe. <i>BMC Infectious Diseases</i> , 2014, 14, 407.	1.3	43
2015	Combining morphological and phylogenetic analyses to unravel systematics in <i>Geastrum</i> sect. <i>Schmidelia</i> . <i>Mycologia</i> , 2014, 106, 1199-1211.	0.8	6
2016	Molecular phylogeny, species limits, and biogeography of the Brazilian endemic lizard genus <i>Enyalius</i> (Squamata: Leiosauridae): An example of the historical relationship between Atlantic Forests and Amazonia. <i>Molecular Phylogenetics and Evolution</i> , 2014, 81, 137-146.	1.2	42
2017	Increasing the number of discrete character states for continuous characters generates well-resolved trees that do not reflect phylogeny. <i>Integrative Zoology</i> , 2014, 9, 531-541.	1.3	12
2018	Phylogeny and classification of the Cuban species of <i>Elaphoglossum</i> (Dryopteridaceae), with description of <i>Elaphoglossum</i> sect. <i>Wrightiana</i> sect. nov.. <i>Plant Systematics and Evolution</i> , 2014, 300, 937-951.	0.3	25
2019	<i>Ilyonectria palmarum</i> sp. nov. causing dry basal stem rot of <i>Arecaceae</i> . <i>European Journal of Plant Pathology</i> , 2014, 138, 347-359.	0.8	19
2020	Molecular evidence on the occurrence of co-infection with <i>Pichia guilliermondii</i> and <i>Wuchereria bancrofti</i> in two filarial endemic districts of India. <i>Infectious Diseases of Poverty</i> , 2014, 3, 13.	1.5	17



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2021	Molecular phylogenetics of the <i>Handleyomys chapmani</i> complex in Mesoamerica. <i>Journal of Mammalogy</i> , 2014, 95, 26-40.	0.6	21
2022	<i>Myxobolus anaticus</i> sp. nov. (Myxozoa) infecting the gill of Anatolian khramulya <i>Capoeta tinca</i> (Cyprinidae) in Turkey. <i>Diseases of Aquatic Organisms</i> , 2014, 109, 213-222.	0.5	6
2023	Three new species of Hyphodontia with peg-like hyphal aggregations. <i>Mycological Progress</i> , 2014, 13, 533-545.	0.5	27
2024	Understanding Sequence Similarity and Framework Analysis Between Centromere Proteins Using Computational Biology. <i>Cell Biochemistry and Biophysics</i> , 2014, 70, 897-906.	0.9	0
2025	Assessment of Genetic Diversity Among Indian Jujube Varieties Based on Nuclear Ribosomal DNA and RAPD Polymorphism. <i>Agricultural Research</i> , 2014, 3, 218-228.	0.9	6
2026	Revision of <i>Entyloma</i> (Entylomatales, Exobasidiomycetes) on <i>Eryngium</i> . <i>Mycologia</i> , 2014, 106, 797-810.	0.8	18
2027	Molecular identification, morphological characterization and new insights into the ecology of larval <i>Pseudoterranova cattani</i> in fishes from the Argentine coast with its differentiation from the Antarctic species, <i>P. decipiens</i> sp. E (Nematoda: Anisakidae). <i>Veterinary Parasitology</i> , 2014, 199, 59-72.	0.7	44
2028	Phylogeny of the family Aglajidae (Pilsbry, 1895) (Heterobranchia: Cephalaspidea) inferred from mtDNA and nDNA. <i>Molecular Phylogenetics and Evolution</i> , 2014, 71, 113-126.	1.2	27
2029	High genetic diversity in the hydroid <i>Plumularia setacea</i> : A multitude of cryptic species or extensive population subdivision?. <i>Molecular Phylogenetics and Evolution</i> , 2014, 76, 1-9.	1.2	67
2030	Phylogenetics of the gastropod genus <i>Nucella</i> (Neogastropoda: Muricidae): species identities, timing of diversification and correlated patterns of life-history evolution. <i>Journal of Molluscan Studies</i> , 2014, 80, 341-353.	0.4	18
2031	The Phylogenetic Placement of <i>Eriosporella bambusicola</i> sp. nov. in <i>Capnodiales</i> . <i>Cryptogamie, Mycologie</i> , 2014, 35, 41-49.	0.2	11
2032	Phylogenetic Relationships among Dracaenoid Genera (Asparagaceae: Nolinoideae) Inferred from Chloroplast DNA Loci. <i>Systematic Botany</i> , 2014, 39, 90-104.	0.2	55
2033	Do freshwater ecoregions and continental shelf width predict patterns of historical gene flow in the freshwater fish <i>Poecilia butleri</i> ?. <i>Biological Journal of the Linnean Society</i> , 2014, 112, 399-416.	0.7	11
2034	Higher-order accuracy of multiscale-double bootstrap for testing regions. <i>Journal of Multivariate Analysis</i> , 2014, 130, 208-223.	0.5	1
2035	Morphologic and molecular description of <i>Metopus fuscus</i> Kahl from North America and new rDNA sequences from seven metopids (Armophorea, Metopidae). <i>European Journal of Protistology</i> , 2014, 50, 213-230.	0.5	28
2036	Molecular phylogeny of the genus <i>Sticta</i> (lichenized Ascomycota: Lobariaceae) in Colombia. <i>Fungal Diversity</i> , 2014, 64, 205-231.	4.7	62
2037	Developed of a method for the genetic identification of ling species ( <i>Genypterus</i> spp.) in seafood products by FINS methodology. <i>Food Chemistry</i> , 2014, 143, 22-26.	4.2	18
2038	The Bromeliaceae tank dweller <i>Bromeliophila</i> (Lejeuneaceae, Porellales) is a member of the <i>Cyclolejeunea-Prionolejeunea</i> clade. <i>Plant Systematics and Evolution</i> , 2014, 300, 63-73.	0.3	14

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2039	Towards a monophyletic classification of Lejeuneaceae I: subtribe Leptolejeuneinae subtr. nov.. Phytotaxa, 2014, 156, 165.	0.1	19
2040	Genetically identical co-housed pigs as models for dietary studies of gut microbiomes. Microbiome Science and Medicine, 2014, 1, .	0.3	3
2041	Integrative taxonomy of Acrapex stem borers (Lepidoptera : Noctuidae : Apameini): combining morphology and Poisson Tree Process analyses. Invertebrate Systematics, 2014, 28, 451.	0.5	20
2042	<span lang="EN-GB">Towards a monophyletic classification of Lejeuneaceae III: the systematic position of Leiolejeunea</span>. Phytotaxa, 2014, 170, 187.	0.1	25
2043	New generic proposal for the European Neogene large testudinids (Cryptodira) and the first phylogenetic hypothesis for the medium and large representatives of the European Cenozoic record. Zoological Journal of the Linnean Society, 2014, 172, 653-719.	1.0	14
2044	Phylogeny and taxonomy of the genus Glioccephalotrichum. Persoonia: Molecular Phylogeny and Evolution of Fungi, 2014, 32, 127-140.	1.6	8
2045	New Phaeoacremonium species isolated from sandalwood trees in Western Australia. IMA Fungus, 2014, 5, 67-77.	1.7	22
2046	Spatiotemporal evolution of Reaumuria (Tamaricaceae) in Central Asia: insights from molecular biogeography. Phytotaxa, 2014, 167, 89.	0.1	13
2047	<p class="p0">A new species of Microthyrium from Yunnan, China. Phytotaxa, 2014, 176, 213.	0.1	11
2048	<i>Tremella cetrariellae</i> (<i>Tremellales</i>, Basidiomycota, Fungi), a new lichenicolous fungus on <i>Cetrariella delisei</i>. Lichenologist, 2015, 47, 359-368.	0.5	15
2049	Phylogeny of tremellomycetous yeasts and related dimorphic and filamentous basidiomycetes reconstructed from multiple gene sequence analyses. Studies in Mycology, 2015, 81, 1-26.	4.5	133
2050	New record of <i>Aplosporella javeedii</i> on five hosts in China based on multi-gene analysis and morphology. Mycotaxon, 2015, 130, 749-756.	0.1	12
2051	Mutinus albotruncatus (Phallales, Agaricomycetes), a new phalloid from the Brazilian semiarid, and a key to the world species. Phytotaxa, 2015, 236, 237.	0.1	7
2052	Multigene phylogeny and morphology reveal Phaeobotryon rhois sp. nov. (Botryosphaerales,) Tj ETQq1 1 0.784314 rgBT /Overlock 10 T	0.1	16
2053	Characterization of Botryosphaeriaceae Species as Causal Agents of Trunk Diseases on Grapevines. Plant Disease, 2015, 99, 1678-1688.	0.7	32
2054	Origins of the Endemic Scaly Tree Ferns on the GalÃ¡pagos and Cocos Islands. International Journal of Plant Sciences, 2015, 176, 869-879.	0.6	6
2055	Molecular Phylogenetics: A Tool for Elucidation of Evolutionary Processes from Biological Data. , 2015, , 203-250.		0
2056	A new predatory leech from Vietnam (Hirudinida : Arhynchobdellida : Salifidae): its phylogenetic position with comments on the classification of the family. Invertebrate Systematics, 2015, 29, 473.	0.5	4

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2057	<i>Sphaerolejeunea</i> (Lejeuneaceae), Tj ETQq0 0 0, rgBT /Overlock 10 Tf 50 147	0.1	19
2058	A new species of <i>Scleroramularia</i> associated with sooty blotch and flyspeck in Southern China. <i>Phytotaxa</i> , 2015, 226, 261.	0.1	3
2059	<i>Clathrosporium retortum</i> , a novel aeroaquatic fungus in the Sordariomycetidae (Ascomycota) from Brazil. <i>Phytotaxa</i> , 2015, 239, 17.	0.1	6
2060	Molecular phylogenetic analysis reveals two new species of <i>Discosia</i> from Italy. <i>Phytotaxa</i> , 2015, 203, 37.	0.1	5
2061	Evidence of bacterioplankton community adaptation in response to long-term mariculture disturbance. <i>Scientific Reports</i> , 2015, 5, 15274.	1.6	45
2062	Phylogenetic analysis of HIV sub-epidemics in Mochudi, Botswana. <i>Epidemics</i> , 2015, 13, 44-55.	1.5	22
2063	Integrative taxonomy reveals an unexpected diversity in <i>Geastrum</i> section <i>Geastrum</i> ( <i>Geastrales</i> , <i>Basidiomycota</i> ). <i>Persoonia: Molecular Phylogeny and Evolution of Fungi</i> , 2015, 34, 130-165.	1.6	31
2064	Elucidating the <i>Ramularia eucalypti</i> species complex. <i>Persoonia: Molecular Phylogeny and Evolution of Fungi</i> , 2015, 34, 50-64.	1.6	27
2065	An integrative taxonomic approach to assess the status of <i>Corsican</i> bumblebees: implications for conservation. <i>Animal Conservation</i> , 2015, 18, 236-248.	1.5	42
2066	Seeing double: pseudocryptic diversity in the <i>Doriopsilla albopunctata</i> – <i>Doriopsilla gemela</i> species complex of the north-eastern Pacific. <i>Zoologica Scripta</i> , 2015, 44, 612-631.	0.7	19
2067	Biogeography of Australasian flightless weevils ( <i>Curculionidae</i> , <i>Celeuthetini</i> ) suggests permeability of Lydekker's and Wallace's Lines. <i>Zoologica Scripta</i> , 2015, 44, 632-644.	0.7	27
2068	Expansion of a novel endogenous retrovirus throughout the pericentromeres of modern humans. <i>Genome Biology</i> , 2015, 16, 74.	3.8	34
2069	Highly polytypic taxon complex: interspecific and intraspecific integrative taxonomic assessment of the widespread pollinator <i>Bombus pascuorum</i> ( <i>Synaldis</i> copoli 1763 ( <i>Hymenoptera</i> : <i>Apididae</i> ). <i>Systematic Entomology</i> , 2015, 40, 881-890.	1.7	19
2070	DNA Barcode Development for Three Recent Exotic Whitefly ( <i>Hemiptera</i> : <i>Aleyrodidae</i> ) Invaders in Florida. <i>Florida Entomologist</i> , 2015, 98, 473-478.	0.2	16
2071	Integrative taxonomy reveals six new species related to the Mediterranean corn stalk borer <i>Sesamia nonagrioides</i> (Lefebvre) ( <i>Lepidoptera</i> , <i>Noctuidae</i> , <i>Sesamiina</i> ). <i>Zoological Journal of the Linnean Society</i> , 2015, 175, 244-270.	1.0	28
2072	A new genus and species of xenodermatid snake ( <i>Squamata</i> ) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 147	0.2	19
2073	Identification of Eastern United States <i>Reticulitermes</i> Termite Species via PCR-RFLP, Assessed Using Training and Test Data. <i>Insects</i> , 2015, 6, 524-537.	1.0	11
2074	The Complete Mitochondrial Genome of the Beet Webworm, <i>Spoladea recurvalis</i> ( <i>Lepidoptera</i> : Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 147	1.1	17

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2075	Three New Species of Cyphellophora (Chaetothyriales) Associated with Sooty Blotch and Flyspeck. PLoS ONE, 2015, 10, e0136857.	1.1	29
2076	Molecular Characterization of Human Respiratory Syncytial Virus in the Philippines, 2012-2013. PLoS ONE, 2015, 10, e0142192.	1.1	55
2077	The Historical Speciation of Mauremys Sensu Lato: Ancestral Area Reconstruction and Interspecific Gene Flow Level Assessment Provide New Insights. PLoS ONE, 2015, 10, e0144711.	1.1	11
2078	Inter-Seasonal Influenza is Characterized by Extended Virus Transmission and Persistence. PLoS Pathogens, 2015, 11, e1004991.	2.1	25
2079	<strong>Molecular phylogeny of Caribbean dipsadid (Xenodontinae: Alsophiini) snakes, including identification of the first record from the Cay Sal Bank, The Bahamas</strong>. Zootaxa, 2015, 4028, 441.	0.2	4
2080	<p><strong>A second New World hoverfly, <em>Toxomerus</em> <em>floralis</em> (Fabricius) (Diptera: Syrphidae), recorded from the Old World, with description of larval pollen-feeding ecology</strong></p>.	0.2	10
2081	<p><strong>Molecular phylogenetics, systematics and host-plant associations of the <em>Bruchidius</em> <em>albosparsus</em> (Fåhræus) species group (Coleoptera, Chrysomelidae, Bruchinae) with the description of four new species</strong></p>.	0.2	5
2082	A revision of the genus Conicofrontia Hampson (Lepidoptera, Noctuidae,  Apameini,) Tj ETQq1 1 0.784314 rgBT /Overlock 10 molecular data. Zootaxa, 2015, 3925, 56-74.	0.2	8
2083	Phylogenetic incongruence inferred with two mitochondrial genes in Mepraia spp. and Triatoma eratyrisiformis (Hemiptera, Reduviidae). Genetics and Molecular Biology, 2015, 38, 390-395.	0.6	11
2084	Genetic Diversity and Conservation of an Endemic Taiwanese Species, Platyeriocheir formosa. , 0, , .		0
2085	Phylogeographic Structure in Anastrepha ludens (Diptera: Tephritidae) Populations Inferred With mtDNA Sequencing. Journal of Economic Entomology, 2015, 108, 1324-1336.	0.8	21
2086	Molecular Phylogenetic and Phylogenomic Approaches in Studies of Lichen Systematics and Evolution. , 2015, , 45-60.		7
2087	Morphological and molecular characterisation of Diaporthe species associated with grapevine trunk disease in China. Fungal Biology, 2015, 119, 283-294.	1.1	62
2088	Pseudo-cryptic speciation in the subterranean medium: A new species of Stylodrilus ClaparÃ“de, 1862, with a revision of the status of Bichaeta Bretscher, 1900 (Annelida, Clitellata, Lumbriculidae). Zoologischer Anzeiger, 2015, 257, 71-86.	0.4	2
2089	Phylogenies from concatenated data: Is the end nigh?. Taxon, 2015, 64, 421-423.	0.4	41
2090	Comparative molecular species delimitation in the charismatic Nawab butterflies (Nymphalidae,) Tj ETQq1 1 0.784314 rgBT /Overlock 10 1.2 22		
2091	Multilocus sequence data reveal dozens of putative cryptic species in a radiation of endemic Californian mygalomorph spiders (Araneae, Mygalomorphae, Nemesiidae). Molecular Phylogenetics and Evolution, 2015, 91, 56-67.	1.2	58
2092	Gut-Associated Bacteria of Dendroctonus valens and their Involvement in Verbenone Production. Microbial Ecology, 2015, 70, 1012-1023.	1.4	91

#	ARTICLE	IF	CITATIONS
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2094	Transcriptomic characterization of the enzymatic antioxidants FeSOD, MnSOD, APX and KatG in the dinoflagellate genus Symbiodinium. BMC Evolutionary Biology, 2015, 15, 48.	3.2	50
2095	Estatus Taxonómico de Caryedon serratus (Olivier) en México. Southwestern Entomologist, 2015, 40, 387-396.	0.1	1
2096	Tracing the evolutionary history of the little-known Mediterranean-Macaronesian genus <i>Andryala</i> (Asteraceae) by multigene sequencing. Taxon, 2015, 64, 535-551.	0.4	11
2097	Burnaia Miller, 2001 (Gastropoda, Heterobranchia, Nudibranchia): a facelinid genus with an Aeolidiidae's outward appearance. Helgoland Marine Research, 2015, 69, 285-291.	1.3	7
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2099	Phylogeography and morphological variation of the northernmost distributed species of the Liolaemus lineomaculatus section (Liolaemini) from Patagonia. Amphibia - Reptilia, 2015, 36, 373-387.	0.1	9
2100	Genetic diversity within ITS-1 region of Eimeria species infecting chickens of north India. Infection, Genetics and Evolution, 2015, 36, 262-267.	1.0	14
2101	Cytospora from Ulmus pumila in Northern China. Mycological Progress, 2015, 14, 1.	0.5	22
2102	Non-invasive fast real-time PCR assay for detection of the enteric parasite Enteromyxum scophthalmi in cultured turbot (Scophthalmus maximus L.). Aquaculture Research, 2015, 46, 2104-2115.	0.9	3
2103	Identification and characterization of Pestalotiopsis-like fungi related to grapevine diseases in China. Fungal Biology, 2015, 119, 348-361.	1.1	43
2104	Phenotypic and molecular diversity of Meyerozyma guilliermondii strains isolated from food and other environmental niches, hints for an incipient speciation. Food Microbiology, 2015, 48, 206-215.	2.1	41
2105	Diversity and potential impact of <i>Calonectria</i> species in <i>Eucalyptus</i> plantations in Brazil. Studies in Mycology, 2015, 80, 89-130.	4.5	60
2106	Molecular phylogenetics and species delimitation of leaf-toed geckos (Phyllodactylidae: Tj ETQq1 1.0.784314 rgBT / Overlock 10 Tf 50.2	1.2	25
2107	New species, hyper-diversity and potential importance of <i>Calonectria</i> spp. from <i>Eucalyptus</i> in South China. Studies in Mycology, 2015, 80, 151-188.	4.5	56
2108	Taking the discovery approach in integrative taxonomy: decrypting a complex of narrow-endemic Alpine harvestmen (Opiliones: Phalangidae: <i>Megabunus</i> ). Molecular Ecology, 2015, 24, 863-889.	2.0	19
2109	Montane and coastal species diversification in the economically important Mexican grasshopper genus Sphenarium (Orthoptera: Pyrgomorphidae). Molecular Phylogenetics and Evolution, 2015, 84, 220-231.	1.2	13
2110	Insect midgut carboxypeptidases with emphasis on <i>S</i> hemipteran and <i>M</i> lepidopteran carboxypeptidases. Insect Molecular Biology, 2015, 24, 222-239.	1.0	19

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2112	Genetic identification and distribution of the parasitic larvae of <i>Anisakis pegreffii</i> and <i>Anisakis simplex</i> (s. s.) in European hake <i>Merluccius merluccius</i> from the Tyrrhenian Sea and Spanish Atlantic coast: Implications for food safety. <i>International Journal of Food Microbiology</i> , 2015, 198, 1-8.	2.1	60
2113	Coevolution between <i>Contraecaecum</i> (Nematoda, Anisakidae) and <i>Austrolebias</i> (Cyprinodontiformes). <i>Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50</i> 913-927.	0.6	3
2114	<i>Protaeolidiella atra</i> Baba, 1955 versus <i>Pleurolidia juliae</i> Burn, 1966: One or two species?. <i>Helgoland Marine Research</i> , 2015, 69, 137-145.	1.3	6
2115	Methods for species delimitation in bumblebees (<sc>H</sc>ymentoptera, <sc>A</sc>pidae). <i>Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50</i>	0.7	51
2116	A mitochondrial genome phylogeny of owl moths (Lepidoptera: Noctuoidea), and examination of the utility of mitochondrial genomes for lepidopteran phylogenetics. <i>Molecular Phylogenetics and Evolution</i> , 2015, 85, 230-237.	1.2	76
2117	Functional Screening and Molecular Characterization of Halophilic and Halotolerant Bacteria by 16S rRNA Gene Sequence Analysis. <i>Proceedings of the National Academy of Sciences India Section B - Biological Sciences</i> , 2015, 85, 957-964.	0.4	6
2118	Phylogeography of the smooth snake <i>Coronella austriaca</i> (Serpentes: Colubridae): evidence for a reduced gene pool and a genetic discontinuity in Central Europe. <i>Biological Journal of the Linnean Society</i> , 2015, 115, 195-210.	0.7	10
2119	<i>Curviclavula</i> , a new genus of anamorphic Helotiales (Leotiomycetes) isolated from air. <i>Mycological Progress</i> , 2015, 14, 1.	0.5	3
2120	Multilocus sequence typing of a dairy-associated <i>Leuconostoc mesenteroides</i> population reveals clonal structure with intragenic homologous recombination. <i>Journal of Dairy Science</i> , 2015, 98, 2284-2293.	1.4	11
2121	<i>Cytospora</i> species associated with walnut canker disease in China, with description of a new species <i>C. gígalocus</i> . <i>Fungal Biology</i> , 2015, 119, 310-319.	1.1	56
2122	<i>Microbacterium proteolyticum</i> sp. nov. isolated from roots of <i>Halimione portulacoides</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2015, 65, 1794-1798.	0.8	16
2123	Next-generation sequencing as means to retrieve tick systematic markers, with the focus on <i>Nuttalliella namaqua</i> (Ixodoidea: Nuttalliellidae). <i>Ticks and Tick-borne Diseases</i> , 2015, 6, 450-462.	1.1	19
2124	Reconciling morphological and molecular classification of predatory ciliates: Evolutionary taxonomy of dileptids (Ciliophora, Litostomatea, Rhynchostomatia). <i>Molecular Phylogenetics and Evolution</i> , 2015, 90, 112-128.	1.2	22
2125	Phylogenetic Diversity of Sponge-Associated Fungi from the Caribbean and the Pacific of Panama and Their In Vitro Effect on Angiotensin and Endothelin Receptors. <i>Marine Biotechnology</i> , 2015, 17, 533-564.	1.1	19
2126	A multilocus molecular phylogeny of the endemic North American camel spider family Eremobatidae (Arachnida: Solifugae). <i>Molecular Phylogenetics and Evolution</i> , 2015, 92, 280-293.	1.2	18
2127	Molecular evaluation of the phylogenetic position of the enigmatic species <i>Trivettea papalotla</i> (Bertsch) (Mollusca : Nudibranchia). <i>Invertebrate Systematics</i> , 2015, 29, 215.	0.5	10
2128	Phylogeny, systematics, and trait evolution of <i>Carex</i> section <i>Glareosae</i> . <i>American Journal of Botany</i> , 2015, 102, 1128-1144.	0.8	19

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2129	Insect midgut Î±-mannosidases from family 38 and 47 with emphasis on those of <i>Tenebrio molitor</i> . <i>Insect Biochemistry and Molecular Biology</i> , 2015, 67, 94-104.	1.2	9
2130	<i>Colletotrichum fructicola</i> is the dominant and one of the most aggressive species causing bitter rot of apple in Uruguay. <i>Tropical Plant Pathology</i> , 2015, 40, 265-274.	0.8	25
2131	Species Selection Favors Dispersive Life Histories in Sea Slugs, but Higher Per-Offspring Investment Drives Shifts to Short-Lived Larvae. <i>Systematic Biology</i> , 2015, 64, 983-999.	2.7	44
2132	<i>Maasoglossum</i> , a basal genus in Geoglossomycetes. <i>Mycoscience</i> , 2015, 56, 572-579.	0.3	4
2133	Evolution of Spermophagus seed beetles (Coleoptera, Bruchinae, Amblycerini) indicates both synchronous and delayed colonizations of host plants. <i>Molecular Phylogenetics and Evolution</i> , 2015, 89, 91-103.	1.2	14
2134	Cultural macroevolution among high latitude hunter-gatherers: a phylogenetic study of the Arctic Small Tool tradition. <i>Journal of Archaeological Science</i> , 2015, 59, 64-79.	1.2	27
2135	Molecular data support <i>Pseudoparmelia</i> as a distinct lineage related to <i>Relicina</i> and <i>Relicinopsis</i> (Ascomycota, Lecanorales). <i>Lichenologist</i> , 2015, 47, 43-49.	0.5	10
2136	Molecular phylogenetic and biogeographical analysis of <i>Nitraria</i> based on nuclear and chloroplast DNA sequences. <i>Plant Systematics and Evolution</i> , 2015, 301, 1897-1906.	0.3	23
2137	Phylogeny of the Acarosporaceae (Lecanoromycetes, Ascomycota, Fungi) and the evolution of carbonized ascomata. <i>Fungal Diversity</i> , 2015, 73, 145-158.	4.7	44
2138	Molecular Characterization of Commercial Varieties of <i>Phyllanthus emblica</i> Using RAPD and Nuclear rDNA SNPs. <i>Proceedings of the National Academy of Sciences India Section B - Biological Sciences</i> , 2015, 85, 971-978.	0.4	3
2139	Phylogeny of <i>Salix</i> subgenus <i>Salix</i> s.l. (Salicaceae): delimitation, biogeography, and reticulate evolution. <i>BMC Evolutionary Biology</i> , 2015, 15, 31.	3.2	77
2140	Comparative genetic diversity of Lyme disease bacteria in Northern Californian ticks and their vertebrate hosts. <i>Ticks and Tick-borne Diseases</i> , 2015, 6, 414-423.	1.1	15
2141	Phylogenetic Molecular Species Delimitations Unravel Potential New Species in the Pest Genus <i>Spodoptera</i> Guenée, 1852 (Lepidoptera, Noctuidae). <i>PLoS ONE</i> , 2015, 10, e0122407.	1.1	67
2142	Molecular investigation of the phylogenetic position of the polar nudibranch <i>Doridoxa</i> (Mollusca, Tj ETQq1 1 0.784314 rgBT / Overlook 0,5 20	0.5	20
2143	Human Endogenous Retrovirus Type K (HERV-K) Particles Package and Transmit HERV-Related Sequences. <i>Journal of Virology</i> , 2015, 89, 7187-7201.	1.5	43
2144	Caulicolous <i>Botryosphaerales</i> from Thailand. <i>Persoonia: Molecular Phylogeny and Evolution of Fungi</i> , 2015, 34, 87-99.	1.6	53
2145	Carpogenic germination of sclerotia of <i>Sclerotinia minor</i> and ascospore infection of pyrethrum flowers. <i>Canadian Journal of Plant Pathology</i> , 2015, 37, 179-187.	0.8	1
2146	Estimating the temporal and spatial extent of gene flow among sympatric lizard populations (genus) Tj ETQq1 1 0.784314 rgBT / Overlook 2,0 16	2.0	16

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2147	Phylogenetic and chemical diversity of fungal endophytes isolated from <i>Silybum marianum</i> (L) Gaertn. (milk thistle). <i>Mycology</i> , 2015, 6, 8-27.	2.0	29
2148	Diverse species of <i>Colletotrichum</i> associated with grapevine anthracnose in China. <i>Fungal Diversity</i> , 2015, 71, 233-246.	4.7	64
2149	<i>Amycolatopsis</i> sp. Poz14 isolated from oil-contaminated soil degrades polycyclic aromatic hydrocarbons. <i>International Biodeterioration and Biodegradation</i> , 2015, 99, 165-173.	1.9	33
2150	Dissecting signal and noise in diatom chloroplast protein encoding genes with phylogenetic information profiling. <i>Molecular Phylogenetics and Evolution</i> , 2015, 89, 28-36.	1.2	81
2151	Long forsaken species diversity in the Middle American lizard <i>Holcosus undulatus</i> (Teiidae). <i>Zoological Journal of the Linnean Society</i> , 2015, 175, 189-210.	1.0	14
2152	DNA Markers in Diversity Analysis. , 2015, , 23-46.		3
2153	<i>Tuber turmericum</i> sp. nov., a Chinese truffle species based on morphological and molecular data. <i>Mycological Progress</i> , 2015, 14, 1.	0.5	10
2154	Integrative taxonomy of <i>Lepidolejeunea</i> (Jungermanniopsida: Porellales): Ocelli allow the recognition of two neglected species. <i>Taxon</i> , 2015, 64, 216-228.	0.4	40
2155	Biogeography of the <i>Phalaenopsis amabilis</i> species complex inferred from nuclear and plastid DNAs. <i>BMC Plant Biology</i> , 2015, 15, 202.	1.6	13
2156	Root of <i>Dictyostelia</i> based on 213 universal proteins. <i>Molecular Phylogenetics and Evolution</i> , 2015, 92, 53-62.	1.2	16
2157	Morphological and molecular characterisation of <i>Geosmithia</i> species on European elms. <i>Fungal Biology</i> , 2015, 119, 1063-1074.	1.1	17
2158	Morphology-based phylogenetic binning to assess a taxonomic challenge: a case study in Graphidaceae (Ascomycota) requires a new generic name for the widespread <i>Lepidotrema wightii</i> . <i>Botanical Journal of the Linnean Society</i> , 2015, 179, 436-443.	0.8	11
2159	Towards a comprehensive phylogeny of the large temperate genus <i>Pedicularis</i> (Orobanchaceae), with an emphasis on species from the Himalaya-Hengduan Mountains. <i>BMC Plant Biology</i> , 2015, 15, 176.	1.6	51
2160	<i>Fonsecaea pugnacius</i> , a Novel Agent of Disseminated Chromoblastomycosis. <i>Journal of Clinical Microbiology</i> , 2015, 53, 2674-2685.	1.8	62
2161	The shrimp mitochondrial FoF1-ATPase inhibitory factor 1 (IF1). <i>Journal of Bioenergetics and Biomembranes</i> , 2015, 47, 383-393.	1.0	5
2162	Morphology, ontogenesis and molecular characterization of <i>Atractos contortus</i> V&agrave;r&agrave;sv&agrave;ry, 1950 and <i>Stichotricha aculeata</i> Wrzesniowski&egrave;, 1866 (Ciliophora, Stichotrichida) with consideration of their systematic positions. <i>European Journal of Protistology</i> , 2015, 51, 351-373.	0.5	14
2163	Mitochondrial DNA diversity, genetic structure, and demographic history of the Neotropical otter ( <i>Lontra longicaudis</i> ) in Mexico. <i>Journal of Mammalogy</i> , 2015, 96, 1162-1173.	0.6	10
2164	Phylogeny and Character Evolution of Papaveraceae s. l. (Ranunculales). <i>Systematic Botany</i> , 2015, 40, 474-488.	0.2	27



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2165	First detection of bovine viral diarrhoea virus type 2 in cattle in Spain. <i>Veterinary Record Open</i> , 2015, 2, e000110.	0.3	12
2166	MtDNA Analysis for Genetic Identification of Forensically Important Sarcophagid Flies (Diptera: Tj ETQq1 1 0.784314 rgBT /Overlock 10	0.9	12
2167	<i>Diaporthe rostrata</i> , a novel ascomycete from <i>Juglans mandshurica</i> associated with walnut dieback. <i>Mycological Progress</i> , 2015, 14, 1.	0.5	26
2168	Minutisphaerales (Dothideomycetes, Ascomycota): a new order of freshwater ascomycetes including a new family, Minutisphaeraceae, and two new species from North Carolina, USA. <i>Mycologia</i> , 2015, 107, 845-862.	0.8	26
2169	Morphological Variation, Niche Divergence, and Phylogeography of Lizards of the <i>Liolaemus lineomaculatus</i> Section (Liolaemini) from Southern Patagonia. <i>Herpetological Monographs</i> , 2015, 29, 65.	1.1	18
2170	The rise of <i>Ramularia</i> from the <i>Mycosphaerella labyrinth</i> . <i>Fungal Biology</i> , 2015, 119, 823-843.	1.1	32
2171	Taxonomy, phylogenetics and biogeography of <i>Chesneya</i> (Fabaceae), evidenced from data of three sequences, ITS, trnS-trnG, and rbcL. <i>Biochemical Systematics and Ecology</i> , 2015, 63, 80-89.	0.6	12
2173	Ancestral reconstruction of reproductive traits shows no tendency toward terrestriality in leptodactyline frogs. <i>BMC Evolutionary Biology</i> , 2015, 15, 91.	3.2	18
2174	Fragiliporiaceae, a new family of Polyporales (Basidiomycota). <i>Fungal Diversity</i> , 2015, 70, 115-126.	4.7	53
2175	Exploring the Genomic Roadmap and Molecular Phylogenetics Associated with MODY Cascades Using Computational Biology. <i>Cell Biochemistry and Biophysics</i> , 2015, 71, 1491-1502.	0.9	2
2177	A new European Late Jurassic pleurosternid (Testudines, Paracryptodira) and a new hypothesis of paracryptodiran phylogeny. <i>Journal of Systematic Palaeontology</i> , 2015, 13, 351-369.	0.6	28
2178	Internal larval characters in anuran systematic studies: a phylogenetic hypothesis for <i>Leptodactylus</i> (Anura, Leptodactylidae). <i>Journal of Zoological Systematics and Evolutionary Research</i> , 2015, 53, 55-66.	0.6	6
2179	Phylogeny of the monarch flycatchers reveals extensive paraphyly and novel relationships within a major Australo-Pacific radiation. <i>Molecular Phylogenetics and Evolution</i> , 2015, 83, 118-136.	1.2	28
2180	Role of Caribbean Islands in the diversification and biogeography of Neotropical <i>Heraclides</i> swallowtails. <i>Cladistics</i> , 2015, 31, 291-314.	1.5	30
2181	Isolation and Characterization of Arsenic-Resistant Bacteria from Contaminated Water-Bodies in West Bengal, India. <i>Geomicrobiology Journal</i> , 2015, 32, 17-26.	1.0	21
2182	Genetic system for traceability of goatfishes by FINS methodology and authentication of mullets ( <i>Mullus barbatus</i> and <i>Mullus surmuletus</i> ) by RT-PCR. <i>European Food Research and Technology</i> , 2015, 240, 423-429.	1.6	4
2183	Coprophilous contributions to the phylogeny of Lasiosphaeriaceae and allied taxa within Sordariales (Ascomycota, Fungi). <i>Fungal Diversity</i> , 2015, 70, 101-113.	4.7	41
2184	Sharpening the species boundaries in the <i>Cladonia mediterranea</i> complex (Cladoniaceae, Ascomycota). <i>Persoonia: Molecular Phylogeny and Evolution of Fungi</i> , 2016, 37, 1-12.	1.6	8

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2185	Finding the missing link: Resolving the Coryneliomycetidae within Eurotiomycetes. <i>Persoonia: Molecular Phylogeny and Evolution of Fungi</i> , 2016, 37, 37-56.	1.6	16
2186	Novel point mutations in the ERG11 gene in clinical isolates of azole resistant <i>Candida</i> species. <i>Memorias Do Instituto Oswaldo Cruz</i> , 2016, 111, 192-199.	0.8	31
2187	Tandem Duplication Events in the Expansion of the Small Heat Shock Protein Gene Family in <i>Solanum lycopersicum</i> (cv. Heinz 1706). <i>G3: Genes, Genomes, Genetics</i> , 2016, 6, 3027-3034.	0.8	13
2188	Analysis of the mitochondrial COI gene and its informative potential for evolutionary inferences in the families Coreidae and Pentatomidae (Heteroptera). <i>Genetics and Molecular Research</i> , 2016, 15, .	0.3	7
2189	After the Ice Age: The Impact of Post-Glacial Dispersal on the Phylogeography of a Small Mammal, <i>Muscardinus avellanarius</i> . <i>Frontiers in Ecology and Evolution</i> , 2016, 4, .	1.1	6
2190	Full Genome Characterization of Human Influenza A/H3N2 Isolates from Asian Countries Reveals a Rare Amantadine Resistance-Confering Mutation and Novel PB1-F2 Polymorphisms. <i>Frontiers in Microbiology</i> , 2016, 7, 262.	1.5	16
2191	First DNA evidence on the occurrence of Pacific bluefin tuna, <i>Thunnus orientalis</i> in northern Philippine waters. <i>Marine Biodiversity Records</i> , 2016, 9, .	1.2	3
2192	Generic hyper-diversity in <i>Stachybotriaceae</i> . <i>Persoonia: Molecular Phylogeny and Evolution of Fungi</i> , 2016, 36, 156-246.	1.6	112
2193	The First Comprehensive Phylogeny of <i>Coptis</i> (Ranunculaceae) and Its Implications for Character Evolution and Classification. <i>PLoS ONE</i> , 2016, 11, e0153127.	1.1	25
2194	Detection, Prevalence and Phylogenetic Relationships of <i>Demodex</i> spp and further Skin Prostigmata Mites (Acari, Arachnida) in Wild and Domestic Mammals. <i>PLoS ONE</i> , 2016, 11, e0165765.	1.1	27
2195	The Model Organism <i>Hermissenda crassicornis</i> (Gastropoda: Heterobranchia) Is a Species Complex. <i>PLoS ONE</i> , 2016, 11, e0154265.	1.1	35
2196	HIV-1 Variants and Drug Resistance in Pregnant Women from Bata (Equatorial Guinea): 2012-2013. <i>PLoS ONE</i> , 2016, 11, e0165333.	1.1	3
2197	A Radical Solution: The Phylogeny of the Nudibranch Family Fionidae. <i>PLoS ONE</i> , 2016, 11, e0167800.	1.1	37
2198	Species delimitation and phylogenetic analyses of some cosmopolitan species of <i>Hypnea</i> (Rhodophyta) reveal synonyms and misapplied names to <i>H. cervicornis</i> , including a new species from Brazil. <i>Journal of Phycology</i> , 2016, 52, 774-792.	1.0	27
2199	The complete mitochondrial genome of <i>Anopheles minimus</i> (Diptera: Culicidae) and the phylogenetics of known <i>Anopheles</i> mitogenomes. <i>Insect Science</i> , 2016, 23, 353-365.	1.5	21
2200	Evolutionary history of a secondary terrestrial Australian diving beetle (Coleoptera, Dytiscidae) reveals a lineage of high morphological and ecological plasticity. <i>Systematic Entomology</i> , 2016, 41, 650-657.	1.7	15
2201	Molecular Analyses of First Collections of <i>Elaphomyces</i> (Elaphomycetaceae, Eurotiales,) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 Polyphyletic. <i>Cryptogamie, Mycologie</i> , 2016, 37, 3-14.	0.2	12
2202	Unusually high genetic diversity in the Bornean <i>Limnonectes kuhlii</i> -like fanged frogs (Anura:) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50	0.2	12

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2203	Evolutionary patterns and physicochemical properties explain macroinvertebrate sensitivity to heavy metals. <i>Ecological Applications</i> , 2016, 26, 1249-1259.	1.8	23
2204	Error distribution modelling of satellite soil moisture measurements for hydrological applications. <i>Hydrological Processes</i> , 2016, 30, 2223-2236.	1.1	10
2205	Hepatitis B virus full-length genomic mutations and quasispecies in hepatocellular carcinoma. <i>Journal of Gastroenterology and Hepatology (Australia)</i> , 2016, 31, 1638-1645.	1.4	10
2206	Complex longitudinal diversification across South China and Vietnam in Stejneger's pit viper, <i>Viridovipera stejnegeri</i> (Schmidt, 1925) (Reptilia: Serpentes: Viperidae). <i>Molecular Ecology</i> , 2016, 25, 2920-2936.	2.0	18
2207	Genetic variation and evolutionary origins of parthenogenetic <i>Artemia</i> (Crustacea: Anostraca) with different ploidies. <i>Zoologica Scripta</i> , 2016, 45, 421-436.	0.7	23
2208	All that glitters is not <i>Ramularia</i> . <i>Studies in Mycology</i> , 2016, 83, 49-163.	4.5	88
2209	<i>Aeolidia papillosa</i> (Linnaeus, 1761) (Mollusca: Heterobranchia: Nudibranchia), single species or a cryptic species complex? A morphological and molecular study. <i>Zoological Journal of the Linnean Society</i> , 2016, 177, 481-506.	1.0	39
2210	Is there more than one Crocodile Lizard? An Integrative Taxonomic Approach Reveals Vietnamese and Chinese <i>Shinisaurus crocodilurus</i> Represent Separate Conservation and Taxonomic Units. <i>Der Zoologische Garten</i> , 2016, 85, 240-260.	0.3	25
2211	Integrative taxonomy helps to reveal the mask of the genus <i>Gynandropaa</i> (Amphibia: Anura). <i>Trends in Ecology and Evolution</i> , 2016, 31, 422-423.	1.3	7
2212	Using nuclear genes to reconstruct angiosperm phylogeny at the species level: A case study with Brassicaceae species. <i>Journal of Systematics and Evolution</i> , 2016, 54, 438-452.	1.6	13
2213	Macroevolution of hyperdiverse flightless beetles reflects the complex geological history of the Sunda Arc. <i>Scientific Reports</i> , 2016, 6, 18793.	1.6	53
2214	Multi-locus phylogeny using topotype specimens sheds light on the systematics of <i>Niviventer</i> (Rodentia, Muridae) in China. <i>BMC Evolutionary Biology</i> , 2016, 16, 261.	3.2	11
2215	Two new species of <i>Helicascus</i> (Morosphaeriaceae) from submerged wood in northern Thailand. <i>Phytotaxa</i> , 2016, 270, 182.	0.1	10
2216	<i>Diatrypella macrospora</i> sp. nov. and new records of diatrypaceous fungi from Iran. <i>Phytotaxa</i> , 2016, 252, 43.	0.1	25
2217	First record of <i>Esox cisalpinus</i> (Teleostea: Esocidae) in Sardinia with insight on its mitochondrial DNA genetic variability. <i>Italian Journal of Zoology</i> , 2016, 83, 514-523.	0.6	2
2218	Ecology of the Atlantic black skipjack <i>Euthynnus alletteratus</i> (Osteichthyes: Scombridae) in the western Mediterranean Sea inferred by parasitological analysis. <i>Parasitology</i> , 2016, 143, 1330-1339.	0.7	10
2219	New species of <i>Sporoschisma</i> (Chaetosphaeriaceae) from aquatic habitats in Thailand. <i>Phytotaxa</i> , 2016, 289, 147.	0.1	18
2220	<i>Ceramothyrium longivolcaniforme</i> sp. nov., a new species of Chaetothyriaceae from northern Thailand. <i>Phytotaxa</i> , 2016, 267, 51.	0.1	6

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2221	Tuber shii sp. nov., a sister species of T. jinshajiangense from China in Puberulum group. Phytotaxa, 2016, 269, 279.	0.1	4
2222	Gelsolin role in microapocrine secretion. Insect Molecular Biology, 2016, 25, 810-820.	1.0	5
2223	Phylogenetic position and independent generic status of Indocypraea (Asteraceae-Heliantheae-Ecliptinae): evidence from chloroplast DNA sequences. Phytotaxa, 2016, 277, 146.	0.1	2
2224	A new phylogenetic tree sampling method for maximum parsimony bootstrapping and proof-of-concept implementation. , 2016, , .		2
2225	Impacts and shortcomings of genetic clustering methods for infectious disease outbreaks. Virus Evolution, 2016, 2, vew031.	2.2	87
2226	Wuacanthus (Acanthaceae), a new Chinese endemic genus segregated from Justicia (Acanthaceae). Plant Diversity, 2016, 38, 312-321.	1.8	15
2227	The forgotten <i>Calonectria</i> collection: Pouring old wine into new bags. Studies in Mycology, 2016, 85, 159-198.	4.5	38
2228	A molecular phylogeny of the <i>Laelia</i> alliance (Orchidaceae) and a reassessment of <i>Laelia</i> and <i>Schomburgkia</i> . Taxon, 2016, 65, 1249-1262.	0.4	14
2229	Phylogeny, morphology and pathogenicity of Botryosphaeriaceae, Diatrypaceae and Gnomoniaceae associated with branch diseases of hazelnut in Sardinia (Italy). European Journal of Plant Pathology, 2016, 146, 259-279.	0.8	37
2230	Botryosphaeriaceae species associated with wood diseases of stone and pome fruits trees: symptoms and virulence across different hosts in Uruguay. European Journal of Plant Pathology, 2016, 146, 519-530.	0.8	27
2231	Botryosphaeriaceae species associated with stem canker, die-back and fruit rot on apple in Uruguay. European Journal of Plant Pathology, 2016, 146, 637-655.	0.8	36
2232	Epidemiological and molecular data on heterophyid trematode metacercariae found in the muscle of grey mullets (Osteichthyes: Mugilidae) from Sardinia (western Mediterranean Sea). Parasitology Research, 2016, 115, 3409-3417.	0.6	18
2233	<strong>A taxonomic revision of the Yasuni Round-eared bat, <i>Lophostoma yasuni</i> (Chiroptera: Phyllostomidae)</strong> . Zootaxa, 2016, 4114, 246.	0.2	6
2234	Taxonomy and molecular phylogeny of Diatrypaceae (Ascomycota, Xylariales) species from the Brazilian semi-arid region, including four new species. Mycological Progress, 2016, 15, 1.	0.5	27
2235	Phylogeography of the Vermilion Flycatcher species complex: Multiple speciation events, shifts in migratory behavior, and an apparent extinction of a Galápagos-endemic bird species. Molecular Phylogenetics and Evolution, 2016, 102, 152-173.	1.2	30
2236	Ligninsphaeria jonesii gen. et. sp. nov., a remarkable bamboo inhabiting ascomycete. Phytotaxa, 2016, 247, 109.	0.1	8
2237	Cryptosporella platyphylla, a new species associated with Betula platyphylla in China. Phytotaxa, 2016, 253, 285.	0.1	5
2238	Are Phenacoccus solani Ferris and P. defectus Ferris (Hemiptera: Pseudococcidae) distinct species?. Zootaxa, 2016, 4093, 539-51.	0.2	10

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2239	Genetic relationships among freshwater mussel species from fifteen Amazonian rivers and inferences on the evolution of the Hyriidae (Mollusca: Bivalvia: Unionida). <i>Molecular Phylogenetics and Evolution</i> , 2016, 100, 148-159.	1.2	21
2240	A phylogeny of Cephaloziaceae (Jungermanniopsida) based on nuclear and chloroplast DNA markers. <i>Organisms Diversity and Evolution</i> , 2016, 16, 727-742.	0.7	18
2241	Understanding lichenicolous heterobasidiomycetes: new taxa and reproductive innovations in <i>Tremella</i> s.l.. <i>Mycologia</i> , 2016, 108, 381-396.	0.8	23
2242	The Primary Structure of Î²I-Chain of Hemoglobin from Snake Sindhi Krait ( <i>Bungarus sindanus</i> ) Tj ETQq1 1 0.784314 rgBT /Overlock 10	0.7	0
2243	Melanconis (Melanconidaceae) associated with <i>Betula</i> spp. in China. <i>Mycological Progress</i> , 2016, 15, 1.	0.5	12
2244	Geographic differentiation of domesticated einkorn wheat and possible Neolithic migration routes. <i>Heredity</i> , 2016, 117, 135-141.	1.2	24
2245	Phylogeography and species delimitation in convict cichlids (Cichlidae: <i>Amatitlania</i> ): implications for taxonomy and Plio-Pleistocene evolutionary history in Central America. <i>Biological Journal of the Linnean Society</i> , 2016, , .	0.7	7
2246	A 4000-species dataset provides new insight into the evolution of ferns. <i>Molecular Phylogenetics and Evolution</i> , 2016, 105, 200-211.	1.2	214
2247	Mitochondrial genome of <i>Cricetulus migratorius</i> (Rodentia: Cricetidae): Insights into the characteristics of the mitochondrial genome and the phylogenetic relationships of <i>Cricetulus</i> species. <i>Gene</i> , 2016, 595, 121-129.	1.0	13
2248	Revising the <i>Schizoparmaceae</i> : <i>Coniella</i> and its synonyms <i>Pilidiella</i> and <i>Schizoparme</i> . <i>Studies in Mycology</i> , 2016, 85, 1-34.	4.5	60
2249	Genetic diversity and phylogeography of Siberian roe deer, <i>Capreolus pygargus</i> , in central and peripheral populations. <i>Ecology and Evolution</i> , 2016, 6, 7286-7297.	0.8	11
2250	Species diversity of <i>Pseudocercospora</i> from Far East Asia. <i>Mycological Progress</i> , 2016, 15, 1093-1117.	0.5	18
2251	<i>Sporoschisma</i> from submerged wood in Yunnan, China. <i>Mycological Progress</i> , 2016, 15, 1145-1155.	0.5	17
2252	The influence of climate on species distribution over time and space during the late Quaternary. <i>Quaternary Science Reviews</i> , 2016, 149, 188-199.	1.4	16
2253	Molecular and morphological systematics of <i>Elysia</i> Risso, 1818 (Heterobranchia: Sacoglossa) from the Caribbean region. <i>Zootaxa</i> , 2016, 4148, 1-137.	0.2	49
2254	Delimitating cryptic species in the <i>Gracilaria domingensis</i> complex (Gracilariaceae, Rhodophyta) using molecular and morphological data. <i>Journal of Phycology</i> , 2016, 52, 997-1017.	1.0	20
2255	Two new species of <i>Fomitiporia</i> (Hymenochaetales, Basidiomycota) from Tibet, southwest China. <i>Mycologia</i> , 2016, 108, 1010-1017.	0.8	11
2256	Phylogeny, distribution and pathogenicity of <i>Lasiodiplodia</i> species associated with dieback of table grape in the main Brazilian exporting region. <i>Plant Pathology</i> , 2016, 65, 92-103.	1.2	40

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2257	Morphological and molecular evidence support a new truffle, <i>Tuber lannaense</i> , from Thailand. <i>Mycological Progress</i> , 2016, 15, 827-834.	0.5	5
2258	Phylogenetic reassessment of the <i>Chaetomium globosum</i> species complex. <i>Persoonia: Molecular Phylogeny and Evolution of Fungi</i> , 2016, 36, 83-133.	1.6	78
2259	Phylogeny and morphology reveal two new species of <i>Diaporthe</i> from <i>Betula</i> spp. in China. <i>Phytotaxa</i> , 2016, 269, 90.	0.1	29
2260	Phylogenetics of extant and fossil Pinaceae: methods for increasing topological stability. <i>Botany</i> , 2016, 94, 863-884.	0.5	21
2261	Patterns of Genetic Divergence among <i>Myotis californicus</i> , <i>M. ciliolabrum</i> , and <i>M. leibii</i> Based on Amplified Fragment Length Polymorphism. <i>Acta Chiropterologica</i> , 2016, 18, 337-347.	0.2	7
2262	Who Let the CAT Out of the Bag? Accurately Dealing with Substitutional Heterogeneity in Phylogenomic Analyses. <i>Systematic Biology</i> , 2017, 66, syw084.	2.7	57
2263	Changing spots: pseudocryptic speciation in the North Pacific dorid nudibranch <i>Diaulula sandiegensis</i> (Cooper, 1863) (Gastropoda: Heterobranchia). <i>Journal of Molluscan Studies</i> , 2016, 82, 564-574.	0.4	13
2264	Phylogeny, biogeography, and diversification of barn owls (Aves: Strigiformes). <i>Biological Journal of the Linnean Society</i> , 2016, 119, 904-918.	0.7	24
2265	A new nothrotheriid xenarthran from the early Pliocene of Pomata-Ayte (Bolivia): new insights into the caniniform-molariform transition in sloths. <i>Zoological Journal of the Linnean Society</i> , 2016, 178, 679-712.	1.0	17
2266	Resolving the phylogenetic placement of <i>Porobeltraniella</i> and allied genera in the Beltraniaceae. <i>Mycological Progress</i> , 2016, 15, 1119-1136.	0.5	18
2267	Characterization of the complete mitochondrial genome of <i>Phodopus roborovskii</i> (Rodentia: Caviidae). <i>Ecology</i> , 2016, 69, 226-235.	0.6	6
2268	Novel chaetosphaeriaceous hyphomycetes from aquatic habitats. <i>Mycological Progress</i> , 2016, 15, 1157-1167.	0.5	26
2269	Taxonomic revision of Bromeliaceae subfam. Tillandsioideae based on a multi-locus DNA sequence phylogeny and morphology. <i>Phytotaxa</i> , 2016, 279, 1.	0.1	132
2270	Characterization and phylogenetic analysis of the complete mitogenome of <i>Allactaga sibirica</i> (Rodentia: Dipodidae). <i>Biochemical Systematics and Ecology</i> , 2016, 69, 195-203.	0.6	3
2271	Genetic characterization of Australian <i>Mycoplasma bovis</i> isolates through whole genome sequencing analysis. <i>Veterinary Microbiology</i> , 2016, 196, 118-125.	0.8	37
2272	Host records of <i>Grammia ursina</i> Schmidt, 2009 (Lepidoptera: Noctuidae: Arctiinae) on San Clemente Island and its potential effect on rare plant restoration. <i>Pan-Pacific Entomologist</i> , 2016, 92, 151-156.	0.1	0
2273	Transfer of the leafy liverwort <i>Xenochila</i> from Plagiochilaceae (Lophocoleineae) to Jungermanniaceae (Jungermanniineae). <i>Plant Systematics and Evolution</i> , 2016, 302, 891-899.	0.3	4
2274	High genetic diversity in the <i>Culex pipiens</i> complex from a West Nile Virus epidemic area in Southern Europe. <i>Parasites and Vectors</i> , 2016, 9, 150.	1.0	7

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2275	Molecular Characterization of <i>Acanthamoeba</i> spp. Occurring in Water Bodies and Patients in Poland and Redefinition of Polish T16 Genotype. <i>Journal of Eukaryotic Microbiology</i> , 2016, 63, 262-270.	0.8	7
2276	Molecular phylogeny of the highly disjunct cliff water beetles from South Africa and China (Coleoptera: Aspidytidae). <i>Zoological Journal of the Linnean Society</i> , 2016, 176, 537-546.	1.0	19
2277	Molecular phylogeography of white-striped tree viper ( <i>Trimeresurus</i> ; Viperidae). <i>Zoologica Scripta</i> , 2016, 45, 252-262.	0.7	17
2278	Utilizing ribosomal DNA gene marker regions to characterize the metacercariae (Trematoda: Digenea) parasitizing piscine intermediate hosts in Manipur, Northeast India. <i>Journal of Parasitic Diseases</i> , 2016, 40, 330-338.	0.4	7
2279	<i>Symplectella rowi</i> (Porifera: Hexactinellida: Lyssacinosa) is a rossellid, not a euplectellid. <i>Journal of the Marine Biological Association of the United Kingdom</i> , 2016, 96, 291-295.	0.4	3
2280	Towards a monophyletic classification of Lejeuneaceae IV: reinstatement of <i>Allorgella</i> , transfer of <i>Microlejeunea aphanella</i> to <i>Vitalianthus</i> and refinements of the subtribal classification. <i>Plant Systematics and Evolution</i> , 2016, 302, 187-201.	0.3	32
2281	Bovine <i>Staphylococcus aureus</i> : Subtyping, evolution, and zoonotic transfer. <i>Journal of Dairy Science</i> , 2016, 99, 515-528.	1.4	86
2282	Can you keep a secret? Introducing the RT-based Concealed Information Test to children. <i>Psychology, Crime and Law</i> , 2016, 22, 276-301.	0.8	6
2283	Towards a revised generic classification of lecanoroid lichens (Lecanoraceae, Ascomycota) based on molecular, morphological and chemical evidence. <i>Fungal Diversity</i> , 2016, 78, 293-304.	4.7	72
2284	Biological role in the transformation of platinum-group mineral grains. <i>Nature Geoscience</i> , 2016, 9, 294-298.	5.4	46
2285	A New Development to Aid Interpretation of Hierarchical Cluster Analysis of Repertory Grid Data. <i>Journal of Constructivist Psychology</i> , 2016, 29, 368-381.	0.7	4
2286	Reduced genetic diversity and increased reproductive isolation follow population-level loss of larval dispersal in a marine gastropod. <i>Evolution; International Journal of Organic Evolution</i> , 2016, 70, 18-37.	1.1	25
2287	Identification of <i>Bacillus megaterium</i> and <i>Microbacterium liquefaciens</i> genes involved in metal resistance and metal removal. <i>Canadian Journal of Microbiology</i> , 2016, 62, 505-513.	0.8	24
2288	Subfamilial and tribal relationships of Ranunculaceae: evidence from eight molecular markers. <i>Plant Systematics and Evolution</i> , 2016, 302, 419-431.	0.3	38
2289	The noncoding trnH-psbA spacer, as an effective DNA barcode for aquatic freshwater plants, reveals prohibited invasive species in aquarium trade in South Africa. <i>South African Journal of Botany</i> , 2016, 102, 208-216.	1.2	12
2290	A phylogeny of Lophocoleaceae-Plagiochilaceae-Brevianthaceae and a revised classification of Plagiochilaceae. <i>Organisms Diversity and Evolution</i> , 2016, 16, 481-495.	0.7	27
2291	Species of the <i>Colletotrichum acutatum</i> complex associated with anthracnose diseases of fruit in Brazil. <i>Fungal Biology</i> , 2016, 120, 547-561.	1.1	71
2292	Maidenhair Ferns, <i>Adiantum</i> , are Indeed Monophyletic and Sister to Shoestring Ferns, Vittarioids (Pteridaceae). <i>Systematic Botany</i> , 2016, 41, 17-23.	0.2	21

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2293	PCR-based assessment of shellfish traceability and sustainability in international Mediterranean seafood markets. <i>Food Chemistry</i> , 2016, 202, 302-308.	4.2	21
2294	Two new species of <i>Coltricia</i> (Hymenochaetaceae, Basidiomycota) from southern China based on evidence from morphology and DNA sequence data. <i>Mycological Progress</i> , 2016, 15, 1.	0.5	7
2295	A new method for studying the evolutionary origin of the SAR11 clade marine bacteria. <i>Molecular Phylogenetics and Evolution</i> , 2016, 98, 271-279.	1.2	13
2296	Phylogenetic relationships of the New World titi monkeys ( <i>Callicebus</i> ): first appraisal of taxonomy based on molecular evidence. <i>Frontiers in Zoology</i> , 2016, 13, 10.	0.9	140
2297	Molecular Phylogeography of Harvest Mice ( <i>Reithrodontomys megalotis</i> ) Based on Cytochrome b DNA Sequences. <i>Journal of Mammalian Evolution</i> , 2016, 23, 297-307.	1.0	8
2298	Pliocene–Pleistocene lineage diversifications in the Eastern Indigo Snake ( <i>Drymarchon couperi</i> ) in the Southeastern United States. <i>Molecular Phylogenetics and Evolution</i> , 2016, 98, 111-122.	1.2	16
2299	<i>Cyphobasidium</i> gen. nov., a new lichen-inhabiting lineage in the Cystobasidiomycetes (Pucciniomycotina, Basidiomycota, Fungi). <i>Fungal Biology</i> , 2016, 120, 1468-1477.	1.1	46
2300	Molecular analysis of Chinese truffles resembling <i>Tuber californicum</i> in morphology reveals a rich pattern of species diversity with emphasis on four new species. <i>Mycologia</i> , 2016, 108, 344-353.	0.8	13
2301	Phylogenetic analyses of Chinese <i>Tuber</i> species that resemble <i>T. borchii</i> reveal the existence of the new species <i>T. hubeiense</i> and <i>T. wumengense</i> . <i>Mycologia</i> , 2016, 108, 354-362.	0.8	9
2302	Multiple Origins of Eukaryotic <i>cox15</i> Suggest Horizontal Gene Transfer from Bacteria to Jakobid Mitochondrial DNA. <i>Molecular Biology and Evolution</i> , 2016, 33, 122-133.	3.5	21
2303	How Should Genes and Taxa be Sampled for Phylogenomic Analyses with Missing Data? An Empirical Study in Iguanian Lizards. <i>Systematic Biology</i> , 2016, 65, 128-145.	2.7	155
2304	Geographic isolation drives divergence of uncorrelated genetic and song variation in the Ruddy-capped Nightingale-Thrush ( <i>Catharus frantzii</i> ; Aves: Turdidae). <i>Molecular Phylogenetics and Evolution</i> , 2016, 94, 74-86.	1.2	28
2305	Molecular diversity of some species belonging to the genus <i>Daphnia</i> O. F. MÅ¼ller, 1785 (Crustacea: Cladocera) in Turkey. <i>Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis</i> , 2017, 28, 424-433.	0.7	5
2306	Frogs at the summits: phylogeny of the Andean frogs of the genus <i>Telmatobius</i> (Anura). <i>Tj ETQq1 1 0.784314 rgBT /Oyerlock 10</i>	1.5	27
2307	First record of the Aeolid <i>Anteaeolidiella fijensis</i> (Nudibranchia, Aeolidiidae) from India. <i>Marine Biodiversity</i> , 2017, 47, 823-830.	0.3	1
2308	Microbiological and molecular identification of bacterial species isolated from nasal and oropharyngeal mucosa of fuel workers in Riyadh, Saudi Arabia. <i>Saudi Journal of Biological Sciences</i> , 2017, 24, 1281-1287.	1.8	7
2309	Which molecular markers for assessing which taxonomic level? The case study of the mite family Phytoseiidae (Acari: Mesostigmata). <i>Cladistics</i> , 2017, 33, 251-267.	1.5	13
2310	Survey for the presence of ascaridoid larvae in the cinnamon flounder <i>Pseudorhombus cinnamomeus</i> (Temminck & Schlegel) (Pleuronectiformes: Paralichthyidae). <i>International Journal of Food Microbiology</i> , 2017, 241, 108-116.	2.1	20



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2312	The phylogenetic position of eriophyoid mites (superfamily Eriophyoidea) in Acariformes inferred from the sequences of mitochondrial genomes and nuclear small subunit (18S) rRNA gene. <i>Molecular Phylogenetics and Evolution</i> , 2017, 109, 271-282.	1.2	58
2313	Evaluation of Systematic Position of Helicoprordontids and Chaeneids (Ciliophora, Litostomatea): An Attempt to Break Long Branches in 18S rRNA Gene Phylogenies. <i>Journal of Eukaryotic Microbiology</i> , 2017, 64, 608-621.	0.8	3
2314	The Calibrated Phylogeny of the <i>Drosophila fasciola</i> Subgroup ( <i>D. repleta</i> Group Wasserman) Indicates Neogene Diversification of Its Internal Branches. <i>Neotropical Entomology</i> , 2017, 46, 537-545.	0.5	2
2315	Comparative morphology and genetics of two populations of spiny lizards (genus <i>Sceloporus</i> ) from Central Mexico. <i>Zoologischer Anzeiger</i> , 2017, 267, 21-30.	0.4	8
2316	Phylogeny and comparative genomic analysis of Pteriomorphia (Mollusca: Bivalvia) based on complete mitochondrial genomes. <i>Marine Biology Research</i> , 2017, 13, 255-268.	0.3	12
2317	Genetic diversity of Indian jujube cultivars using SCoT, ISSR, and rDNA markers. <i>Tree Genetics and Genomes</i> , 2017, 13, 1.	0.6	20
2318	Molecular identification of honey entomological origin based on bee mitochondrial 16S rRNA and COI gene sequences. <i>Food Control</i> , 2017, 78, 150-159.	2.8	37
2319	Reconstructing the geographic origin of the New World jays. <i>Neotropical Biodiversity</i> , 2017, 3, 80-92.	0.2	12
2320	The role of ecological factors in determining phylogeographic and population genetic structure of two sympatric island skinks ( <i>Plestiodon kishinouyei</i> and <i>P. stimpsonii</i> ). <i>Genetica</i> , 2017, 145, 223-234.	0.5	3
2321	A New Genus and Two New Species of Arboreal Toads from the Highlands of Sumatra with a Phylogeny of Sundaland Toad Genera. <i>Herpetologica</i> , 2017, 73, 63-75.	0.2	11
2322	Species diversity and molecular systematics of <i>Fibroporia</i> (Polyporales, Basidiomycota) and its related genera. <i>Mycological Progress</i> , 2017, 16, 521-533.	0.5	19
2323	Phylogeny and taxonomy of the scab and spot anthracnose fungus <i>Elsinoë</i> ( <i>Myriangiiales</i> ) Tj ETQq0,0,0 rgBT /Overlock 1	4.5	59
2324	The evolutionary history and taxonomic reevaluation of the Japanese coral snake, <i>Sinomicrurus japonicus</i> (Serpentes, Elapidae), endemic to the Ryukyu Archipelago, Japan, by use of molecular and morphological analyses. <i>Journal of Zoological Systematics and Evolutionary Research</i> , 2017, 55, 156-166.	0.6	6
2325	Revisiting phylogenetic relationships in Phoradendreae (Viscaceae): utility of the <i>trnL-F</i> region of chloroplast DNA and presence of a homoplasious inversion in the intergenic spacer. <i>Botany</i> , 2017, 95, 247-258.	0.5	4
2326	Genome sequencing of an Indian peste des petits ruminants virus isolate, Izatnagar/94, and its implications for virus diversity, divergence and phylogeography. <i>Archives of Virology</i> , 2017, 162, 1677-1693.	0.9	23
2327	Four new species of <i>Tubeufia</i> (Tubeufiaceae, Tubeufiales) from Thailand. <i>Mycological Progress</i> , 2017, 16, 403-417.	0.5	23
2328	<i>Steinernema ralatorein</i> , sp. Isolated from Sugarcane Areas at Veracruz, Mexico. <i>Southwestern Entomologist</i> , 2017, 42, 171-190.	0.1	11

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2329	Complex biogeographic scenarios revealed in the diversification of the largest woodpecker radiation in the New World. <i>Molecular Phylogenetics and Evolution</i> , 2017, 112, 53-67.	1.2	15
2330	Genetic evidence supporting the taxonomic separation of the Arabian and Northwest African subspecies of the desert hedgehog ( <i>Paraechinus aethiopicus</i> ). <i>Gene</i> , 2017, 620, 54-65.	1.0	10
2331	A new species of <i>Plestiodon</i> (Squamata: Scincidae) from the Senkaku Group, Ryukyu Archipelago, Japan. <i>Zootaxa</i> , 2017, 4254, 520.	0.2	3
2332	The affinities of <i>Homo floresiensis</i> based on phylogenetic analyses of Cranial, dental, and postcranial characters. <i>Journal of Human Evolution</i> , 2017, 107, 107-133.	1.3	89
2333	Strong spatial genetic congruence between a wood-feeding cockroach and its bacterial endosymbiont, across a topographically complex landscape. <i>Journal of Biogeography</i> , 2017, 44, 1500-1511.	1.4	17
2334	Integrative systematic analyses of the genus <i>Chodsigoa</i> (Mammalia: Eulipotyphla: Soricidae), with descriptions of new species. <i>Zoological Journal of the Linnean Society</i> , 2017, 180, 694-713.	1.0	17
2335	Like a bat out of heaven: the phylogeny and diversity of the bat-winged slugs (Heterobranchia: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 3	1.0	7
2336	Redescription of <i>Keronopsis helluo</i> Penard, 1922 from Antarctica and <i>Paraholosticha pannonica</i> Gell�rt and Tam�s, 1959 from Alaska (Ciliophora, Hypotricha). <i>European Journal of Protistology</i> , 2017, 60, 102-118.	0.5	15
2337	Contribution of filamentous fungi to the musty odorant 2,4,6-trichloroanisole in water supply reservoirs and associated drinking water treatment plants. <i>Chemosphere</i> , 2017, 182, 223-230.	4.2	23
2338	Multi-locus phylogeny and morphology reveal five new species of <i>Fomitiporia</i> (Hymenochaetaceae) from China. <i>Mycological Progress</i> , 2017, 16, 687-701.	0.5	14
2339	Systematics and Molecular Phylogeny of the Ciliate Genus <i>Pseudokeronopsis</i> (Ciliophora,) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 3	0.8	16
2340	A new species of Rock Gecko genus <i>Cnemaspis</i> (Squamata: Gekkonidae) from Western Sarawak, Malaysia. <i>Zootaxa</i> , 2017, 4258, 525.	0.2	11
2341	Phylogenetic analysis and evolution of morphological characters in <i>Metrodorea</i> and related species in Rutoideae (Rutaceae). <i>Plant Systematics and Evolution</i> , 2017, 303, 927-943.	0.3	6
2342	Ribosomal DNA analysis of isolates of the liver fluke <i>Opisthorchis pedicellata</i> (Verma, 1927) from two siluroid fish species in India. <i>Journal of Helminthology</i> , 2017, 91, 302-311.	0.4	4
2343	Inferring Trees. <i>Methods in Molecular Biology</i> , 2017, 1525, 349-377.	0.4	5
2344	CIE Standard Sky classification by accessible climatic indices. <i>Renewable Energy</i> , 2017, 113, 347-356.	4.3	29
2345	First report of <i>Phyllosticta citricarpa</i> and description of two new species, <i>P. Aparacapitalensis</i> and <i>P. Aparacitricarpa</i> , from citrus in Europe. <i>Studies in Mycology</i> , 2017, 87, 161-185.	4.5	79
2346	Monoblepharidomycetes diversity includes new parasitic and saprotrophic species with highly intronized rDNA. <i>Fungal Biology</i> , 2017, 121, 729-741.	1.1	23

#	ARTICLE	IF	CITATIONS
2347	A new color pattern of the <i>Bungarus candidus</i> complex (Squamata: Elapidae) from Vietnam based on morphological and molecular data. <i>Zootaxa</i> , 2017, 4268, 563.	0.2	3
2348	<i>Chloridium terricola</i> sp. nov. from China. <i>Mycotaxon</i> , 2017, 132, 79-86.	0.1	2
2349	Morphologic and phylogenetic studies of two hypotrichous ciliates, with notes on morphogenesis in <i>Gastrostyla steinii</i> Engelmann, 1862 (Ciliophora, Hypotrichia). <i>European Journal of Protistology</i> , 2017, 60, 119-133.	0.5	16
2350	A new genus of megalonychid ground sloth (Mammalia, Xenarthra) from the late Pleistocene of Quintana Roo, Mexico. <i>Journal of Vertebrate Paleontology</i> , 2017, 37, e1307206.	0.4	21
2351	Integrative taxonomy of the Central African forest chameleon, <i>Kinyongia adolfifriederici</i> (Sauria: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 58 the Linnean Society, 0, , .	1.0	6
2352	A newly recognised Australian endemic species of <i>Austrolecanium</i> Gullan & Hodgson 1998 (Hemiptera: Coccidae) from Queensland. <i>Zootaxa</i> , 2017, 4272, 119.	0.2	4
2353	An old familiar face: <i>Tremella anaptychia</i> sp. nov. (Tremellales, Basidiomycota). <i>Phytotaxa</i> , 2017, 307, 254.	0.1	13
2354	Structure and in vitro activities of a Copper II-chelating anionic peptide from the venom of the scorpion <i>Tityus stigmurus</i> . <i>Peptides</i> , 2017, 94, 91-98.	1.2	14
2355	A new species of <i>Leptotalax</i> (Anura: Megophryidae) from Vietnam. <i>Zootaxa</i> , 2017, 4273, 61-79.	0.2	21
2356	Phylogenetic Analyses. , 2017, , 143-172.		1
2357	A new species of <i>Pseudocalotes</i> (Squamata: Agamidae) from the Bukit Barisan Range of Sumatra with an Estimation of its phylogeny. <i>Zootaxa</i> , 2017, 4276, 215-232.	0.2	14
2358	Ultraconserved elements show utility in phylogenetic inference of <sc>A</sc>dephaga (<sc>C</sc>oleoptera) and suggest paraphyly of "Hydradephaga"™. <i>Systematic Entomology</i> , 2017, 42, 786-795.	1.7	77
2359	Identification of bakanae disease resistance loci in japonica rice through genome wide association study. <i>Rice</i> , 2017, 10, 29.	1.7	43
2360	<i>Diaporthe</i> species occurring on <i>Senna bicapsularis</i> in southern China, with descriptions of two new species. <i>Phytotaxa</i> , 2017, 302, 145.	0.1	19
2361	Six new soil-inhabiting <i>Cladosporium</i> species from plateaus in China. <i>Mycologia</i> , 2017, 109, 244-260.	0.8	19
2362	<i>Diatrypella tectonae</i> and <i>Peroneutypa mackenziei</i> spp. nov. (Diatrypaceae) from northern Thailand. <i>Mycological Progress</i> , 2017, 16, 463-476.	0.5	25
2363	Genotypic and Phenotypic Characterization of <i>Streptomyces</i> Species Causing Potato Common Scab in Uruguay. <i>Plant Disease</i> , 2017, 101, 1362-1372.	0.7	18
2364	A new species of <i>Antrodia</i> (Basidiomycota, Polyporales) from juniper forest of Uzbekistan. <i>Phytotaxa</i> , 2017, 303, 47.	0.1	5

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2365	The accuracy of confidence intervals for field normalised indicators. <i>Journal of Informetrics</i> , 2017, 11, 530-540.	1.4	9
2366	Molecular phylogenetic diversity in the widespread lizard <i>Cercosaura ocellata</i> (Reptilia: Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50	0.5	5
2367	Diversity of culturable fungi inhabiting petroleum-contaminated soils in Southern Iran. <i>Antonie Van Leeuwenhoek</i> , 2017, 110, 903-923.	0.7	11
2368	Additions to <i>Lindgomyces</i> (Lindgomycetaceae, Pleosporales, Dothideomycetes), including two new species occurring on submerged wood from North Carolina, USA, with notes on secondary metabolite profiles. <i>Mycological Progress</i> , 2017, 16, 535-552.	0.5	14
2369	Phylogenetic relationships of weaverbirds (Aves: Ploceidae): A first robust phylogeny based on mitochondrial and nuclear markers. <i>Molecular Phylogenetics and Evolution</i> , 2017, 109, 21-32.	1.2	19
2370	Exploring the shell-based taxonomy of the Sri Lankan land snail <i>Corilla</i> H. and A. Adams, 1855 (Pulmonata: Corillidae) using mitochondrial DNA. <i>Molecular Phylogenetics and Evolution</i> , 2017, 107, 609-618.	1.2	6
2371	A molecular phylogeny of <i>Dichocarpum</i> (Ranunculaceae): Implications for eastern Asian biogeography. <i>Molecular Phylogenetics and Evolution</i> , 2017, 107, 594-604.	1.2	28
2372	Ecomorphological adaptations in Collembola in relation to feeding strategies and microhabitat. <i>European Journal of Soil Biology</i> , 2017, 78, 82-91.	1.4	35
2373	Wood-rotting basidiomycetes associated with declining native trees in timber harvesting compartments of the Garden Route National Park of South Africa. <i>Austral Ecology</i> , 2017, 42, 947-963.	0.7	9
2374	New species of <i>Talaromyces</i> isolated from maize, indoor air, and other substrates. <i>Mycologia</i> , 2017, 109, 1-20.	0.8	16
2375	Integrative taxonomy of the <i>Felimare californiensis</i> and <i>F. ghiselini</i> species complex (Nudibranchia: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50	0.4	7
2376	A New Species and New Records of <i>Cyphostemma</i> (Vitaceae) from China and Vietnam Based on Morphological and Molecular Evidence. <i>Systematic Botany</i> , 2017, 42, 449-457.	0.2	9
2377	Identification of Forensically Important Blow Flies (Diptera: Calliphoridae) in China Based on COI. <i>Journal of Medical Entomology</i> , 2017, 54, 1193-1200.	0.9	13
2378	Taxonomy assignment approach determines the efficiency of identification of OTUs in marine nematodes. <i>Royal Society Open Science</i> , 2017, 4, 170315.	1.1	36
2379	Associations between socio-demographic characteristics and chemical concentrations contributing to cumulative exposures in the United States. <i>Journal of Exposure Science and Environmental Epidemiology</i> , 2017, 27, 544-550.	1.8	26
2380	Longitudinal sequencing of HIV-1 infected patients with low-level viremia for years while on ART shows no indications for genetic evolution of the virus. <i>Virology</i> , 2017, 510, 185-193.	1.1	25
2381	Molecular phylogenetic analysis and character evolution in <i>Pseudostellaria</i> (Caryophyllaceae) and description of a new genus, <i>Hartmaniella</i> , in North America. <i>Botanical Journal of the Linnean Society</i> , 2017, 184, 444-456.	0.8	12
2382	Generic recircumscription of <i>Parasenecio</i> (Asteraceae: Senecioneae) based on nuclear ribosomal and plastid DNA sequences, with descriptions of two new genera. <i>Botanical Journal of the Linnean Society</i> , 2017, 184, 418-443.	0.8	12

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2383	<i>Coniella vitis</i> sp. nov. Is the Common Pathogen of White Rot in Chinese Vineyards. <i>Plant Disease</i> , 2017, 101, 2123-2136.	0.7	29
2384	Molecular phylogenetics and definition of the <i>Acrapex minima</i> Janse group (Lepidoptera, Noctuidae.) <i>Tj ETQq1 1 0.784314 rgBT /Overl</i> <i>Societe Entomologique De France</i> , 2017, 53, 219-235.	0.4	4
2385	Three new species of <i>Phylloporia</i> (Hymenochaetales) with dimitic hyphal systems from tropical China. <i>Mycologia</i> , 2017, 109, 951-964.	0.8	8
2386	Aligning evidence: concerns regarding multiple sequence alignments in estimating the phylogeny of the Nudibranchia suborder Doridina. <i>Royal Society Open Science</i> , 2017, 4, 171095.	1.1	31
2387	Global population structure and adaptive evolution of aflatoxin-producing fungi. <i>Ecology and Evolution</i> , 2017, 7, 9179-9191.	0.8	25
2388	Preliminary phylogeny of <i>Coemansia</i> (Kickxellales), with descriptions of four new species from Taiwan. <i>Mycologia</i> , 2017, 109, 1-17.	0.8	4
2389	<i>Mycosphaerellaceae</i> : Chaos or clarity?. <i>Studies in Mycology</i> , 2017, 87, 257-421.	4.5	119
2390	A Consensus Method for Ancestral Recombination Graphs. <i>Journal of Molecular Evolution</i> , 2017, 84, 129-138.	0.8	0
2391	Integrative taxonomy and biogeography of the genus <i>Bulbaeolidia</i> (Nudibranchia: Aeolidida). <i>Journal of Molluscan Studies</i> , 2017, 83, 440-450.	0.4	2
2392	Visualizing phylogenetic tree landscapes. <i>BMC Bioinformatics</i> , 2017, 18, 85.	1.2	18
2393	<i>Cantharellus</i> (Cantharellales, Basidiomycota) revisited in Europe through a multigene phylogeny. <i>Fungal Diversity</i> , 2017, 83, 263-292.	4.7	38
2394	Neogene diversification in the temperate lichen-forming fungal genus <i>Parmelia</i> (Parmeliaceae.) <i>Tj ETQq1 1 0.784314 rgBT /Overl</i> <i>0.5 14</i>	0.5	14
2395	Phylogeny of anaerobic fungi (phylum Neocallimastigomycota), with contributions from yak in China. <i>Antonie Van Leeuwenhoek</i> , 2017, 110, 87-103.	0.7	47
2396	A first higher-level time-calibrated phylogeny of antlions (Neuroptera: Myrmeleontidae). <i>Molecular Phylogenetics and Evolution</i> , 2017, 107, 103-116.	1.2	30
2397	Phylogenetic relationships of Mediterranean and North-East Atlantic Cantharidinae and notes on Stomatellinae (Vetigastropoda: Trochidae). <i>Molecular Phylogenetics and Evolution</i> , 2017, 107, 64-79.	1.2	23
2398	Segmentation, Splitting, and Classification of Overlapping Bacteria in Microscope Images for Automatic Bacterial Vaginosis Diagnosis. <i>IEEE Journal of Biomedical and Health Informatics</i> , 2017, 21, 1095-1104.	3.9	36
2399	Analysis of phylogeny, distribution, and pathogenicity of Botryosphaeriaceae species associated with gummosis of <i>Anacardium</i> in Brazil, with a new species of <i>Lasiodiplodia</i> . <i>Fungal Biology</i> , 2017, 121, 437-451.	1.1	47
2400	From a lost world: an integrative phylogenetic analysis of <i>Ansonia Stoliczka, 1870</i> (Lissamphibia:) <i>Tj ETQq1 1 0.784314 rgBT /Overl</i> <i>0.7 10</i>	0.7	10

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2401	DNA barcoding for identification of consumer-relevant mushrooms: A partial solution for product certification?. <i>Food Chemistry</i> , 2017, 214, 383-392.	4.2	68
2402	<scp>DNA</scp> barcoding for the species identification of commercially important fishery products in Indonesian markets. <i>International Journal of Food Science and Technology</i> , 2017, 52, 266-274.	1.3	18
2403	Families, genera, and species of Botryosphaerales. <i>Fungal Biology</i> , 2017, 121, 322-346.	1.1	134
2404	Phylogenetic relationships of the largest lungless tetrapod ( <i>Gymnophiona</i> , <i>Atretochoana</i> ) and the evolution of lunglessness in caecilians. <i>Zoologica Scripta</i> , 2017, 46, 255-263.	0.7	5
2405	Leapfrogging into new territory: How Mascarene ridged frogs diversified across Africa and Madagascar to maintain their ecological niche. <i>Molecular Phylogenetics and Evolution</i> , 2017, 106, 254-269.	1.2	44
2406	<i>Dictyosporium wuyiense</i> sp. nov. from Wuyi Mountain China. <i>Phytotaxa</i> , 2017, 314, 251.	0.1	6
2407	<i>Heterobasidion amyloideopsis</i> sp. nov. (Basidiomycota, Russulales) evidenced by morphological characteristics and phylogenetic analysis. <i>Phytotaxa</i> , 2017, 317, 199.	0.1	11
2408	<i>Gyrodon suthepensis</i> (Boletales, Basidiomycota), a new ectomycorrhizal fungus from northern Thailand and its ecomycorrhizal association. <i>Phytotaxa</i> , 2017, 321, 181.	0.1	0
2409	Morphological and phylogenetic insights resolve <i>Plenodomus sinensis</i> (Leptosphaeriaceae) as a new species. <i>Phytotaxa</i> , 2017, 324, 73.	0.1	8
2410	Multiple gene genealogy reveals high genetic diversity and evidence for multiple origins of Chinese <i>Plasmopara viticola</i> population. <i>Scientific Reports</i> , 2017, 7, 17304.	1.6	16
2411	Establishing the phylogeny of <i>Prochlorococcus</i> with a new alignment-free method. <i>Ecology and Evolution</i> , 2017, 7, 11057-11065.	0.8	7
2412	<i>Novomicrothelia pandanicola</i> sp. nov., a non-lichenized Trypetheliaceae species from Pandanus. <i>Phytotaxa</i> , 2017, 321, 254.	0.1	4
2413	High species diversity in <i>Colletotrichum</i> associated with citrus diseases in Europe. <i>Persoonia: Molecular Phylogeny and Evolution of Fungi</i> , 2017, 39, 32-50.	1.6	86
2414	<div class="page" title="Page 1"><div class="layoutArea"><div class="column">Species circumscription of the <i>Caltha leptosepala</i> polyploid complex (Ranunculaceae) based on molecular and morphological data</div></div></div>. <i>Phytotaxa</i> , 2017, 316, 201.	0.1	1
2415	Clarifying the identity of <i>Geastrum campestre</i> var. <i>famatinum</i> (Geastrales, Basidiomycota). <i>Phytotaxa</i> , 2017, 328, 159.	0.1	1
2416	<i>Gymnosporangium przewalskii</i> sp. nov. (Pucciniales, Basidiomycota) from China and its life cycle. <i>Phytotaxa</i> , 2017, 311, 67.	0.1	6
2417	The new Hispaniolan genus <i>Tainus</i> (Rubiaceae) constitutes an isolated lineage in the Caribbean biodiversity hotspot. <i>Willdenowia</i> , 2017, 47, 259.	0.5	8
2418	Unravelling the phylogeny of the root-emiparasitic genus <i>Odontites</i> (tribe Rhinanthaeae,) Tj ETQq1 1 0.784314 rgBT <sub>0,4</sub> /Overlook	0.4	24

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2419	<i>Tuber griseolivaceum</i> sp. nov., a new olive-gray truffle species from China based on morphological and DNA data. <i>Phytotaxa</i> , 2017, 309, 166.	0.1	1
2420	<i>Neoalbatrellus odorus</i> sp. nov. (Albatrellaceae, Russulales) from Southwest China. <i>Phytotaxa</i> , 2017, 309, 217.	0.1	5
2421	<i>Helicosporium luteosporum</i> sp. nov. and <i>Acanthohelicospora aurea</i> (Tubeufiaceae, Tubeufiales) from terrestrial habitats. <i>Phytotaxa</i> , 2017, 319, 241.	0.1	16
2422	Host and geographic range extensions of <i>Melanconiella</i> , with a new species <i>M. cornuta</i> in China. <i>Phytotaxa</i> , 2017, 327, 252.	0.1	10
2423	Ecotypes and evolutionary significant units in endangered North African gazelles. <i>Biological Journal of the Linnean Society</i> , 2017, 122, 286-300.	0.7	9
2424	A newly recognised species that has been confused with the global polyphagous pest scale insect, <i>Coccus hesperidum</i> Linnaeus (Hemiptera: Coccoidea: Coccidae). <i>Zootaxa</i> , 2017, 4320, .	0.2	5
2425	A new species of <i>Plestiodon</i> (Squamata: Scincidae) from the Balsas Basin, Mexico. <i>Zootaxa</i> , 2017, 4365, 149-172.	0.2	7
2426	Robust Phylogeny of <i>Tetrastigma</i> (Vitaceae) Based on Ten Plastid DNA Regions: Implications for Infrageneric Classification and Seed Character Evolution. <i>Frontiers in Plant Science</i> , 2017, 8, 590.	1.7	22
2427	Emerging citrus diseases in Europe caused by species of <i>Diaporthe</i> . <i>IMA Fungus</i> , 2017, 8, 317-334.	1.7	98
2428	A morphological and molecular study of <i>Psilops</i> , a replacement name for the Brazilian microteiid lizard genus <i>Psilophthalmus</i> Rodrigues 1991 (Squamata, Gymnophthalmidae), with the description of two new species. <i>Zootaxa</i> , 2017, 4286, .	0.2	11
2429	Demulsification of crude oil-in-water emulsions by means of fungal spores. <i>PLoS ONE</i> , 2017, 12, e0170985.	1.1	18
2430	Characterisation of the nicotianamine aminotransferase and deoxymugineic acid synthase genes essential to Strategy II iron uptake in bread wheat ( <i>Triticum aestivum</i> L.). <i>PLoS ONE</i> , 2017, 12, e0177061.	1.1	55
2431	High prevalence and diversity of HIV-1 non-B genetic forms due to immigration in southern Spain: A phylogeographic approach. <i>PLoS ONE</i> , 2017, 12, e0186928.	1.1	7
2432	Insight into Central Asian flora from the Cenozoic Tianshan montane origin and radiation of <i>Lagochilus</i> (Lamiaceae). <i>PLoS ONE</i> , 2017, 12, e0178389.	1.1	10
2433	Phylogenetic placement within <i>Lecanoromycetes</i> of lichenicolous fungi associated with <i>Cladonia</i> and some other genera. <i>Persoonia: Molecular Phylogeny and Evolution of Fungi</i> , 2017, 39, 91-117.	1.6	40
2434	DNA barcoding evaluation and implications for phylogenetic relationships in Lauraceae from China. <i>PLoS ONE</i> , 2017, 12, e0175788.	1.1	38
2435	<i>Periconia thailandica</i> (Periconiaceae), a new species from Thailand. <i>Phytotaxa</i> , 2017, 323, 253.	0.1	9
2436	Total evidence phylogeny and evolutionary timescale for Australian faunivorous marsupials (Dasyuromorphia). <i>BMC Evolutionary Biology</i> , 2017, 17, 240.	3.2	57

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2437	Phylogenetic reconstruction of the genus <i>Helianthemum</i> (Cistaceae) using plastid and nuclear DNA sequences: Systematic and evolutionary inferences. <i>Taxon</i> , 2017, 66, 868-885.	0.4	25
2438	Landscape Genetics of <i>Aedes mcintoshii</i> (Diptera: Culicidae), an Important Vector of Rift Valley Fever Virus in Northeastern Kenya. <i>Journal of Medical Entomology</i> , 2017, 54, 1258-1265.	0.9	3
2439	<i>Calonectria</i> species isolated from Eucalyptus plantations and nurseries in South China. <i>IMA Fungus</i> , 2017, 8, 259-286.	1.7	37
2440	Species-level phylogenetic analysis in the <i>Relhania</i> clade and a new generic treatment of species previously assigned to <i>Macowania</i> and <i>Arrowsmithia</i> (Asteraceae.) <i>Tj ETQq1 1 0.784314 rgrBT /Overlock 10 Tf 5</i>	0.7	15
2441	<i>Jenynsia lineata</i> species complex, revision and new species description (Cyprinodontiformes.) <i>Tj ETQq0 0 0 rgrBT /Overlock 10 Tf 5</i>	0.7	15
2442	Diversity and evolution of four-domain voltage-gated cation channels of eukaryotes and their ancestral functional determinants. <i>Scientific Reports</i> , 2018, 8, 3539.	1.6	24
2443	Phylogenomic analysis demonstrates a pattern of rare and long-lasting concerted evolution in prokaryotes. <i>Communications Biology</i> , 2018, 1, 12.	2.0	16
2444	Renewing Felsenstein's phylogenetic bootstrap in the era of big data. <i>Nature</i> , 2018, 556, 452-456.	13.7	513
2445	Phylogenetic signature of lateral exchange of genes for antibiotic production and resistance among bacteria highlights a pattern of global transmission of pathogens between humans and livestock. <i>Molecular Phylogenetics and Evolution</i> , 2018, 125, 255-264.	1.2	7
2446	Cyanotrophic and arsenic oxidizing activities of <i>Pseudomonas mendocina</i> P6115 isolated from mine tailings containing high cyanide concentration. <i>Archives of Microbiology</i> , 2018, 200, 1037-1048.	1.0	5
2447	Diaporthe from walnut tree ( <i>Juglans regia</i> ) in China, with insight of the <i>Diaporthe eres</i> complex. <i>Mycological Progress</i> , 2018, 17, 841-853.	0.5	34
2448	A New Species of Long-glanded Coralsnake of the Genus <i>Calliophis</i> (Squamata: Elapidae) from Dinagat Island, with Notes on the Biogeography and Species Diversity of Philippine <i>Calliophis</i> and <i>Hemibungarus</i> . <i>Herpetologica</i> , 2018, 74, 89-104.	0.2	10
2449	Two New Species of <i>Ansonia</i> from Thailand (Anura: Bufonidae). <i>Zoological Science</i> , 2018, 35, 39-48.	0.3	8
2450	Continued propagation of the CRF19_cpx variant among HIV-positive MSM patients in Spain. <i>Journal of Antimicrobial Chemotherapy</i> , 2018, 73, 1031-1038.	1.3	12
2451	Comparative analysis of the efficiency of intron-length polymorphism of $\beta$ -tubulin genes and microsatellite loci for flax varieties genotyping. <i>Cytology and Genetics</i> , 2018, 52, 1-10.	0.2	15
2452	Comprehensive Assessment of Global Surface Net Radiation Products and Uncertainty Analysis. <i>Journal of Geophysical Research D: Atmospheres</i> , 2018, 123, 1970-1989.	1.2	49
2453	Rickettsial infection in ticks (Acari: Ixodidae) from reptiles in the Colombian Caribbean. <i>Ticks and Tick-borne Diseases</i> , 2018, 9, 623-628.	1.1	21
2454	Multigene phylogeny and morphology reveal <i>Cytospora spiraeae</i> sp. nov. (Diaporthales, Ascomycota) in China. <i>Phytotaxa</i> , 2018, 338, 49.	0.1	16



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2455	Muscular evolution of hemipenis in Imantodini snakes (Squamata: Dipsadidae). <i>Zoological Journal of the Linnean Society</i> , 2018, 183, 966-980.	1.0	0
2456	Phylogeography of the <i>Oenanthe hispanica-pleschanka-cypriaca</i> complex (Aves, Muscicapidae): Tj ETQq1 1 0.784314 rgBT /Ove data, and morphometric data. <i>Journal of Zoological Systematics and Evolutionary Research</i> , 2018, 56, 408-427.	0.6	13
2457	A tale of two soft-shell clams: an integrative taxonomic analysis confirms <i>Mya japonica</i> as a valid species distinct from <i>Mya arenaria</i> (Bivalvia: Myidae). <i>Zoological Journal of the Linnean Society</i> , 2018, 184, 605-622.	1.0	10
2458	Species diversity in the marine microturbellarian <i>Astrotrorhynchus bifidus sensu lato</i> (Platyhelminthes: Rhabdozoa) from the Northeast Pacific Ocean. <i>Molecular Phylogenetics and Evolution</i> , 2018, 120, 259-273.	1.2	16
2459	Phylogenomics. <i>Methods in Molecular Biology</i> , 2018, 1704, 103-187.	0.4	15
2460	Two new species of <i>Phanerochaete</i> (Basidiomycota) and redescription of <i>P. robusta</i> . <i>Mycological Progress</i> , 2018, 17, 425-435.	0.5	14
2461	Multi-gene phylogenetic analyses reveals <i>Neohelicosporium</i> gen. nov. and five new species of helicosporous hyphomycetes from aquatic habitats. <i>Mycological Progress</i> , 2018, 17, 631-646.	0.5	24
2462	Evolution of Early Thule Material Culture: Cultural Transmission and Terrestrial Ecology. <i>Human Ecology</i> , 2018, 46, 633-650.	0.7	10
2463	Diaporthosporiaceae, a novel family of Diaporthales (Sordariomycetes, Ascomycota). <i>Mycoscience</i> , 2018, 59, 229-235.	0.3	15
2464	Host Specificity in Subarctic Aphids. <i>Environmental Entomology</i> , 2018, 47, 77-86.	0.7	0
2465	Following the cold: geographical differentiation between interglacial refugia and speciation in the arctoalpine species complex <i>Bombus monticola</i> (Hymenoptera: Apidae). <i>Systematic Entomology</i> , 2018, 43, 200-217.	1.7	40
2466	<i>Xanthagaricus thailandensis</i> sp. nov. (Agaricales, Basidiomycota), from northern Thailand. <i>Phytotaxa</i> , 2018, 348, 109.	0.1	4
2467	Skin mites in mice ( <i>Mus musculus</i> ): high prevalence of <i>Myobia</i> sp. (Acari, Arachnida) in Robertsonian mice. <i>Parasitology Research</i> , 2018, 117, 2139-2148.	0.6	5
2468	Genetic diversity studies in endangered desert teak [ <i>Tecomella undulata</i> (Sm) Seem] using arbitrary (RAPD), semi-arbitrary (ISSR) and sequence based (nuclear rDNA) markers. <i>Trees - Structure and Function</i> , 2018, 32, 1083-1101.	0.9	11
2469	Phylogenetic position and taxonomy of <i>Kusaghiporia usambarensis</i> gen. et sp. nov. (Polyporales). <i>Mycology</i> , 2018, 9, 136-144.	2.0	8
2470	Quartet Sampling distinguishes lack of support from conflicting support in the green plant tree of life. <i>American Journal of Botany</i> , 2018, 105, 385-403.	0.8	194
2471	<i>Leucangium carthusianum</i> var. <i>purpureum</i> , a new purple truffle from China. <i>Phytotaxa</i> , 2018, 347, 165.	0.1	0
2472	Historical biogeography of Loranthaceae (Santalales): Diversification agrees with emergence of tropical forests and radiation of songbirds. <i>Molecular Phylogenetics and Evolution</i> , 2018, 124, 199-212.	1.2	47

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2473	Recent radiation and dispersal of an ancient lineage: The case of <i>Fouquieria</i> (Fouquieriaceae, Ericales) in North American deserts. <i>Molecular Phylogenetics and Evolution</i> , 2018, 126, 92-104.	1.2	19
2474	A review of the genus <i>Berthellina</i> (Mollusca: Heterobranchia: Pleurobranchidae) in the Red Sea based on molecular and anatomical data. <i>Marine Biodiversity</i> , 2018, 48, 1121-1134.	0.3	2
2475	Mitogenomics phylogenetic relationships of the current slothâ€™s genera and species (Bradyrodidae and) Tj ETQq0 0 0 rgBT /Overlock 0.7	0.7	9
2476	Molecular phylogeny of <i>Glossodoris</i> (Ehrenberg, 1831) nudibranchs and related genera reveals cryptic and pseudocryptic species complexes. <i>Cladistics</i> , 2018, 34, 41-56.	1.5	12
2477	Phylogeny and historical biogeography of silky lacewings (<sc>N</sc>europtera:) Tj ETQq0 0 0 rgBT /Overlock 1.7 If 50 582 Td (<sc>	1.7	5
2478	Molecular identification of species and production origins of edible bird's nest using FINS and SYBR green I based real-time PCR. <i>Food Control</i> , 2018, 84, 118-127.	2.8	29
2479	UFBoot2: Improving the Ultrafast Bootstrap Approximation. <i>Molecular Biology and Evolution</i> , 2018, 35, 518-522.	3.5	5,798
2480	Diversity of Diatrypaceae Species Associated with Dieback of Grapevines in South Africa, with the Description of <i>Eutypa cremea</i> sp. nov.. <i>Plant Disease</i> , 2018, 102, 220-230.	0.7	26
2481	First Record of <i>Tylosurus crocodilus</i> (Péron & Lesueur, 1821) (Beloniformes: Belonidae) from Odisha Coast, Bay of Bengal, India: Exploration of a Biological Invasion Using DNA Barcoding. <i>Thalassas</i> , 2018, 34, 209-217.	0.1	4
2482	<i>Purpureocorticium microsporum</i> (Basidiomycota) gen. et sp. nov. from East Asia. <i>Mycological Progress</i> , 2018, 17, 357-364.	0.5	3
2483	Species delimitation and biogeography of the Ryukyu ground geckos, <i>Goniurosaurus kuroiwa</i> ssp. (Squamata: Eublepharidae), by use of mitochondrial and nuclear DNA analyses. <i>Journal of Zoological Systematics and Evolutionary Research</i> , 2018, 56, 209-222.	0.6	7
2484	Subspecies dynamics in space and time: A study of the red deer complex using ancient and modern <sc>DNA</sc> and morphology. <i>Journal of Biogeography</i> , 2018, 45, 367-380.	1.4	30
2485	Global variation in the cost of increasing ecosystem carbon. <i>Nature Climate Change</i> , 2018, 8, 38-42.	8.1	10
2486	Cryptic diversity in <i>Rhampholeon boulengeri</i> (Sauria: Chamaeleonidae), a pygmy chameleon from the Albertine Rift biodiversity hotspot. <i>Molecular Phylogenetics and Evolution</i> , 2018, 122, 125-141.	1.2	17
2487	Simplified and efficient DNA extraction protocol for Meliolaceae specimens. <i>Mycological Progress</i> , 2018, 17, 403-415.	0.5	10
2488	Morphology, cell-division, and phylogeny of <i>Schmidingerothrix elongata</i> spec. nov. (Ciliophora,) Tj ETQq1 1 0.784314 rgBT /Overlock 0.5 30 Protistology, 2018, 62, 24-42.	0.5	30
2489	Genetic Diversity and Structure through Three Cycles of a <i>Eucalyptus urophylla</i> S.T.Blake Breeding Program. <i>Forests</i> , 2018, 9, 372.	0.9	10
2490	Using DNA Markers in Studies of Chigger Mites (Acariformes, Trombiculidae). <i>Entomological Review</i> , 2018, 98, 1351-1368.	0.1	2

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2491	Complete mitochondrial genome of the nesting Colombian Caribbean loggerhead turtle: first approach of tRNAs and phylogenetic analysis. <i>Universitas Scientiarum</i> , 2018, 23, 355-381.	0.2	3
2492	<i>Archaeomyiodon sampedinensis</i> , gen. et sp. nov., a new mylodontine from the middle Pleistocene of Pampean Region, Argentina. <i>Journal of Vertebrate Paleontology</i> , 2018, 38, e1542308.	0.4	5
2493	Challenges and changes in the parenting experiences of Korean immigrants in New Zealand. <i>Asian and Pacific Migration Journal</i> , 2018, 27, 431-450.	0.5	4
2494	Phylogenetic Analysis: Basic Concepts and Its Use as a Tool for Virology and Molecular Epidemiology. <i>Acta Scientiae Veterinariae</i> , 2018, 44, 20.	0.2	0
2495	The first set of universal nuclear protein-coding loci markers for avian phylogenetic and population genetic studies. <i>Scientific Reports</i> , 2018, 8, 15723.	1.6	16
2496	<i>Teuwoa saxicola</i> and <i>T. alpina</i> spp. nov. and the genus in China. <i>Mycotaxon</i> , 2018, 133, 79-87.	0.1	1
2497	A new species and new record of <i>Lophiotrema</i> (Lophiotremataceae, Dothideomycetes) from karst landforms in southwest China. <i>Phytotaxa</i> , 2018, 379, 169.	0.1	5
2498	<i>Cytospora piceae</i> sp. nov. associated with canker disease of <i>Picea crassifolia</i> in China. <i>Phytotaxa</i> , 2018, 383, 181.	0.1	8
2499	Description of <i>Dermea persica</i> (Dermateaceae, Helotiales), a new asexual Ascomycete from Iran, and an updated key to <i>Dermea</i> species. <i>Phytotaxa</i> , 2018, 367, 25.	0.1	4
2500	<i>Trichoderma polyalthiae</i> sp. nov., an endophytic fungus from <i>Polyalthia debilis</i> . <i>Phytotaxa</i> , 2018, 371, 273.	0.1	2
2501	A New Species of <i>Enyalius</i> (Squamata, Leiosauridae) Endemic to the Brazilian Cerrado. <i>Herpetologica</i> , 2018, 74, 355-369.	0.2	6
2502	A new endophytic fungus, <i>Tulasnella phuhinrongklaensis</i> (Cantharellales, Basidiomycota) isolated from roots of the terrestrial orchid, <i>Phalaenopsis pulcherrima</i> . <i>Phytotaxa</i> , 2018, 374, 99.	0.1	9
2503	Two new species of <i>Barssia</i> from China. <i>Phytotaxa</i> , 2018, 374, 129.	0.1	2
2504	<i>Triadelphia fusiformis</i> sp. nov. from a freshwater habitat in Thailand. <i>Phytotaxa</i> , 2018, 374, 231.	0.1	4
2505	Species delimitation of <i>Stemona</i> (Stemonaceae) based on sequences of five plastid DNA regions. <i>Phytotaxa</i> , 2018, 374, 291.	0.1	1
2506	A taxonomic reassessment of Tubeufiales based on multi-locus phylogeny and morphology. <i>Fungal Diversity</i> , 2018, 92, 131-344.	4.7	49
2507	Two new species of <i>Barssia</i> from China. <i>Phytotaxa</i> , 2018, 374, 129.	0.1	2
2508	<i>Neopestalotiopsis rosicola</i> sp. nov. causing stem canker of <i>Rosa chinensis</i> in China. <i>Mycotaxon</i> , 2018, 133, 271-283.	0.1	14

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2509	The challenges and potential utility of phenotypic specimen-level phylogeny based on maximum parsimony. <i>Earth and Environmental Science Transactions of the Royal Society of Edinburgh</i> , 2018, 109, 301-323.	0.3	11
2510	Diversity of Mesopelagic Fishes in the Southern Ocean - A Phylogeographic Perspective Using DNA Barcoding. <i>Frontiers in Ecology and Evolution</i> , 2018, 6, .	1.1	23
2511	A new species of Parafimbrios from northern Vietnam (Squamata: Xenodermatidae). <i>Zootaxa</i> , 2018, 4527, 269.	0.2	7
2512	An Introduction to the Systematics of Small-Bodied Neacomys (Rodentia: Cricetidae) from Peru with Descriptions of Two New Species. <i>American Museum Novitates</i> , 2018, 3913, 1-38.	0.2	12
2513	How Many Species, Taxa, or Lineages of <i>Cebus albifrons</i> (Platyrrhini, Primates) Inhabit Ecuador? Insights from Mitogenomics. <i>International Journal of Primatology</i> , 2018, 39, 1068-1104.	0.9	4
2514	Divergence of Fecal Microbiota and Their Associations With Host Phylogeny in Cervinae. <i>Frontiers in Microbiology</i> , 2018, 9, 1823.	1.5	9
2515	Building (Viral) Phylogenetic Trees Using a Maximum Likelihood Approach. <i>Current Protocols in Microbiology</i> , 2018, 51, e63.	6.5	5
2516	Phylogenetic approaches resolve taxonomical confusion in <i>Pedicularis</i> (Orobanchaceae): Reinstatement of <i>Pedicularis delavayi</i> and discovering a new species <i>Pedicularis milliana</i> . <i>PLoS ONE</i> , 2018, 13, e0200372.	1.1	4
2517	Phylogeography of the gall-inducing micromoth <i>Eucecidoses minutanus</i> BrÅ'thes (Cecidosidae) reveals lineage diversification associated with the Neotropical Peripampasic Orogenic Arc. <i>PLoS ONE</i> , 2018, 13, e0201251.	1.1	8
2518	Associating larvae and adults of the Neotropical caddisfly genus <i>Synoestropsis</i> Ulmer (Trichoptera: Tj ETQq1 1 0.784314 rgBT /Overlock 10 TF 277, 169-189.	0.4	3
2519	Chinese black truffles: <i>Tuber yigongense</i> sp. nov., taxonomic reassessment of <i>T. indicum</i> s.l., and re-examination of the <i>T. sinense</i> isotype. <i>Mycotaxon</i> , 2018, 133, 183-196.	0.1	1
2520	Equine Transport and Changes in Equid Herpesvirus' Status. <i>Frontiers in Veterinary Science</i> , 2018, 5, 224.	0.9	17
2521	Genome Sequence of Peacock Reveals the Peculiar Case of a Glittering Bird. <i>Frontiers in Genetics</i> , 2018, 9, 392.	1.1	32
2522	A newly recognised species of <i>Cryptes</i> Maskell 1892 (Hemiptera: Coccidae) from Western Australia. <i>Zootaxa</i> , 2018, 4508, 101.	0.2	1
2523	Two new and potentially highly threatened <i>Megophrys</i> Horned frogs (Amphibia: Megophryidae) from Indochina's highest mountains. <i>Zootaxa</i> , 2018, 4508, 301.	0.2	27
2524	Testing the validity of two putative sympatric species from <i>Sinocyclocheilus</i> (Cypriniformes: Tj ETQq1 1 0.784314 rgBT /Overlock 10 TF 0.2	0.2	9
2525	Transmission Dynamics of Highly Pathogenic Avian Influenza Virus A(H5Nx) Clade 2.3.4.4, North America, 2014-2015. <i>Emerging Infectious Diseases</i> , 2018, 24, 1840-1848.	2.0	41
2526	Colonization and divergence: phylogeography and population genetics of the Atlantic coast beach mice. <i>Systematics and Biodiversity</i> , 2018, 16, 757-773.	0.5	3

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2528	A. W.ÂF. Edwards and the Origin of Bayesian Phylogenetics. , 0, , 352-362.		0
2529	Four new <i>Tuber</i> species added to the <i>Rufum</i> group from China based on morphological and molecular evidence. Mycologia, 2018, 110, 771-779.	0.8	4
2530	<i>Acuminatispora palmarum</i> gen. et sp. nov. from mangrove habitats. Mycological Progress, 2018, 17, 1173-1188.	0.5	8
2531	Integrative species delimitation in practice: Revealing cryptic lineages within the short-nosed skink <i>Plestiodon brevirostris</i> (Squamata: Scincidae). Molecular Phylogenetics and Evolution, 2018, 129, 242-257.	1.2	17
2532	Phylogenetic relationships in Malesian Pacific <i>Piper</i> (Piperaceae) and their implications for systematics. Taxon, 2018, 67, 693-724.	0.4	15
2533	Phylogeny and morphology reveal two new species of <i>Diaporthe</i> from Traditional Chinese Medicine in Northeast China. Phytotaxa, 2018, 336, 159.	0.1	19
2534	Hemogregarine and Rickettsial infection in ticks of toads from northeastern Colombia. International Journal for Parasitology: Parasites and Wildlife, 2018, 7, 237-242.	0.6	12
2535	A new species of <i>Leptolalax</i> (Anura: Megophryidae) from Son Tra Peninsula, central Vietnam. Zootaxa, 2018, 4388, 1-21.	0.2	25
2536	Manganese (Mn <sup>2+</sup> ) tolerance and biosorption by <i>Meyerozyma guilliermondii</i> and <i>Meyerozyma caribbica</i> strains. Journal of Environmental Chemical Engineering, 2018, 6, 4538-4545.	3.3	16
2537	Reassessment of the taxonomic status of <i>Craseomys</i> and three controversial species of <i>Myodes</i> and <i>Alticola</i> (Rodentia: Arvicolinae). Zootaxa, 2018, 4429, 1-52.	0.2	5
2538	A new cryptic species of <i>Oreobates</i> (Anura: Craugastoridae) from the seasonally dry tropical forest of central Brazil. Zootaxa, 2018, 4441, 89.	0.2	4
2539	Diversity of fungal latent pathogens and true endophytes associated with fruit trees in Uruguay. Journal of Phytopathology, 2018, 166, 633-647.	0.5	32
2540	An updated phylogeny and morphological study of the <i>Phymaturus vociferator</i> clade (Iguania: Tj ETQq1 1 0.784314 rgBT /Overlock 10	0.2	5
2541	A Phylogeny of the Genus <i>Amaranthus</i> (Amaranthaceae) Based on Several Low-Copy Nuclear Loci and Chloroplast Regions. Systematic Botany, 2018, 43, 439-458.	0.2	38
2542	<i>Translucidityrium thailandicum</i> gen. et sp. nov.: a new genus in Phaeothecoidiaceae. Mycological Progress, 2018, 17, 1087-1096.	0.5	6
2543	On the distinctiveness of <i>Amapasaurus</i> , its relationship with <i>Loxopholis</i> Cope 1869, and description of a new genus for <i>L. guianensis</i> and <i>L. hoogmoedi</i> (Gymnophthalmoidea/Ecpleopodini: Squamata). Zootaxa, 2018, 4441, 332-346.	0.2	10
2544	Morphological and molecular evidence support a new endophytic fungus, <i>Chaetomella endophytica</i> from Japan. Mycoscience, 2018, 59, 473-478.	0.3	3

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2545	The Hemiparasitic Plant <i>Phtheirospermum</i> (Orobanchaceae) Is Polyphyletic and Contains Cryptic Species in the Hengduan Mountains of Southwest China. <i>Frontiers in Plant Science</i> , 2018, 9, 142.	1.7	38
2546	Chimeric Structure of Plant Malic Enzyme Family: Different Evolutionary Scenarios for NAD- and NADP-Dependent Isoforms. <i>Frontiers in Plant Science</i> , 2018, 9, 565.	1.7	22
2547	<i>Phaeoclavulina pseudozippelii</i> sp. nov. (Gomphales, Basidiomycota) from Northern Thailand. <i>Phytotaxa</i> , 2018, 362, 211.	0.1	1
2548	<i>Diaporthe</i> diversity and pathogenicity revealed from a broad survey of grapevine diseases in Europe. <i>Persoonia: Molecular Phylogeny and Evolution of Fungi</i> , 2018, 40, 135-153.	1.6	107
2549	Divergent evolution and clade-specific duplications of the Insulin-like Receptor in malacostracan crustaceans. <i>General and Comparative Endocrinology</i> , 2018, 268, 34-39.	0.8	9
2550	Phylogeny and evolution of the genus <i>Ctenocolum</i> Kingsolver & Whitehead (Coleoptera, Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 Evolution, 2018, 50, 1-35.	0.2	0
2551	Comparative genomics and the nature of placozoan species. <i>PLoS Biology</i> , 2018, 16, e2005359.	2.6	73
2552	Transfer of <i>Senecio karelinioides</i> (Asteraceae~Senecioneae) to <i>Synotis</i> based on evidence from morphology, karyology and ITS/ETS sequence data. <i>Nordic Journal of Botany</i> , 2018, 36, e01838.	0.2	2
2553	DNA analysis reveals rich diversity of <i>Hydnotrya</i> with emphasis on the species found in China. <i>Mycological Progress</i> , 2018, 17, 1123-1137.	0.5	2
2554	Genomic Comparison Among Global Isolates of <i>L. interrogans</i> Serovars Copenhageni and <i>Icterohaemorrhagiae</i> Identified Natural Genetic Variation Caused by an Indel. <i>Frontiers in Cellular and Infection Microbiology</i> , 2018, 8, 193.	1.8	39
2555	Effect of Oxygen on Verbenone Conversion From cis-Verbenol by Gut Facultative Anaerobes of <i>Dendroctonus valens</i> . <i>Frontiers in Microbiology</i> , 2018, 9, 464.	1.5	14
2556	The complete mitochondrial genome of <i>Clostera anastomosis</i> (Lepidoptera: Notodontidae) and implication for the phylogenetic relationships of Noctuoidea species. <i>International Journal of Biological Macromolecules</i> , 2018, 118, 1574-1583.	3.6	22
2557	Phylogeography and population genomics of a lotic water beetle across a complex tropical landscape. <i>Molecular Ecology</i> , 2018, 27, 3346-3356.	2.0	12
2558	Fossils know it best: Using a new set of fossil calibrations to improve the temporal phylogenetic framework of murid rodents (Rodentia: Muridae). <i>Molecular Phylogenetics and Evolution</i> , 2018, 128, 98-111.	1.2	61
2559	Combining complete chloroplast genome sequences with target loci data and morphology to resolve species limits in <i>Triplostegia</i> (Caprifoliaceae). <i>Molecular Phylogenetics and Evolution</i> , 2018, 129, 15-26.	1.2	40
2560	Exploring data processing strategies in NGS target enrichment to disentangle radiations in the tribe <i>Cardueae</i> (Compositae). <i>Molecular Phylogenetics and Evolution</i> , 2018, 128, 69-87.	1.2	38
2561	MPBoot: fast phylogenetic maximum parsimony tree inference and bootstrap approximation. <i>BMC Evolutionary Biology</i> , 2018, 18, 11.	3.2	129
2562	Detection of ascaridoid nematode parasites in the important marine food-fish <i>Conger myriaster</i> (Brevoort) (Anguilliformes: Congridae) from the Zhoushan Fishery, China. <i>Parasites and Vectors</i> , 2018, 11, 274.	1.0	22

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2563	Morphological and mitochondrial DNA data reshuffle the taxonomy of the genera <i>Atopochetus</i> Attems, <i>Litostrophus</i> Chamberlin and <i>Tonkinbolus</i> Verhoeff (Diplopoda: Spirobolida: Pachybolidae), with descriptions of nine new species. <i>Invertebrate Systematics</i> , 2018, 32, 159.	0.5	12
2564	Nectar protection in arid-adapted flowers of Zygothylaceae-Zygothylloideae. <i>Perspectives in Plant Ecology, Evolution and Systematics</i> , 2018, 34, 37-50.	1.1	3
2565	Families and genera of diaporthean fungi associated with canker and dieback of tree hosts. <i>Persoonia: Molecular Phylogeny and Evolution of Fungi</i> , 2018, 40, 119-134.	1.6	57
2566	Phylogeny of <i>Spiraea</i> (Rosaceae) based on plastid and nuclear molecular data: Implications for morphological character evolution and systematics. <i>Perspectives in Plant Ecology, Evolution and Systematics</i> , 2018, 34, 109-119.	1.1	12
2567	Complete genome sequence analysis of rare G4P[6] rotavirus strains from human and pig reveals the evidence for interspecies transmission. <i>Infection, Genetics and Evolution</i> , 2018, 65, 357-368.	1.0	13
2568	Convex hull analysis of evolutionary and phylogenetic relationships between biological groups. <i>Journal of Theoretical Biology</i> , 2018, 456, 34-40.	0.8	19
2569	Parahelicops, Pararhabdophis, Paraphyly: Phylogenetic Relationships among Certain Southeast Asian Natricine Snakes (Hebius). <i>American Museum Novitates</i> , 2018, 3906, 1-7.	0.2	10
2570	Phragmidium species parasitizing species of Rosaceae in Tibet, China, with descriptions of three new species. <i>Mycological Progress</i> , 2018, 17, 967-988.	0.5	7
2571	Using mating-type loci to improve taxonomy of the <i>Tuber indicum</i> complex, and discovery of a new species, <i>T. longispinosum</i> . <i>PLoS ONE</i> , 2018, 13, e0193745.	1.1	13
2572	Bootstrap and Rogue Identification Tests for Phylogenetic Analyses. <i>Molecular Biology and Evolution</i> , 2018, 35, 2327-2333.	3.5	31
2573	Biogeographic patterns and diversification dynamics of the genus <i>Cardiodactylus</i> Saussure (Orthoptera, Grylloidea, Eneopterinae) in Southeast Asia. <i>Molecular Phylogenetics and Evolution</i> , 2018, 129, 1-14.	1.2	22
2574	<i>Spissiomycetes endophytica</i> (Dothideomycetes, Ascomycota), a new endophytic fungus from Thailand. <i>Phytotaxa</i> , 2018, 333, 219.	0.1	2
2575	An appendage-bearing coelomycete <i>Pseudotruncatella arezzoensis</i> gen. and sp. nov. (Amphisphaeriales) <i>Tj ETQq0 0,0 rgBT /Qverlock 10</i>	0.1	5
2576	Coleopuccinia in China and its relationship to <i>Gymnosporangium</i> . <i>Phytotaxa</i> , 2018, 347, 235.	0.1	5
2577	Studies of botryosphaeralean fungi associated with canker and dieback of tree hosts in Dongling Mountain of China. <i>Phytotaxa</i> , 2018, 348, 63.	0.1	20
2578	DNA barcodes uncover hidden taxonomic diversity behind the variable wing patterns in the Neotropical butterfly genus <i>Zaretis</i> (Lepidoptera: Nymphalidae: Charaxinae). <i>Zoological Journal of the Linnean Society</i> , 2019, 185, 132-192.	1.0	7
2579	Information from the mitochondrial genomes of two egg parasitoids, <i>Gonatocerus</i> sp. and <i>Telenomus</i> sp., reveals a controversial phylogenetic relationship between Mymaridae and Scelionidae. <i>Genomics</i> , 2019, 111, 1059-1065.	1.3	11
2580	In vitro establishment of shoot meristems of <i>Ilex paraguariensis</i> and identification of endophytic bacteria. <i>Journal of Forestry Research</i> , 2019, 30, 1765-1777.	1.7	3

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2581	Tree Evaluation and Robustness Testing. , 2019, , 736-745.		0
2582	The Munduruku marmoset: a new monkey species from southern Amazonia. PeerJ, 2019, 7, e7019.	0.9	19
2583	Phylogenetic relationships within the flatworm genus Matuxia (Platyhelminthes, Tricladida,) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 667 T genus. Organisms Diversity and Evolution, 2019, 19, 377-390.	0.7	0
2584	Improvement of time forecasting models using a novel hybridization of bootstrap and double bootstrap artificial neural networks. Applied Soft Computing Journal, 2019, 84, 105676.	4.1	18
2585	Phylogenetic relationships within the genus <i>Hypnea</i> (Cystocloniaceae, Rhodophyta): convergent evolution and its implications in the infrageneric classification. Botanica Marina, 2019, 62, 563-575.	0.6	5
2586	<i>Conioscypha tenebrosa</i> sp. nov. (Conioscyphaceae) from China and notes on <i>Conioscypha</i> species. Phytotaxa, 2019, 413, 159-171.	0.1	5
2587	A new species of <i>Acanthosaura</i> Gray 1831 (Reptilia: Agamidae) from central Vietnam. Zootaxa, 2019, 4612, 555.	0.2	3
2588	A cytochrome P450 monooxygenase gene required for biosynthesis of the trichothecene toxin harzianum A in <i>Trichoderma</i> . Applied Microbiology and Biotechnology, 2019, 103, 8087-8103.	1.7	13
2589	A Systematist's Guide to Estimating Bayesian Phylogenies From Morphological Data. Insect Systematics and Diversity, 2019, 3, 2.	0.7	28
2590	Mitochondrial genome reorganization provides insights into the relationship between oribatid mites and astigmatid mites (Acari: Sarcoptiformes: Oribatida). Zoological Journal of the Linnean Society, 2019, 187, 585-598.	1.0	17
2591	The Complete Mitochondrial Genome of <i>Platysternon megacephalum peguense</i> and Molecular Phylogenetic Analysis. Genes, 2019, 10, 487.	1.0	11
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2595	A molecular phylogeny of the Indo-West Pacific species of <i>Haloa</i> sensu lato gastropods (Cephalaspidea: Haminoeidae): Tethyan vicariance, generic diversity, and ecological specialization. Molecular Phylogenetics and Evolution, 2019, 139, 106557.	1.2	19
2596	Canker and Wood Rot Pathogens Present in Young Apple Trees and Propagation Material in the Western Cape of South Africa. Plant Disease, 2019, 103, 3129-3141.	0.7	20
2597	Form and relationship of the social networks of the New Testament. Social Network Analysis and Mining, 2019, 9, 1.	1.9	4
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2600	Phylogenetic relationships in the cricket tribe Xenogryllini (Orthoptera, Gryllidae, Eneopterinae) and description of the Indian genus <i>Indigryllus</i> gen. nov.. Journal of Zoological Systematics and Evolutionary Research, 2019, 57, 789-805.	0.6	4
2601	Mini DNA-barcode as molecular marker for heavily processed hairtail fish products authentication. IOP Conference Series: Earth and Environmental Science, 2019, 278, 012001.	0.2	2
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2603	Mitochondrial genome reorganization characterizes various lineages of mesostigmatid mites (Acari: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5	0.7	24
2604	<i>Cytospora elaeagnicola</i> sp. nov. Associated with Narrow-leaved Oleaster Canker Disease in China. Mycobiology, 2019, 47, 319-328.	0.6	7
2605	Three new species of the snake genus <i>Achalinus</i> from Vietnam (Squamata: Xenodermatidae). Zootaxa, 2019, 4590, zootaxa.4590.2.3.	0.2	16
2606	Phylogeny, origin and dispersal of <i>Saussurea</i> (Asteraceae) based on chloroplast genome data. Molecular Phylogenetics and Evolution, 2019, 141, 106613.	1.2	39
2607	Freshwater Sordariomycetes. Fungal Diversity, 2019, 99, 451-660.	4.7	119
2608	Botryosphaeralean fungi causing canker and dieback of tree hosts from Mount Yudu in China. Mycological Progress, 2019, 18, 1341-1361.	0.5	13
2609	Species Diversity in <i>Colletotrichum</i> Causing Anthracnose of Aromatic and Ornamental Lamiaceae in Italy. Agronomy, 2019, 9, 613.	1.3	28
2610	Crustose Calicioid Lichens and Fungi in Mountain Cloud Forests of Tanzania. Microorganisms, 2019, 7, 491.	1.6	3
2611	New species of <i>Septoria</i> associated with leaf spot diseases in Iran. Mycologia, 2019, 111, 1056-1071.	0.8	9
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2613	<i>Baeolidia moebii</i> Bergh, 1888 (Mollusca: Gastropoda: Nudibranchia) is spreading in the eastern Mediterranean Sea. Regional Studies in Marine Science, 2019, 32, 100830.	0.4	5
2614	<i>Ganoderma shanxiense</i> , a new species from northern China based on morphological and molecular evidence. Phytotaxa, 2019, 406, 129-136.	0.1	9
2615	TreeCluster: Clustering biological sequences using phylogenetic trees. PLoS ONE, 2019, 14, e0221068.	1.1	99
2616	Integrative taxonomy of an arctic bumblebee species complex highlights a new cryptic species (Apidae: Tj ETQq1 1,0,784314 rgBT /Ove	1.0	23

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2617	A new species of <i>Opisthotropis</i> from northern Vietnam previously misidentified as the Yellow-spotted Mountain Stream Keelback <i>O. maculosa</i> Stuart & Chuaynkern, 2007 (Squamata: Natricidae). <i>Zootaxa</i> , 2019, 4613, 578.	0.2	3
2618	Two new species of <i>Eutypella</i> and a new combination in the genus <i>Peroneutypa</i> (Diatrypaceae). <i>Mycological Progress</i> , 2019, 18, 1057-1069.	0.5	12
2619	Diversity of <i>Empruthotrema</i> Johnston and Tiegs, 1992 parasitizing batoids (Chondrichthyes: Rajiformes) Tj ETQq0 0 0 rgBT /Overlock 10 Parasitology Research, 2019, 118, 3113-3127.	0.6	5
2620	Foliar pathogens of eucalypts. <i>Studies in Mycology</i> , 2019, 94, 125-298.	4.5	66
2621	Two novel species of <i>Marasmius</i> (Marasmiaceae, Agaricales) from lower northern Thailand. <i>Phytotaxa</i> , 2019, 403, 111.	0.1	1
2622	&lt;p&gt;&lt;strong&gt;&lt;em&gt;Tremateia murispora sp. nov.&lt;/em&gt;&lt;/strong&gt;&lt;strong&gt; (&lt;em&gt;Didymosphaeriaceae&lt;/em&gt;, Pleosporales) from Guizhou, China&lt;/strong&gt;&lt;p&gt;. <i>Phytotaxa</i> , 2019, 416, 79-87.	0.1	8
2623	Additions to the genus <i>Savoryella</i> (Savoryellaceae), with the asexual morphs <i>Savoryella nypae</i> comb. nov. and <i>S. sarushimana</i> sp. nov.. <i>Phytotaxa</i> , 2019, 408, 195-207.	0.1	11
2624	Two new species of <i>Pachyphlodes</i> from China. <i>Phytotaxa</i> , 2019, 411, 105-115.	0.1	1
2625	Present-Day Distribution and Potential Spread of the Invasive Green Alga <i>Avrainvillea amadelpha</i> Around the Main Hawaiian Islands. <i>Frontiers in Marine Science</i> , 2019, 6, .	1.2	8
2626	Molecular identification of mosquitoes of the <i>Anopheles maculatus</i> group of subgenus <i>Cellia</i> (Diptera: Culicidae) in the Indonesian Archipelago. <i>Acta Tropica</i> , 2019, 199, 105124.	0.9	8
2627	New species, diversity, systematics, and conservation assessment of the Puppet Toads of Sumatra (Anura: Bufonidae: <i>Sigalegalephrynus</i> ). <i>Zootaxa</i> , 2019, 4679, zootaxa.4679.2.9.	0.2	6
2628	&lt;p&gt;&lt;strong&gt;Multigene phylogenetic characterisation of&lt;em&gt; Colletotrichum artocarpicola sp. nov.&lt;/em&gt; from &lt;em&gt;Artocarpus heterophyllus&lt;/em&gt; in northern Thailand&lt;/strong&gt;&lt;p&gt;. <i>Phytotaxa</i> , 2019, 418, 273-286.	0.1	11
2629	Unexpected phylogenetic relationships within the world's largest limbless skink species ( <i>Acontias</i> ) Tj ETQq0 0 0 rgBT /Overlock 10 T Journal of Zoological Systematics and Evolutionary Research, 2019, 57, 445-460.	0.6	3
2630	Contrasting morphology with population genetics approach: An insight to revision of the Neotropical annual fish <i>Austrolebias robustus</i> species group based on a taxonomic integrative framework. <i>Journal of Zoological Systematics and Evolutionary Research</i> , 2019, 57, 591-605.	0.6	4
2631	Taxonomic revision of the dolphin genus <i>Lagenorhynchus</i> . <i>Marine Mammal Science</i> , 2019, 35, 957-1057.	0.9	22
2632	Not always young: The first vertebrate ancient origin of true parthenogenesis found in an Amazon leaf litter lizard with evidence of mitochondrial haplotypes surfing on the wave of a range expansion. <i>Molecular Phylogenetics and Evolution</i> , 2019, 135, 105-122.	1.2	17
2633	<i>Lasiodiplodia theobromae</i> and <i>L. pseudotheobromae</i> causing leaf necrosis on <i>Camellia sinensis</i> in Fujian Province, China. <i>Canadian Journal of Plant Pathology</i> , 2019, 41, 277-284.	0.8	7
2634	Identification assisted by molecular markers of larval parasites in two limpet species (Patellogastropoda: <i>Nacella</i> ) inhabiting Antarctic and Magellan coastal systems. <i>Polar Biology</i> , 2019, 42, 1175-1182.	0.5	7

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2636	Evolution of the Piscine orthoreovirus Genome Linked to Emergence of Heart and Skeletal Muscle Inflammation in Farmed Atlantic Salmon ( <i>Salmo salar</i> ). Viruses, 2019, 11, 465.	1.5	24
2637	The pioneer lichen Placopsis in maritime Antarctica: Genetic diversity of their mycobionts and green algal symbionts, and their correlation with deglaciation time. Symbiosis, 2019, 79, 1-24.	1.2	19
2638	Advanced understanding of phylogenetic relationships, morphological evolution and biogeographic history of the mega-diverse plant genus Myrcia and its relatives (Myrtaceae: Myrteae). Molecular Phylogenetics and Evolution, 2019, 138, 65-88.	1.2	24
2639	Identification of Caragana arborescens shoot blight disease caused by Phaeobotryon caraganae sp. nov. (Botryosphaerales) in China. European Journal of Plant Pathology, 2019, 155, 537-544.	0.8	1
2640	Relict distribution of <i>Microhyla</i> (Amphibia: Microhylidae) in the Ryukyu Archipelago: High diversity in East Asia maintained by insularization. Zoologica Scripta, 2019, 48, 440-453.	0.7	12
2641	Isolation and characterization of marine bacteria from East Coast of India: functional screening for salt stress tolerance. Heliyon, 2019, 5, e01869.	1.4	14
2642	Optimization and characterization of red pigment production from an endophytic fungus, Nigrospora aurantiaca CMU-ZY2045, and its potential source of natural dye for use in textile dyeing. Applied Microbiology and Biotechnology, 2019, 103, 6973-6987.	1.7	24
2643	Untangling a mess of worms: Species delimitations reveal morphological crypsis and variability in Southeast Asian semi-aquatic earthworms (Almidae, Glyphidrilus). Molecular Phylogenetics and Evolution, 2019, 139, 106531.	1.2	12
2644	Characterization of antifungal metabolites produced by novel lactic acid bacterium and their potential application as food biopreservatives. Annals of Agricultural Sciences, 2019, 64, 71-78.	1.1	93
2645	The little Aplysia coming of age: from one species to a complex of species complexes in Aplysia parvula (Mollusca: Gastropoda: Heterobranchia). Zoological Journal of the Linnean Society, 2019, 187, 279-330.	1.0	25
2646	The changing spectrum of Saccharomycotina yeasts causing candidemia: phylogeny mirrors antifungal susceptibility patterns for azole drugs and amphotericin B. FEMS Yeast Research, 2019, 19, .	1.1	30
2647	New insights into the tetrameric family of the Fur metalloregulators. BioMetals, 2019, 32, 501-519.	1.8	14
2648	An extensive molecular phylogeny of weaverbirds (Aves: Ploceidae) unveils broad nonmonophyly of traditional genera and new relationships. Auk, 2019, 136, .	0.7	6
2649	Distribution and asymptotic behavior of the phylogenetic transfer distance. Journal of Mathematical Biology, 2019, 79, 485-508.	0.8	2
2650	Partitioning of morphospace in larval and adult reed frogs (Anura: Hyperoliidae: Hyperolius) of the Central African Albertine Rift. Zoologischer Anzeiger, 2019, 280, 65-77.	0.4	11
2651	Payments for adding ecosystem carbon are mostly beneficial to biodiversity. Environmental Research Letters, 2019, 14, 054001.	2.2	2
2652	What is really out there? Review of the genus Okenia Menke, 1830 (Nudibranchia: Goniadorididae) in the Mediterranean Sea with description of two new species. PLoS ONE, 2019, 14, e0215037.	1.1	19

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2653	Low-copy nuclear sequence data confirm complex patterns of farina evolution in notholaenid ferns (Pteridaceae). <i>Molecular Phylogenetics and Evolution</i> , 2019, 138, 139-155.	1.2	14
2654	3RAD-based systematics of the transitional Nearctic-Neotropical lubber grasshopper genus <i>Taeniopoda</i> (Orthoptera: Romaleidae). <i>Molecular Phylogenetics and Evolution</i> , 2019, 137, 64-75.	1.2	7
2655	Nuclear and plastid DNA phylogeny of tribe Cardueae (Compositae) with Hyb-Seq data: A new subtribal classification and a temporal diversification framework. <i>Molecular Phylogenetics and Evolution</i> , 2019, 137, 313-332.	1.2	58
2656	<i>Kevinhydea brevistipitata</i> gen. et sp. nov. and <i>Helicoma hydei</i> sp. nov., (Tubeufiaceae) from decaying wood habitats. <i>Mycological Progress</i> , 2019, 18, 671-682.	0.5	7
2657	<i>Neoastrisphaeriella aquatica</i> sp. nov. (Aigialaceae), a new species from freshwater habitat in southern Thailand. <i>Phytotaxa</i> , 2019, 391, 197.	0.1	6
2658	Molecular detection of rabies virus strain with N-gene that clustered with China lineage 2 co-circulating with Africa lineages in Monrovia, Liberia: first reported case in Africa. <i>Epidemiology and Infection</i> , 2019, 147, e85.	1.0	11
2659	The mystery of the origins of <i>Cebus albifrons malitiosus</i> and <i>Cebus albifrons hypoleucus</i> : mitogenomics and microsatellite analyses revealed an amazing evolutionary history of the Northern Colombian white-fronted capuchins. <i>Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis</i> , 2019, 30, 525-547.	0.7	2
2660	A new species of the genus <i>Hilethera</i> Uvarov, 1923 (Orthoptera: Acrididae: Oedipodinae) from China and its complete mitochondrial genome. <i>Zootaxa</i> , 2019, 4564, 514.	0.2	0
2661	Four new species of <i>Hydnobolites</i> (sequestrate Pezizaceae, Ascomycota) from China. <i>Mycological Progress</i> , 2019, 18, 405-414.	0.5	1
2662	Towards resolving and redefining Amphipyrrinae (Lepidoptera, Noctuoidea, Noctuidae): a massively polyphyletic taxon. <i>Systematic Entomology</i> , 2019, 44, 451-464.	1.7	11
2663	<i>Cladosporium hebeiense</i> sp. nov., pathogenic on grape leaves in China. <i>Mycotaxon</i> , 2019, 133, 643-654.	0.1	1
2664	Genetic heterogeneity of two bioeconomically important kelp species along the Norwegian coast. <i>Conservation Genetics</i> , 2019, 20, 615-628.	0.8	17
2665	Molecular detection of <i>Rickettsia</i> spp., <i>Anaplasma platys</i> and <i>Theileria equi</i> in ticks collected from horses in Tayrona National Park, Colombia. <i>Experimental and Applied Acarology</i> , 2019, 77, 411-423.	0.7	18
2666	Multiple radiations of spiny mice (Rodentia: Acomys) in dry open habitats of Afro-Arabia: evidence from a multi-locus phylogeny. <i>BMC Evolutionary Biology</i> , 2019, 19, 69.	3.2	31
2667	Updated phylogenetic and systematics of the <i>Acrapex albivena</i> Hampson, 1910 and <i>A. stygiata</i> (Hampson, 1910) species groups (Lepidoptera, Noctuidae, Noctuinae, Apameini, Sesamiina), with the description of nine new species from the Afrotropics. <i>Annales De La Societe Entomologique De France</i> , 2019, 55, 219-248.	0.4	4
2668	On the occurrence and molecular identification of <i>Contraecaecum</i> larvae (Nematoda: Anisakidae) in <i>Mugil cephalus</i> from Turkish waters. <i>Parasitology Research</i> , 2019, 118, 1393-1402.	0.6	13
2669	Physiological characterization and sequence analysis of a syringate-consuming Actinobacterium. <i>Bioresource Technology</i> , 2019, 285, 121327.	4.8	13
2670	The complete mitochondrial genomes of three species of <i>Macridiscus</i> Dall, 1902 (Bivalvia: Veneroidea: Tj ETQq1 1 0.784314 rgBT /Over	0.4	4

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2671	Welcome back Janolidae and <i>Antiopella</i> : Improving the understanding of Janolidae and Madrellidae (Cladobronchia, Heterobronchia) with description of four new species. Journal of Zoological Systematics and Evolutionary Research, 2019, 57, 345-368.	0.6	5
2672	On stability measures and effects of data structure in the recognition of areas of endemism. Biological Journal of the Linnean Society, 2019, 127, 143-155.	0.7	5
2673	Secondary Metabolite Dereplication and Phylogenetic Analysis Identify Various Emerging Mycotoxins and Reveal the High Intra-Species Diversity in <i>Aspergillus flavus</i> . Frontiers in Microbiology, 2019, 10, 667.	1.5	24
2674	How soil granulometry, temperature, and water predict genetic differentiation in Namibian spiders () Tj ETQq1 1 0.784314 rgBT /Overlock 10 T	0.8	11
2675	Genetic differences in <i>Chlamydia pecorum</i> between neighbouring sub-populations of koalas ( <i>Phascolarctos cinereus</i> ). Veterinary Microbiology, 2019, 231, 264-270.	0.8	14
2676	An Integrative Systems Biology and Experimental Approach Identifies Convergence of Epithelial Plasticity, Metabolism, and Autophagy to Promote Chemoresistance. Journal of Clinical Medicine, 2019, 8, 205.	1.0	17
2677	Food and feeding habits of <i>Octopus insularis</i> in the Veracruz Reef System National Park and confirmation of its presence in the southwest Gulf of Mexico. Marine Ecology, 2019, 40, e12535.	0.4	19
2678	One stop shop II: taxonomic update with molecular phylogeny for important phytopathogenic genera: 26â€“50 (2019). Fungal Diversity, 2019, 94, 41-129.	4.7	69
2679	Biodiesel from <i>Saccharomyces cerevisiae</i> : fuel property analysis and comparative economics. SN Applied Sciences, 2019, 1, 1.	1.5	9
2680	Eputyfication of <i>Fusarium oxysporum</i> â€“ clearing the taxonomic chaos. Persoonia: Molecular Phylogeny and Evolution of Fungi, 2019, 43, 1-47.	1.6	131
2681	Ciliate Environmental Diversity Can Be Underestimated by the V4 Region of SSU rDNA: Insights from Species Delimitation and Multilocus Phylogeny of <i>Pseudokeronopsis</i> (Ciliophora, Spirotrichea). Microorganisms, 2019, 7, 493.	1.6	14
2682	Demographic Histories and Genome-Wide Patterns of Divergence in Incipient Species of Shorebirds. Frontiers in Genetics, 2019, 10, 919.	1.1	14
2683	Digital Filiation Studies: Phylogenetic Analysis in the Study of Tibetan Buddhist Canonical Texts. , 2019, 209-226.		1
2684	Molecular phylogenetics of Black Cobra ( <i>Naja naja</i> ) in Pakistan. Electronic Journal of Biotechnology, 2019, 42, 23-29.	1.2	4
2685	Maximize Resolution or Minimize Error? Using Genotyping-By-Sequencing to Investigate the Recent Diversification of <i>Helianthemum</i> (Cistaceae). Frontiers in Plant Science, 2019, 10, 1416.	1.7	15
2686	<p><strong><em>Diaporthe</em></strong> from China based on morphological characters and DNA sequence data analyses</strong></p>. Phytotaxa, 2019, 422, 157-174.	0.1	22
2687	<p><strong><em>Leptolejeunea nigra</em></strong> (Lejeuneaceae), a new species with brownish black ocelli based upon morphology and DNA sequences</strong></p>. Phytotaxa, 2019, 427, 31-42.	0.1	3
2688	Mitogenomics suggests a sister relationship of <i>Relicanthus daphneae</i> (Cnidaria: Anthozoa:) Tj ETQq1 1 0.784314 rgBT /Overlock 10 T	1.6	11

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2689	Taxonomy and phylogenetic position of <i>Phragmidium altaicum</i> , a newly described rust fungus on <i>Rosa</i> , based on molecular and morphological data. <i>Phytotaxa</i> , 2019, 423, 187-194.	0.1	4
2690	Two new species of <i>Phyllachora</i> (Phyllachoraceae). <i>Tropical Plant Biology</i> , 2019, 12, 78-86.	0.1	5
2691	Unravelling the phylogeny, cryptic diversity and morphological evolution of <i>Diptilomiopus</i> mites (Acari: Eriophyoidea). <i>Experimental and Applied Acarology</i> , 2019, 79, 323-344.	0.7	9
2692	Neotypification of <i>Fusarium chlamydosporum</i> - a reappraisal of a clinically important species complex. <i>Fungal Systematics and Evolution</i> , 2019, 4, 183-200.	0.9	20
2693	Fish diversification at the pace of geomorphological changes: evolutionary history of western Iberian Leuciscinae (Teleostei: Leuciscidae) inferred from multilocus sequence data. <i>Molecular Phylogenetics and Evolution</i> , 2019, 133, 263-285.	1.2	19
2694	Molecular characterization and phylogenetic analysis of <i>Trypanosoma evansi</i> from Northern India based on 18S ribosomal gene. <i>Veterinary Parasitology: Regional Studies and Reports</i> , 2019, 15, 100259.	0.3	2
2695	An integrative approach elucidates the systematics of <i>Sea Hayward</i> and <i>Cybdelis Boisduval</i> (Lepidoptera: Nymphalidae: Biblidinae). <i>Systematic Entomology</i> , 2019, 44, 226-250.	1.7	4
2696	VARV B22R homologue as phylogenetic marker gene for <i>Capripoxvirus</i> classification and divergence time dating. <i>Virus Genes</i> , 2019, 55, 51-59.	0.7	7
2697	Families in Botryosphaerales: a phylogenetic, morphological and evolutionary perspective. <i>Fungal Diversity</i> , 2019, 94, 1-22.	4.7	63
2698	Morphological and genetic characterization of <i>Syphabulea tjanschani</i> (Nematoda: Oxyuridae), with phylogenetic position of <i>Syphabulea</i> in Oxyuridae. <i>Infection, Genetics and Evolution</i> , 2019, 67, 159-166.	1.0	4
2699	Phylogenetic relationships among Bombycidae s.l. (Lepidoptera) based on analyses of complete mitochondrial genomes. <i>Systematic Entomology</i> , 2019, 44, 490-498.	1.7	27
2700	The Complete Chloroplast Genomes of <i>Echinacanthus</i> Species (Acanthaceae): Phylogenetic Relationships, Adaptive Evolution, and Screening of Molecular Markers. <i>Frontiers in Plant Science</i> , 2018, 9, 1989.	1.7	38
2701	Molecular and morphological evidence for a new species of liverwort, <i>Lejeunea heinrichsii</i> (Marchantiophyta: Lejeuneaceae) from Taveuni, Fiji. <i>Journal of Systematics and Evolution</i> , 2019, 57, 361-370.	1.6	6
2702	Authentication of nine snapper species by single-strand conformation polymorphism (SSCP) and forensically informative nucleotide sequencing (FINS) methods. <i>Food Control</i> , 2019, 99, 124-130.	2.8	6
2703	Molecular Detection of Zoonotic Microsporidia in Domestic Cats in Turkey: A Preliminary Study. <i>Acta Parasitologica</i> , 2019, 64, 13-18.	0.4	17
2704	Phylogeny of the family Cladoniaceae (Lecanoromycetes, Ascomycota) based on sequences of multiple loci. <i>Cladistics</i> , 2019, 35, 351-384.	1.5	29
2705	Molecular phylogeny of <i>Chondrocyclus</i> (Gastropoda: Cyclophoridae), a widespread genus of sedentary, restricted-range snails. <i>Molecular Phylogenetics and Evolution</i> , 2019, 131, 193-210.	1.2	7
2706	Integrative systematics of <i>Placida cremoniana</i> (Trinchese, 1892) (Gastropoda, Heterobranchia). <i>Tropical Plant Biology</i> , 2019, 12, 100259.	0.3	10

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2707	Intron length polymorphism of $\beta$ -tubulin genes of <i>Aegilops biuncialis</i> . <i>Vis. Cell Biology International</i> , 2019, 43, 1031-1039.	1.4	13
2708	The <i>Colletotrichum dracaenophilum</i> , <i>C. magnum</i> and <i>C. orchidearum</i> species complexes. <i>Studies in Mycology</i> , 2019, 92, 1-46.	4.5	165
2709	Seroepidemiology and phylogenetic analysis of human herpesvirus type 8 in injection drug users and men who have sex with men in northern Taiwan. <i>Journal of International Medical Research</i> , 2020, 48, 030006051876474.	0.4	4
2710	Morphological and molecular characterization of <i>Megalobatrachonema hainanensis</i> sp. nov. (Nematoda: Ascaridida), with phylogenetic position of <i>Megalobatrachonema</i> in Cosmocercoidea. <i>Journal of Helminthology</i> , 2020, 94, e19.	0.4	5
2711	Genetic diversity evidence a mixed reproduction mode in <i>Venturia oleaginea</i> populations in Uruguay. <i>Journal of Plant Pathology</i> , 2020, 102, 123-133.	0.6	3
2712	Root nodules of <i>Genista germanica</i> harbor <i>Bradyrhizobium</i> and <i>Rhizobium</i> bacteria exchanging nodC and nodZ genes. <i>Systematic and Applied Microbiology</i> , 2020, 43, 126026.	1.2	6
2713	Phylogenetic analysis of the Archaeocidaridae and Palaeozoic Miocidaridae (Echinodermata). <i>Tj ETQq0 0 0 rgBT /Overlock 10 Jf 50 502 T</i>	0.7	8
2714	<i>Metschnikowia miensis</i> f.a., sp. nov., isolated from flowers in Mie prefecture, Japan. <i>Antonie Van Leeuwenhoek</i> , 2020, 113, 321-329.	0.7	3
2715	Mathematical proof of the third order accuracy of the speedy double bootstrap method. <i>Communications in Statistics - Theory and Methods</i> , 2020, 49, 3950-3964.	0.6	0
2716	Dynamics of the cerebral blood flow response to brief neural activity in human visual cortex. <i>Journal of Cerebral Blood Flow and Metabolism</i> , 2020, 40, 1823-1837.	2.4	13
2717	The most heat-resistant conidia observed to date are formed by distinct strains of <i>Paecilomyces variotii</i> . <i>Environmental Microbiology</i> , 2020, 22, 986-999.	1.8	26
2718	Two novel cricetine mitogenomes: Insight into the mitogenomic characteristics and phylogeny in Cricetinae (Rodentia: Cricetidae). <i>Genomics</i> , 2020, 112, 1716-1725.	1.3	9
2719	Obligate intracellular bacteria diversity in unfed <i>Leptotrombidium scutellare</i> larvae highlights novel bacterial endosymbionts of mites. <i>Microbiology and Immunology</i> , 2020, 64, 1-9.	0.7	13
2720	Multilocus phylogeny of a cryptic radiation of Afrotropical long-fingered bats (Chiroptera). <i>Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 17</i>	0.7	17
2721	Molecular epidemiology of infectious bronchitis virus in Poland from 1980 to 2017. <i>Infection, Genetics and Evolution</i> , 2020, 80, 104177.	1.0	16
2722	<i>Candida auris</i> candidaemia in an intensive care unit – Prospective observational study to evaluate epidemiology, risk factors, and outcome. <i>Journal of Critical Care</i> , 2020, 57, 42-48.	1.0	55
2723	First genetic evidence that invasive bullhead ( <i>Cottus</i> L. 1758) in Scotland is of English origin and the difficulty of resolving the European <i>Cottus</i> species taxonomy. <i>Journal of Fish Biology</i> , 2020, 96, 617-630.	0.7	5
2724	Mitochondrial DNA-Based Identification of Forensically Important Flesh Flies (Diptera: Sarcophagidae) in Thailand. <i>Insects</i> , 2020, 11, 2.	1.0	7

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2725	Description of <i>Aedes (Hulecoeteomyia) bhutanensis</i> n. sp. (Diptera: Culicidae) from Bhutan. <i>Acta Tropica</i> , 2020, 203, 105280.	0.9	7
2726	The enigmatic Leiosaurae clade: Phylogeography, species delimitation, phylogeny and historical biogeography of its southernmost species. <i>Molecular Phylogenetics and Evolution</i> , 2020, 144, 106725.	1.2	10
2727	Phylogenomic analysis of the beetle suborder Adephaga with comparison of tailored and generalized ultraconserved element probe performance. <i>Systematic Entomology</i> , 2020, 45, 552-570.	1.7	35
2728	Divergence and support among slightly suboptimal likelihood gene trees. <i>Cladistics</i> , 2020, 36, 322-340.	1.5	20
2729	Ectopic expression of LoSVP, a MADS-domain transcription factor from lily, leads to delayed flowering in transgenic <i>Arabidopsis</i> . <i>Plant Cell Reports</i> , 2020, 39, 289-298.	2.8	13
2730	Morphology, morphogenesis, and molecular phylogeny of a new freshwater ciliate, <i>Gonostomum jangbogoensis</i> n. sp. (Ciliophora, Hypotricha), from Victoria Land, Antarctica. <i>European Journal of Protistology</i> , 2020, 73, 125669.	0.5	28
2731	A morphological and genetic comparison of <i>Septifer bilocularis</i> , <i>Mytilisepta virgata</i> and <i>Brachidontes variabilis</i> (Bivalvia: Mytiloidea) from Hong Kong and erection of the Mytiliseptiferinae sub-fam. nov.. <i>Regional Studies in Marine Science</i> , 2020, 34, 100981.	0.4	5
2732	Molecular phylogeny of the family Coccidae (Hemiptera, Coccomorpha), with a discussion of their waxy ovisacs. <i>Systematic Entomology</i> , 2020, 45, 396-414.	1.7	13
2733	Too strict or too loose? Integrative taxonomic assessment of <i>Bombus lapidarius</i> complex (Hymenoptera: Apidae). <i>Zoologica Scripta</i> , 2020, 49, 187-196.	0.7	6
2734	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
2735	Born migrators: Historical biogeography of the cosmopolitan family Cannabaceae. <i>Journal of Systematics and Evolution</i> , 2020, 58, 461-473.	1.6	21
2736	Far away, so close! Integrative taxonomy reveals a new genus and species of land flatworm (Platyhelminthes: Geoplanidae) from southern South America. <i>Zoological Journal of the Linnean Society</i> , 2020, 189, 722-744.	1.0	8
2737	Systematic review of the <i>Chromodoris quadricolor</i> group of East Africa, with descriptions of two new species of the genus <i>Chromodoris</i> Alder & Hancock, 1855 (Heterobranchia). <i>Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 10</i>	1.6	16
2738	Evolutionary household archaeology: Inter-generational cultural transmission at housepit 54, Bridge River site, British Columbia. <i>Journal of Archaeological Science</i> , 2020, 124, 105260.	1.2	5
2739	Variation in Botryosphaeriaceae from Eucalyptus plantations in YunNan Province in southwestern China across a climatic gradient. <i>IMA Fungus</i> , 2020, 11, 22.	1.7	25
2740	<i>Mycoenterolobium aquadictyosporium</i> sp. nov. (Pleosporomycetidae, Dothideomycetes) from a freshwater habitat in Thailand. <i>Mycological Progress</i> , 2020, 19, 1031-1042.	0.5	5
2741	Morphology and Molecular Phylogenetic Analysis of Deep-Sea Purple Gorgonians (Octocorallia: Scleractinia) Species. <i>Frontiers in Marine Science</i> , 2020, 7, .	1.2	4
2742	Influence of silver nanoparticles on a common contaminant isolated during the establishment of <i>Stevia rebaudiana</i> Bertoni culture. <i>Plant Cell, Tissue and Organ Culture</i> , 2020, 143, 609-618.	1.2	7



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2743	A phylogeny for the plant pathogen <i>Piptoporellus baudonii</i> using a multigene data set. <i>Mycologia</i> , 2020, 112, 1017-1025.	0.8	7
2744	Generic boundaries in subtribe Saussureinae (Compositae: Cardueae): Insights from HybSeq data. <i>Taxon</i> , 2020, 69, 694-714.	0.4	8
2745	Genotyping <i>Mycoplasma hyorhinitis</i> by multi-locus sequence typing and multiple-locus variable-number tandem-repeat analysis. <i>Veterinary Microbiology</i> , 2020, 249, 108836.	0.8	7
2746	Functional and structural characterization of a novel GH3 $\beta$ -glucosidase from the gut metagenome of the Brazilian Cerrado termite <i>Syntermes wheeleri</i> . <i>International Journal of Biological Macromolecules</i> , 2020, 165, 822-834.	3.6	9
2747	Molecular characterization and gene expression modulation of the alternative oxidase in a scuticociliate parasite by hypoxia and mitochondrial respiration inhibitors. <i>Scientific Reports</i> , 2020, 10, 11880.	1.6	5
2748	Gut microbiome signatures of nursing home residents carrying <i>Enterobacteria</i> producing extended-spectrum $\beta$ -lactamases. <i>Antimicrobial Resistance and Infection Control</i> , 2020, 9, 107.	1.5	12
2749	Phylogenetic relationships of the carnivorous terrestrial snail family Streptaxidae (Stylommatophora: Achatinina) in Thailand and surrounding areas of Southeast Asia. <i>Systematics and Biodiversity</i> , 2020, 18, 720-738.	0.5	7
2750	<i>Neofusicoccum</i> species causing branch cankers on avocado in Crete (Greece). <i>Journal of Plant Pathology</i> , 2020, 102, 1251-1255.	0.6	8
2751	Genetic characterisation of the intermediate hosts of schistosomiasis in its southern distribution limit in South America. <i>Molluscan Research</i> , 2020, 40, 267-275.	0.2	0
2752	<i>Hedyotis</i> , <i>Oldenlandia</i> and related genera (Rubiaceae: Spermacoceae) in Australia: New genera and new combinations in an Asian–Australian–Pacific lineage. <i>Taxon</i> , 2020, 69, 515-542.	0.4	7
2753	A New Report on Edible Tropical Bolete, <i>Phlebopus spongiosus</i> in Thailand and Its Fruiting Body Formation without the Need for a Host Plant. <i>Mycobiology</i> , 2020, 48, 263-275.	0.6	4
2754	Multilocus Sequence Typing Reveals Clonality of Fluconazole-Nonsusceptible <i>Candida tropicalis</i> : A Study From Wuhan to the Global. <i>Frontiers in Microbiology</i> , 2020, 11, 554249.	1.5	13
2755	Adding another piece to the southern African <i>Cercopithecus</i> monkey phylogeography puzzle. <i>African Zoology</i> , 2020, 55, 351-362.	0.2	2
2756	A revision of species of the <i>Parmelia saxatilis</i> complex in the Iberian Peninsula with the description of <i>P. rojoi</i> , a new potentially relict species. <i>Lichenologist</i> , 2020, 52, 365-376.	0.5	5
2757	Two new species and a new record of <i>Nigrograna</i> (Nigrogranaceae, Pleosporales) from China and Thailand. <i>Mycological Progress</i> , 2020, 19, 1365-1375.	0.5	5
2758	Studies in <i>Gyromitra</i> I: the <i>Gyromitra gigas</i> species complex. <i>Mycological Progress</i> , 2020, 19, 1459-1473.	0.5	7
2759	Climatic and topographic changes since the Miocene influenced the diversification and biogeography of the tent tortoise ( <i>Psammobates tentorius</i> ) species complex in Southern Africa. <i>BMC Evolutionary Biology</i> , 2020, 20, 153.	3.2	7
2760	Out of the Hengduan Mountains: Molecular phylogeny and historical biogeography of the Asian water snake genus <i>Trimerodytes</i> (Squamata: Colubridae). <i>Molecular Phylogenetics and Evolution</i> , 2020, 152, 106927.	1.2	11

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2761	Evaluation and utility of mitochondrial ribosomal genes for molecular systematics of parasitic nematodes. <i>Parasites and Vectors</i> , 2020, 13, 364.	1.0	24
2762	A Multiplex PCR Based on Mitochondrial COI Sequences for Identification of Members of the <i>Anopheles barbirostris</i> Complex (Diptera: Culicidae) in Thailand and Other Countries in the Region. <i>Insects</i> , 2020, 11, 409.	1.0	22
2763	The insights into the systematic relationship of <i>Gastrostyla</i> -affinitive genera, with report on a new saline soil ciliate genus and new species (Protozoa, Ciliophora). <i>BMC Evolutionary Biology</i> , 2020, 20, 92.	3.2	18
2764	Fourteen complete mitochondrial genomes of butterflies from the genus <i>Lethe</i> (Lepidoptera,) Tj ETQq1 1 0.784314 rgBT / Overlock 10	1.3	28
2765	<i>Cylindrocladiella peruviana</i> and <i>Pleiocarpon algeriense</i> causing stem and crown rot on avocado ( <i>Persea americana</i> ). <i>European Journal of Plant Pathology</i> , 2020, 158, 419-430.	0.8	6
2766	First report of diazotrophic <i>Brevundimonas</i> spp. as growth enhancer and root colonizer of potato. <i>Scientific Reports</i> , 2020, 10, 12893.	1.6	62
2767	In Silico Study of the Structure and Ligand Interactions of Alcohol Dehydrogenase from <i>Cyanobacterium Synechocystis</i> Sp. PCC 6803 as a Key Enzyme for Biofuel Production. <i>Applied Biochemistry and Biotechnology</i> , 2020, 192, 1346-1367.	1.4	14
2768	Mating strategy and mating type distribution in six global populations of the <i>Eucalyptus</i> foliar pathogen <i>Teratosphaeria destructans</i> . <i>Fungal Genetics and Biology</i> , 2020, 137, 103350.	0.9	19
2769	<i>Mallocybe velutina</i> (Agaricales, Inocybaceae), a new species from Pakistan. <i>Mycoscience</i> , 2020, 61, 348-352.	0.3	6
2770	ITS2 ribotyping, in vitro anti-inflammatory screening, and metabolic profiling of fungal endophytes from the Mexican species <i>Crescentia alata</i> Kunth. <i>South African Journal of Botany</i> , 2020, 134, 213-224.	1.2	8
2771	Infestation Pattern and Population Dynamics of the Tropical Bed Bug, <i>Cimex hemipterus</i> (F.) (Hemiptera:) Tj ETQq0,0,0 rgBT / Overlock 1	1.0	4
2772	A revision of the genus <i>Poconoma</i> Tams & Bowden (Lepidoptera: Noctuidae: Apameini: Sesamiina) with the description of a new genus and two new sesamiine species from the Afrotropical region. <i>Annales De La Societe Entomologique De France</i> , 2020, 56, 313-331.	0.4	1
2773	One-step partial synthesis of (±)-asperteretone B and related hPTP1B1â€‘400 inhibitors from butyrolactone I. <i>Bioorganic and Medicinal Chemistry</i> , 2020, 28, 115817.	1.4	5
2774	Phylogeny and systematics of the <i>Sesamia coniota</i> Hampson species group (Lepidoptera: Noctuidae:) Tj ETQq1 1 0.784314 rgBT / Overlock 1	0.4	1
2775	<i>Profilicollis chasmagnathi</i> (Acanthocephala) parasitizing freshwater fishes: paratenicity and an exception to the phylogenetic conservatism of the genus?. <i>Parasitology Research</i> , 2020, 119, 3957-3966.	0.6	9
2776	The return of the clown: pseudocryptic speciation in the North Pacific clown nudibranch, <i>Triopha catalinae</i> (Cooper, 1863) sensu lato identified by integrative taxonomic approaches. <i>Marine Biodiversity</i> , 2020, 50, 1.	0.3	4
2777	Regional drivers of diversification in the late Quaternary in a widely distributed generalist species, the common pheasant <i>Phasianus colchicus</i> . <i>Journal of Biogeography</i> , 2020, 47, 2714-2727.	1.4	10
2778	<p><strong>Additions to Karst Fungi 4:<em> Botryosphaeria</em> spp. associated with woody hosts in Guizhou province, China including <em>B. guttulata</em> <em>sp. nov.</em></strong></p>. <i>Phytotaxa</i> , 2020, 454, 186-202.	0.1	10

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2779	Involvement of hpap2 and dgkA Genes in Colistin Resistance Mediated by mcr Determinants. Antibiotics, 2020, 9, 531.	1.5	9
2780	The complete mitochondrial genome sequence of <i>Cletus rubidiventris</i> (Heteroptera: Coreidae). Mitochondrial DNA Part B: Resources, 2020, 5, 3075-3076.	0.2	1
2781	The Ability of Taxonomic Identification of Bifidobacteria Based on the Variable Regions of 16S rRNA Gene. Russian Journal of Genetics, 2020, 56, 926-934.	0.2	6
2782	Circumscription and phylogenetic position of <i>Ligularia</i> sect. <i>Stenostegia</i> (Asteraceae: Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 587 Td (Raphi) 0.4 5	0.4	5
2783	Morphological and molecular identification of Hysterothylacium larvae (Nematoda: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 587 Td (Raphi) 0.6 2	0.6	2
2784	Integrative taxonomy of the new millipede genus <i>Coxobolellus</i> , gen. nov. (Diplopoda : Spirobolida : Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 587 Td (Raphi) 0.5 10	0.5	10
2785	An Evolving View of Phylogenetic Support. Systematic Biology, 2022, 71, 921-928.	2.7	19
2786	<strong>Taxonomic evaluation of two similar bent-toed geckos</strong> <strong>Squamata:</strong> 2020, 4830, 186-196.	0.2	4
2787	Piscine Orthoreovirus-1 Isolates Differ in Their Ability to Induce Heart and Skeletal Muscle Inflammation in Atlantic Salmon ( <i>Salmo salar</i> ). Pathogens, 2020, 9, 1050.	1.2	28
2788	Comparative genomics of <i>Klebsiella michiganensis</i> BD177 and related members of <i>Klebsiella</i> sp. reveal the symbiotic relationship with <i>Bactrocera dorsalis</i> . BMC Genetics, 2020, 21, 138.	2.7	5
2789	Biological and Genomic Characterization of a Novel Jumbo Bacteriophage, vB_VhaM_pir03 with Broad Host Lytic Activity against <i>Vibrio harveyi</i> . Pathogens, 2020, 9, 1051.	1.2	20
2790	<strong>Roussoella guttulata</strong> (Roussoellaceae, Pleosporales), a novel bambusicolous ascomycete from Thailand. Phytotaxa, 2020, 471, 221-233.	0.1	6
2791	Target-capture phylogenomics provide insights on gene and species tree discordances in Old World treefrogs (Anura: Rhacophoridae). Proceedings of the Royal Society B: Biological Sciences, 2020, 287, 20202102.	1.2	17
2792	<strong>A new species of <i>Clavariadelphus</i> (Basidiomycota)</strong> Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 587 Td (Raphi) 0.1 1	0.1	1
2793	Dieback of <i>Euonymus alatus</i> (Celastraceae) Caused by <i>Cytospora haidianensis</i> sp. nov. in China. Forests, 2020, 11, 524.	0.9	9
2794	Independent Recruitment of Duplicated $\hat{2}$ -Subunit-Coding NAD-ME Genes Aided the Evolution of C4 Photosynthesis in Cleomaceae. Frontiers in Plant Science, 2020, 11, 572080.	1.7	12
2795	Gene Sequences of Potential Targets of Insecticidal PF2 Lectin Identified from the Larval De Novo Transcriptome of the Mexican Bean Weevil ( <i>Zabrotes Subfasciatus</i> ; Boheman 1833). Insects, 2020, 11, 736.	1.0	3
2796	Clinical Characteristics of Acute Hepatitis E and Their Correlation with HEV Genotype 3 Subtypes in Italy. Pathogens, 2020, 9, 832.	1.2	9

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2797	Integrative systematics reveals the new land-snail genus <i>Taphrenalla</i> (Eupulmonata: Ariophantidae) with a description of nine new species from Thailand. <i>Contributions To Zoology</i> , 2020, 90, 21-69.	0.2	13
2798	Taxonomic revision of the <i>Sylvarum</i> group of bumblebees using an integrative approach. <i>Systematics and Biodiversity</i> , 2020, 18, 12-28.	0.5	6
2799	A new species of phytotelm breeding frog (Anura: Rhacophoridae) from the Central Highlands of Vietnam. <i>Zootaxa</i> , 2020, 4779, zootaxa.4779.3.3.	0.2	7
2800	Phylogenetic placement of <i>Paratrachaptum</i> and reconsideration of <i>Gloeophyllales</i> . <i>Fungal Systematics and Evolution</i> , 2020, 5, 119-130.	0.9	3
2801	Molecular and morphological evidence of sibling species in <i>Anopheles baileyi</i> Edwards (Diptera: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 58	0.9	9
2802	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
2803	Genomics and High-Resolution Typing Confirm Predominant Clonal Evolution Down to a Microevolutionary Scale in <i>Trypanosoma cruzi</i> . <i>Pathogens</i> , 2020, 9, 356.	1.2	3
2804	Nuclear phylogenomic analyses of asterids conflict with plastome trees and support novel relationships among major lineages. <i>American Journal of Botany</i> , 2020, 107, 790-805.	0.8	75
2805	The Cycad Genus <i>Cycas</i> May Have Diversified From Indochina and Occupied Its Current Ranges Through Vicariance and Dispersal Events. <i>Frontiers in Ecology and Evolution</i> , 2020, 8, .	1.1	13
2806	An integrative approach to the systematics of the <i>Berthella californica</i> species complex (Heterobranchia: Pleurobranchidae). <i>Journal of Molluscan Studies</i> , 2020, 86, 186-200.	0.4	2
2807	Diversity in Chlamydial plasmids. <i>PLoS ONE</i> , 2020, 15, e0233298.	1.1	15
2808	Integrated systematics of <i>Anopheles subpictus</i> (Diptera: Culicidae) in the Oriental Region, with emphasis on forms in Thailand and Sulawesi, Indonesia. <i>Acta Tropica</i> , 2020, 208, 105503.	0.9	7
2809	Dimeric phenalenones from <i>Talaromyces</i> sp. (IQ-313) inhibit hPTP1B1-400: Insights into mechanistic kinetics from in vitro and in silico studies. <i>Bioorganic Chemistry</i> , 2020, 101, 103893.	2.0	16
2810	Large-Scale Hybridisation as an Extinction Threat to the Suweon Treefrog (Hylidae: Dryophytes) Tj ETQq1 1 0.784314 rgBT /Overlock 10 8	1.0	8
2811	A new species of <i>Phrynobatrachus</i> (Amphibia: Anura: Phrynobatrachidae) from the Northern Mountains of Tanzania. <i>Journal of Natural History</i> , 2020, 54, 63-85.	0.2	3
2812	LtEPG1, a Secretory Endopolygalacturonase Protein, Regulates the Virulence of <i>Lasiodiplodia theobromae</i> in <i>Vitis vinifera</i> and Is Recognized as a Microbe-Associated Molecular Patterns. <i>Phytopathology</i> , 2020, 110, 1727-1736.	1.1	13
2813	A new species of the genus <i>Zhangixalus</i> (Amphibia: Rhacophoridae) from Vietnam. <i>Journal of Natural History</i> , 2020, 54, 257-273.	0.2	7
2814	New haplochromine cichlid from the upper Miocene (9–10 MYA) of Central Kenya. <i>BMC Evolutionary Biology</i> , 2020, 20, 65.	3.2	8

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2815	The enigmatic case of the genus <i>Argyresthia</i> in the Azores Islands (Lepidoptera: Tj ETQq0 0 0 rgBT /Overlock 10 Tf	0.2	2
2816	Modeling the Evolution of Ceramic Traditions Through a Phylogenetic Analysis of the Chañes Opatoires: the European Bronze Age as a Case Study. Journal of Archaeological Method and Theory, 2020, 27, 992-1039.	1.4	15
2817	Morphology and phylogenetic analysis of two new species of deep-sea golden gorgonians (Cnidaria: Tj ETQq0 0 0 rgBT /Overlock 10 Tf zootaxa.4731.2.4.	0.2	5
2818	Description of the immature stages of the Neotropical whirligig beetle <i>Gyrinus</i> ( <i>Neogyrinus</i> ) <i>rozei</i> Ochs, 1953 (Coleoptera: Gyrinidae) and first report of the parasitoid wasp <i>Melanosmicra</i> sp. (Hymenoptera: Chalcididae) on a <i>Gyrinus</i> species. Zootaxa, 2020, 4732, 99-116.	0.2	1
2819	Phylogenetic Analysis and Development of Molecular Tool for Detection of Diaporthe citri Causing Melanose Disease of Citrus. Plants, 2020, 9, 329.	1.6	16
2820	High genetic connectivity in a gastropod with long-lived planktonic larvae. Journal of Molluscan Studies, 2020, 86, 42-55.	0.4	5
2821	A morphometric and molecular phylogenetic analysis of the African orchid genus <i>Stenoglottis</i> (Orchidaceae: Orchidoideae). Botanical Journal of the Linnean Society, 2020, 193, 340-362.	0.8	5
2822	Taxonomy and phylogeny of hyaline-spored coelomycetes. Fungal Diversity, 2020, 100, 279-801.	4.7	58
2823	Phylogenetic relationships of Asian freshwater Mytiloidea (Bivalvia): a morphological and genetic comparison of <i>Sinomytilus harmandi</i> , <i>Limnoperna fortunei</i> and <i>Septifer bilocularis</i> . Molluscan Research, 2020, 40, 120-129.	0.2	5
2824	A new species of <i>Dixonius</i> (Squamata: Gekkonidae) from the karst forest of Khammouane Province, central Laos. Zootaxa, 2020, 4759, zootaxa.4759.4.4.	0.2	3
2825	Phylogeny and classification of Leptophlebiidae (Ephemeroptera) with an emphasis on Neotropical fauna. Systematic Entomology, 2020, 45, 415-429.	1.7	15
2826	Economic and feasibility comparison of the dRIT and DFA for decentralized rabies diagnosis in resource-limited settings: The use of Nigerian dog meat markets as a case study. PLoS Neglected Tropical Diseases, 2020, 14, e0008088.	1.3	7
2827	Phylogeny and Historical Biogeography of <i>Paphiopedilum Pfitzer</i> (Orchidaceae) Based on Nuclear and Plastid DNA. Frontiers in Plant Science, 2020, 11, 126.	1.7	10
2828	Morphological, Molecular, and Pathological Appraisal of <i>Hymenolepis nana</i> (Hymenolepididae) Infecting Laboratory Mice ( <i>Mus musculus</i> ). Microscopy and Microanalysis, 2020, 26, 348-362.	0.2	4
2829	Keratinophilic fungi: Specialized fungal communities in a desert ecosystem identified using cultured-based and Illumina sequencing approaches. Microbiological Research, 2020, 239, 126530.	2.5	12
2830	Essential oil analysis of eight 'Nepeta' taxa in Iran. Mediterranean Botany, 2020, 41, 43-53.	0.9	6
2831	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). Journal of Molluscan Studies, 2020, 86, 240-248.	0.4	1
2832	From Folk Taxonomy to Species Confirmation of <i>Acorus</i> (Acoraceae): Evidences Based on Phylogenetic and Metabolomic Analyses. Frontiers in Plant Science, 2020, 11, 965.	1.7	24

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2833	Taxonomic revision of the Afrotropical <i>Phytomia Guérin-Méneville</i> (Diptera: Syrphidae). <i>Zootaxa</i> , 2020, 4803, zootaxa.4803.2.1.	0.2	4
2834	Molecular phylogeny and species delimitation of Stachyuraceae: Advocating a herbarium specimen-based phylogenomic approach in resolving species boundaries. <i>Journal of Systematics and Evolution</i> , 2020, 58, 710-724.	1.6	6
2835	High Diversity of <i>Cytospora</i> Associated With Canker and Dieback of Rosaceae in China, With 10 New Species Described. <i>Frontiers in Plant Science</i> , 2020, 11, 690.	1.7	29
2836	Insight into Evolution and Conservation Patterns of B1-Subfamily Members of GPCR. <i>International Journal of Peptide Research and Therapeutics</i> , 2020, 26, 2505-2517.	0.9	3
2837	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 aenigma</i> group. <i>Annales De La Societe Entomologique De France</i> , 2020, 56, 29-91.	0.4	4
2838	Molecular and Morphological Characterization of Two Novel Species Collected from Soil in Korea. <i>Mycobiology</i> , 2020, 48, 9-19.	0.6	6
2839	Investigating the amphiatlantic status of <i>Facelina bostoniensis</i> (Couthouy, 1838) (Nudibranchia: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 18).	0.4	3
2840	It is not a disaster: molecular and morphologically based phylogenetic analysis of <i>Rondeletieae</i> and the <i>Rondeletia</i> complex (Cinchonoideae, Rubiaceae). <i>Plant Systematics and Evolution</i> , 2020, 306, 1.	0.3	6
2841	Worldwide Phylogeography of <i>Ceratitis capitata</i> (Diptera: Tephritidae) Using Mitochondrial DNA. <i>Journal of Economic Entomology</i> , 2020, 113, 1455-1470.	0.8	12
2842	Diversity of <i>Mycobacteriaceae</i> from aquatic environment at the São Paulo Zoological Park Foundation in Brazil. <i>PLoS ONE</i> , 2020, 15, e0227759.	1.1	4
2843	On the origin and diversification history of the African genus <i>Encephalartos</i> . <i>South African Journal of Botany</i> , 2020, 130, 231-239.	1.2	12
2844	Stargazing under the sea: molecular and morphological data reveal a constellation of species in the <i>Berthella stellata</i> (Risso, 1826) species complex (Mollusca, Heterobranchia, Pleurobranchidae). <i>Marine Biodiversity</i> , 2020, 50, 1.	0.3	6
2845	Molecular Detection of Rabies Lyssaviruses from Dogs in Southeastern Nigeria: Evidence of Transboundary Transmission of Rabies in West Africa. <i>Viruses</i> , 2020, 12, 134.	1.5	9
2846	Anthraxnose Disease of Carpetgrass ( <i>Axonopus compressus</i> ) Caused by <i>Colletotrichum hainanense</i> sp. nov.. <i>Plant Disease</i> , 2020, 104, 1744-1750.	0.7	8
2847	A taxonomic conundrum: Characterizing a cryptic radiation of Asian gracile skinks (Squamata: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 18).	1.2	2
2848	A bushel of viruses: Identification of seventeen novel putative viruses by RNA-seq in six apple trees. <i>PLoS ONE</i> , 2020, 15, e0227669.	1.1	36
2849	Assessment of species boundaries of the <i>Moringa ovalifolia</i> in Namibia using nuclear its DNA sequence data. <i>South African Journal of Botany</i> , 2020, 131, 335-341.	1.2	1
2850	A revision of <i>Neusticomys peruviansis</i> (Rodentia: Cricetidae) with the description of a new subspecies. <i>Journal of Mammalogy</i> , 2020, 101, 858-871.	0.6	4

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2852	Morphology, DNA Phylogeny, and Pathogenicity of <i>Wilsonomyces carpophilus</i> Isolate Causing Shot-Hole Disease of <i>Prunus divaricata</i> and <i>Prunus armeniaca</i> in Wild-Fruit Forest of Western Tianshan Mountains, China. <i>Forests</i> , 2020, 11, 319.	0.9	5
2853	Candidate cases of poecilogony in Neogastropoda: implications for the systematics of the genus <i>Raphitoma</i> Bellardi, 1847. <i>Invertebrate Systematics</i> , 2020, , .	0.5	1
2854	Unexpected predicted length variation for the coding sequence of the sleep related gene, <i>BHLHE41</i> in gorilla amidst strong purifying selection across mammals. <i>PLoS ONE</i> , 2020, 15, e0223203.	1.1	0
2855	Rabies in the African Civet: An Incidental Host for Lyssaviruses?. <i>Viruses</i> , 2020, 12, 368.	1.5	9
2856	Molecular and morphological evidence for sibling species within <i>Anopheles</i> ( <i>Anopheles</i> ) <i>lindesayi</i> Giles (Diptera: Culicidae) in Bhutan. <i>Acta Tropica</i> , 2020, 207, 105455.	0.9	15
2857	Phenotypic and molecular characterisation of <i>Sporothrix globosa</i> of diverse origin from India. <i>Brazilian Journal of Microbiology</i> , 2021, 52, 91-100.	0.8	14
2858	New Burmese amber fossils clarify the evolution of bethylid wasps (Hymenoptera: Chrysididae). <i>Zoological Journal of the Linnean Society</i> , 2021, 191, 1044-1058.	1.0	10
2859	Another stripe on the tiger makes no difference? Unexpected diversity in the widespread tiger tarantula <i>Davus pentalaris</i> (Araneae: Theraphosidae: Theraphosinae). <i>Zoological Journal of the Linnean Society</i> , 2021, 192, 75-104.	1.0	8
2860	Molecular phylogeny, species delimitation and biogeographic history of the <i>Stegana</i> ( <i>Steganina</i> ) <i>shirozui</i> species group (Diptera: Drosophilidae) from East Asia. <i>Zoological Journal of the Linnean Society</i> , 2021, 192, 998-1016.	1.0	2
2861	A review of extant <i>Tudivasum</i> Rosenberg & Petit, 1987 (Neogastropoda: Turbinellidae) and description of three new species from Western Australia. <i>Journal of Molluscan Studies</i> , 2021, 87, .	0.4	2
2862	Ionizing-radiation-resistant <i>Kocuria rhizophila</i> PT10 isolated from the Tunisian Sahara xerophyte <i>Panicum turgidum</i> : Polyphasic characterization and proteogenomic arsenal. <i>Genomics</i> , 2021, 113, 317-330.	1.3	7
2863	Screening and characterization of polyhydroxyalkanoate granules, and phylogenetic analysis of polyhydroxyalkanoate synthase gene <i>PhaC</i> in cyanobacteria. <i>Journal of Phycology</i> , 2021, 57, 754-765.	1.0	6
2864	Colpodean ciliate phylogeny and reference alignments for phylogenetic placements. <i>European Journal of Protistology</i> , 2021, 77, 125747.	0.5	5
2865	Genetic population structure of the monogenean parasite <i>Gyrodactylus thymalli</i> and its host European grayling ( <i>Thymallus thymallus</i> ) in a large Norwegian lake. <i>Hydrobiologia</i> , 2021, 848, 547-561.	1.0	3
2866	<i>Colletotrichum</i> spp. causing anthracnose on ornamental plants in northern Italy. <i>Journal of Plant Pathology</i> , 2021, 103, 127-137.	0.6	15
2867	Phylogenetics of <i>Paepalanthus</i> (Eriocaulaceae), a diverse Neotropical monocot lineage. <i>Botanical Journal of the Linnean Society</i> , 2021, 195, 34-52.	0.8	12
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2870	Phytochemical Analysis and Evaluation of Antioxidant and Biological Activities of Extracts from Three Clausenae Plants in Northern Thailand. <i>Plants</i> , 2021, 10, 117.	1.6	7
2871	Two new species of <i>Panellus</i> (Agaricales, Basidiomycota) from China. <i>Mycological Progress</i> , 2021, 20, 51-60.	0.5	2
2872	Intraspecific Differentiation in White Mistletoe ( <i>Viscum album</i> L.) Using the Analysis of Intron Length Polymorphism of $\beta$ -Tubulin Genes and the SSR Analysis. <i>Cytology and Genetics</i> , 2021, 55, 1-9.	0.2	3
2873	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
2874	Undressing <i>Lophodoris danielsseni</i> (Friele & Hansen, 1878) (Nudibranchia: Goniadorididae). <i>Organisms Diversity and Evolution</i> , 2021, 21, 107-117.	0.7	7
2875	Identification of a predominant genotype of <i>Mycobacterium tuberculosis</i> in Brazilian indigenous population. <i>Scientific Reports</i> , 2021, 11, 1224.	1.6	3
2876	Models in parasite and pathogen evolution: Genomic analysis reveals predominant clonality and progressive evolution at all evolutionary scales in parasitic protozoa, yeasts and bacteria. <i>Advances in Parasitology</i> , 2021, 111, 75-117.	1.4	10
2877	Wildlife Forensic Genetics and Biodiversity Conservation: The Intersection of Science, Species Management, and the Law. , 2021, , 163-191.		0
2878	Using heritability of stellar chemistry to reveal the history of the Milky Way. <i>Monthly Notices of the Royal Astronomical Society</i> , 2021, 502, 32-47.	1.6	6
2879	A new family and two new genera of calcaxonian octocoral, including a redescription of. <i>Invertebrate Systematics</i> , 2021, 35, 282-297.	0.5	2
2880	Revisiting the phylogenetic predicament of the genus <i>Huia</i> (Amphibia: Ranidae) using molecular data and tadpole morphology. <i>Zoological Journal of the Linnean Society</i> , 2021, 193, 673-699.	1.0	5
2881	Diversity unearthed by the estimated molecular phylogeny and ecologically quantitative characteristics of uncultured Ehrlichia bacteria in Haemaphysalis ticks, Japan. <i>Scientific Reports</i> , 2021, 11, 687.	1.6	9
2882	<i>Reinertia</i> , a New Subgenus of the Genus <i>Aedes</i> Meigen and Its Type Species <i>Aedes</i> ( <i>Reinertia</i> ) <i>suffusus</i> (Diptera: Culicidae), Newly Recorded From Bhutan. <i>Journal of Medical Entomology</i> , 2021, 58, 1138-1148.	0.9	5
2883	Reanalysis and Revision of the Complete Mitochondrial Genome of <i>Artemia urmiana</i> G&A1/4nther, 1899 (Crustacea: Anostraca). <i>Diversity</i> , 2021, 13, 14.	0.7	8
2884	Species Diversity and Molecular Phylogeny of <i>Cyanosporus</i> (Polyporales, Basidiomycota). <i>Frontiers in Microbiology</i> , 2021, 12, 631166.	1.5	16
2885	Absence of metabotropic glutamate receptor homolog(s) accelerates acetylcholine neurotransmission in <i>Caenorhabditis elegans</i> . <i>Neuroscience Letters</i> , 2021, 746, 135666.	1.0	5
2886	Genomic islands mediate environmental adaptation and the spread of antibiotic resistance in multiresistant Enterococci - evidence from genomic sequences. <i>BMC Microbiology</i> , 2021, 21, 55.	1.3	14



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2888	Two new <i>Morinia</i> species from palms ( <i>Arecaceae</i> ) in Portugal. <i>Mycological Progress</i> , 2021, 20, 83-94.	0.5	1
2889	Taxonomy and Phylogeny of the <i>Fomitopsis pinicola</i> Complex With Descriptions of Six New Species From East Asia. <i>Frontiers in Microbiology</i> , 2021, 12, 644979.	1.5	24
2890	First report and molecular prevalence of potential zoonotic <i>Enterocytozoon bienersi</i> in Turkish tumbler pigeons ( <i>Columba livia domestica</i> ). <i>Medical Mycology</i> , 2021, 59, 864-868.	0.3	8
2891	An Integrative Approach for the Characterization of Plant-Pathogenic <i>Streptomyces</i> spp. Strains Based on Metabolomic, Bioactivity, and Phylogenetic Analysis. <i>Frontiers in Microbiology</i> , 2021, 12, 643792.	1.5	3
2892	Four novel <i>Myxobolus</i> species (Cnidaria: Myxozoa) infecting Anatolian <i>khramulya</i> <i>Capoeta tinca</i> (Cyprinidae) in northern Turkey. <i>Diseases of Aquatic Organisms</i> , 2021, 144, 41-54.	0.5	0
2893	Genome-wide identification and expression profiling revealed tissue-specific inducible expression of cytochrome P450s conferring cadmium tolerance in the Pacific oyster, <i>Crassostrea gigas</i> . <i>Aquaculture Reports</i> , 2021, 19, 100582.	0.7	1
2894	<i>Placobdelloides tridens</i> sp. n., a new species of glossiphoniid leech (Hirudinea: Rhynchobdellida) found feeding on captive <i>Oritia borneensis</i> in Thailand, and an update to the host distribution of <i>P. siamensis</i> . <i>Systematic Parasitology</i> , 2021, 98, 141-154.	0.5	5
2895	Disentangling <i>Antirhea</i> (Rubiaceae): resurrection of <i>Guettardella</i> and description of the new genus <i>Achilleanthus</i> . <i>Botanical Journal of the Linnean Society</i> , 2021, 197, 85-103.	0.8	0
2896	Multiple lines of evidence reveal a new species of Krait (Squamata, Elapidae, <i>Bungarus</i> ) from Southwestern China and Northern Myanmar. <i>ZooKeys</i> , 2021, 1025, 35-71.	0.5	10
2897	Morphometric and Molecular Diversity among Seven European Isolates of <i>Pratylenchus penetrans</i> . <i>Plants</i> , 2021, 10, 674.	1.6	4
2898	Isolation, Identification, and Analysis of Potential Functions of Culturable Bacteria Associated with an Invasive Gall Wasp, <i>Leptocybe invasa</i> . <i>Microbial Ecology</i> , 2022, 83, 151-166.	1.4	7
2899	Volatile Organic Compound from <i>Trichoderma asperelloides</i> TSU1: Impact on Plant Pathogenic Fungi. <i>Journal of Fungi</i> (Basel, Switzerland), 2021, 7, 187.	1.5	38
2900	Mitochondrial analysis of oribatid mites provides insights into their atypical tRNA annotation, genome rearrangement and evolution. <i>Parasites and Vectors</i> , 2021, 14, 221.	1.0	2
2901	Identification of Bloodmeals from Sand Flies (Diptera: Psychodidae) Collected in the Parque Nacional do Viruãj, State of Roraima, Brazil. <i>Journal of Medical Entomology</i> , 2021, 58, 2488-2494.	0.9	4
2902	Phylogeny, divergence times, and diversification in <i>Calophyllaceae</i> : Linking key characters and habitat changes to the evolution of Neotropical <i>Calophylleae</i> . <i>Molecular Phylogenetics and Evolution</i> , 2021, 157, 107041.	1.2	7
2903	Invasion of <i>Eragrostis albensis</i> in Central Europe: distribution patterns, taxonomy and phylogenetic insight into the <i>Eragrostis pilosa</i> complex. <i>Biological Invasions</i> , 2021, 23, 2305-2327.	1.2	2
2904	Re-Description of <i>Sillago malabarica</i> , Silver Whiting from Southern Indian Waters. <i>Thalassas</i> , 2021, 37, 627-640.	0.1	1

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2906	Additions to the Genus <i>Arthrinium</i> (Apiosporaceae) From Bamboos in China. <i>Frontiers in Microbiology</i> , 2021, 12, 661281.	1.5	20
2907	Back from the past: Molecular and morphological support for <i>Simulium mutucuna</i> Nunes de Mello & Vieira da Silva, 1974 (Diptera: Simuliidae) as a valid species. <i>Acta Tropica</i> , 2021, 216, 105846.	0.9	1
2908	DNA barcoding of important fruit tree species of agronomic interest in the genus <i>Garcinia</i> L. from the Western Ghats. <i>Genetic Resources and Crop Evolution</i> , 2021, 68, 3161-3177.	0.8	2
2909	Two new species of <i>Litostigma</i> (Gesneriaceae) from the China-Vietnam border area based on morphological and molecular data, adding new stigma characters for the genus. <i>Nordic Journal of Botany</i> , 2021, 39, .	0.2	0
2910	Phylogeny and divergence dating of the ladybird beetle tribe Coccinellini Latreille (Coleoptera: Tj ETQq1 1 0.784314,rgBT /Overlock 10	1.7	8
2911	Soil Metabarcoding Offers a New Tool for the Investigation and Hunting of Truffles in Northern Thailand. <i>Journal of Fungi</i> (Basel, Switzerland), 2021, 7, 293.	1.5	2
2912	Out of the trap: A new phytothelma breeding species of <i>Philautus</i> and an updated phylogeny of Bornean bush frogs (Anura: Rhacophoridae). <i>Journal of Zoological Systematics and Evolutionary Research</i> , 2021, 59, 1064-1096.	0.6	3
2913	A new species of <i>Ahaetulla</i> Link, 1807 (Squamata: Colubridae) from the Mekong Delta, Vietnam. <i>Zootaxa</i> , 2021, 4966, 290304.	0.2	2
2914	A morphological and molecular review of the genus <i>Goniurosaurus</i> , including an identification key. <i>European Journal of Taxonomy</i> , 0, 751, .	0.6	3
2915	A new species of <i>Sakuraeolis</i> from Mozambique, described using 3D reconstruction of anatomy and phylogenetic analysis. <i>Journal of Molluscan Studies</i> , 2021, 87, .	0.4	4
2916	Two new cryptic species of <i>Microhyla</i> Tschudi, 1838 (Amphibia, Anura, Microhylidae) related to the <i>M. heymonsi</i> group from central Vietnam. <i>ZooKeys</i> , 2021, 1036, 47-74.	0.5	6
2917	The Hidden Diversity of Diatrypaceous Fungi in China. <i>Frontiers in Microbiology</i> , 2021, 12, 646262.	1.5	12
2918	A worthy conservation target? Revising the status of the rarest bumblebee of Europe. <i>Insect Conservation and Diversity</i> , 2021, 14, 661-674.	1.4	13
2919	Morphometric Analysis of <i>Coptotermes</i> spp. Soldier Caste (Blattodea: Rhinotermitidae) in Indonesia and Evidence of <i>Coptotermes gestroi</i> Extreme Head-Capsule Shapes. <i>Insects</i> , 2021, 12, 477.	1.0	5
2920	Building of an Internal Transcribed Spacer (ITS) Gene Dataset to Support the Italian Health Service in Mushroom Identification. <i>Foods</i> , 2021, 10, 1193.	1.9	7
2921	Phylogenetic assessment and taxonomic revision of <i>Halobyssothecium</i> and <i>Lentithecium</i> (Lentitheciaceae, Pleosporales). <i>Mycological Progress</i> , 2021, 20, 701-720.	0.5	12
2922	DNA barcoding of <i>Thrixspermum longipilosum</i> based on Internal Transcribed Spacer 2 (ITS2) region. <i>IOP Conference Series: Earth and Environmental Science</i> , 2021, 743, 012092.	0.2	0

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2923	Toward a Stable Global Noctuidae (Lepidoptera) Taxonomy. <i>Insect Systematics and Diversity</i> , 2021, 5, .	0.7	24
2924	Occurrence of Canker and Wood Rot Pathogens on Stone Fruit Propagation Material and Nursery Trees in the Western Cape of South Africa. <i>Plant Disease</i> , 2021, 105, 3586-3599.	0.7	5
2925	Testing the Utility of Alternative Metrics of Branch Support to Address the Ancient Evolutionary Radiation of Tunas, Stromateoids, and Allies (Teleostei: Pelagiaria). <i>Systematic Biology</i> , 2021, 70, 1123-1144.	2.7	19
2926	Varadia, a new helicarionoidean semi-slug genus from India's Western Ghats (Stylommatophora: Tj ETQq1 1 0.784314 rgBT /Ove	0.6	4
2927	Phylogeny of terraranan frogs based on 2,665 loci and impacts of missing data on phylogenomic analyses. <i>Systematics and Biodiversity</i> , 2021, 19, 818-833.	0.5	10
2928	Using stratified Bayesian model averaging in probabilistic forecasts of precipitation over the middle and lower Yangtze River region. <i>Meteorology and Atmospheric Physics</i> , 2021, 133, 961-972.	0.9	3
2929	Revision of the Afrotropical species of the hover fly genus <i>Mesembrius</i> Rondani (Diptera, Syrphidae) using morphological and molecular data. <i>ZooKeys</i> , 2021, 1046, 1-141.	0.5	2
2930	Rhizobia and endophytic bacteria isolated from rainforest fragments within an iron ore mining site of the Eastern Brazilian Amazon. <i>Brazilian Journal of Microbiology</i> , 2021, 52, 1461-1474.	0.8	5
2931	Revising dating estimates and the antiquity of eusociality in termites using the fossilized birth-death process. <i>Systematic Entomology</i> , 2021, 46, 592-610.	1.7	25
2932	Rapid radiation of angraecoids (Orchidaceae, Angraecinae) in tropical Africa characterised by multiple karyotypic shifts under major environmental instability. <i>Molecular Phylogenetics and Evolution</i> , 2021, 159, 107105.	1.2	7
2933	Molecular and Morphological Assessment of <i>Septoria</i> Species Associated with Ornamental Plants in Yunnan Province, China. <i>Journal of Fungi (Basel, Switzerland)</i> , 2021, 7, 483.	1.5	0
2934	Evolutionary relationships among the snakelike pygopodid lizards: a review of phylogenetic studies of an enigmatic Australian adaptive radiation. <i>PeerJ</i> , 2021, 9, e11502.	0.9	0
2935	Morphological and Molecular Characterization of Five Species Including Three New Species of Golden Gorgonians (Cnidaria: Octocorallia) from Seamounts in the Western Pacific. <i>Biology</i> , 2021, 10, 588.	1.3	1
2936	Molecular and morphological analyses reveal pseudocryptic diversity in <i>Micromelodanus</i> (Bruguière, 1792) (Gastropoda: Heterobranchia: Aplustridae). <i>Systematics and Biodiversity</i> , 2021, 19, 834-858.	0.5	5
2937	Characterization, Comparative Analysis and Phylogenetic Implications of Mitogenomes of Fulgoridae (Hemiptera: Fulgoromorpha). <i>Genes</i> , 2021, 12, 1185.	1.0	3
2938	<i>Biscogniauxia rosacearum</i> the charcoal canker agent as a pathogen associated with grapevine trunk diseases in Zagros region of Iran. <i>Scientific Reports</i> , 2021, 11, 14098.	1.6	13
2939	Molecular phylogeny of European Runcinida (Gastropoda, Heterobranchia): the discover of an unexpected pool of complex species, with special reference to the case of <i>Runcina coronata</i> . <i>Zoological Journal of the Linnean Society</i> , 2022, 194, 761-788.	1.0	7
2940	Concordance-Based Approaches for the Inference of Relationships and Molecular Rates with Phylogenomic Data Sets. <i>Systematic Biology</i> , 2022, 71, 943-958.	2.7	11

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2942	Molecular phylogenetic analysis of the <i>Paguristes tortugae</i> Schmitt, 1933 complex and selected other Paguroidea (Crustacea: Decapoda: Anomura). <i>Zootaxa</i> , 2021, 4999, 301-324.	0.2	2
2943	Parsimony analysis of phylogenomic datasets (II): evaluation of PAUP*, MEGA and MPBoot. <i>Cladistics</i> , 2022, 38, 126-146.	1.5	11
2944	Including fossils in phylogeny: a glimpse into the evolution of the superfamily Evanioidea (Hymenoptera: Apocrita) under tip-dating and the fossilized birth-death process. <i>Zoological Journal of the Linnean Society</i> , 2022, 194, 1396-1423.	1.0	19
2945	Phylogenetics of mud snakes (Squamata: Serpentes: Homalopsidae): A paradox of both undescribed diversity and taxonomic inflation. <i>Molecular Phylogenetics and Evolution</i> , 2021, 160, 107109.	1.2	17
2946	A novel reference dated phylogeny for the genus <i>Spodoptera</i> Guenée (Lepidoptera: Noctuidae): Tj ETQq1 1 0.784314 rgBT/Overlook Evolution, 2021, 161, 107161.	1.2	30
2947	Biological control activity of <i>Trichoderma asperelloides</i> PSU-P1 against gummy stem blight in muskmelon ( <i>Cucumis melo</i> ). <i>Physiological and Molecular Plant Pathology</i> , 2021, 115, 101663.	1.3	17
2948	Novel saprobic <i>Hermatomyces</i> species (Hermatomycetaceae, Pleosporales) from China (Yunnan) Tj ETQq1 1 0.784314 rgBT/Overlook 0.8 8	0.8	8
2949	Accuracy in Near-Perfect Virus Phylogenies. <i>Systematic Biology</i> , 2022, 71, 426-438.	2.7	8
2950	Taxonomic revision of the <i>Peyssonneliales</i> (Rhodophyta): Circumscribing the authentic <i>Peyssonnelia</i> clade and proposing four new genera and seven new species. <i>Journal of Phycology</i> , 2021, 57, 1749-1767.	1.0	9
2951	New species of <i>Yamadazyma</i> from rotting wood in China. <i>MycKeys</i> , 2021, 83, 69-84.	0.8	5
2952	An integrative analysis uncovers a new, pseudo-cryptic species of Amazonian marmoset (Primates): Tj ETQq1 1 0.784314 rgBT/Overlook 1.6 17	1.6	17
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3089	Species of <i>Hymenochaete</i> (Hymenochaetales, Basidiomycota) on bamboos from East Asia, with descriptions of two new species. <i>MycKeys</i> , 0, 20, 51-65.	0.8	6
3090	High diversity of <i>Diaporthe</i> species associated with dieback diseases in China, with twelve new species described. <i>MycKeys</i> , 2018, 39, 97-149.	0.8	83
3091	<i>Apophysomyces thailandensis</i> (Mucorales, Mucoromycota), a new species isolated from soil in northern Thailand and its solubilization of non-soluble minerals. <i>MycKeys</i> , 2019, 45, 75-92.	0.8	12
3092	<i>Neostagonosporella sichuanensis</i> gen. et sp. nov. (Phaeosphaeriaceae, Pleosporales) on <i>Phyllostachys heteroclada</i> (Poaceae) from Sichuan Province, China. <i>MycKeys</i> , 2019, 46, 119-150.	0.8	17
3093	Four new species of <i>Tremella</i> (Tremellales, Basidiomycota) based on morphology and DNA sequence data. <i>MycKeys</i> , 2019, 47, 75-95.	0.8	10
3094	Species of <i>Dendrostoma</i> (Erythrogloeaceae, Diaporthales) associated with chestnut and oak canker diseases in China. <i>MycKeys</i> , 2019, 48, 67-96.	0.8	22
3095	Striatiguttulaceae, a new pleosporalean family to accommodate <i>Longicorpus</i> and <i>Striatiguttula</i> gen. nov. from palms. <i>MycKeys</i> , 2019, 49, 99-129.	0.8	15
3096	Morphology and phylogeny reveal two novel <i>Coryneum</i> species from China. <i>MycKeys</i> , 2019, 56, 67-80.	0.8	4
3097	<i>Clitopilus lampangensis</i> (Agaricales, Entolomataceae), a new species from northern Thailand. <i>MycKeys</i> , 2019, 58, 69-82.	0.8	4
3098	Diaporthalean fungi associated with canker and dieback of trees from Mount Dongling in Beijing, China. <i>MycKeys</i> , 2019, 59, 67-94.	0.8	12
3099	The genus <i>Simplicillium</i> . <i>MycKeys</i> , 2019, 60, 69-92.	0.8	34
3100	Taxonomy of two synnematal fungal species from <i>Rhus chinensis</i> , with <i>Flavignomonium</i> gen. nov. described. <i>MycKeys</i> , 2019, 60, 17-29.	0.8	5
3101	Identification of six <i>Cytospora</i> species on Chinese chestnut in China. <i>MycKeys</i> , 2020, 62, 1-25.	0.8	24
3102	Discovery of <i>Cytospora</i> species associated with canker disease of tree hosts from Mount Dongling of China. <i>MycKeys</i> , 2020, 62, 97-121.	0.8	14
3103	Lignicolous freshwater ascomycota from Thailand: Phylogenetic and morphological characterisation of two new freshwater fungi: <i>Tingoldiigo hydei</i> sp. nov. and <i>T. clavata</i> sp. nov. from Eastern Thailand. <i>MycKeys</i> , 2020, 65, 119-138.	0.8	7
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3105	Threats from wildlife trade: The importance of genetic data in safeguarding the endangered Four-eyed Turtle ( <i>Sacalia quadriocellata</i> ). <i>Nature Conservation</i> , 0, 41, 91-111.	0.0	10

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3107	A new quadrannulate species of <i>Orobdella</i> (Hirudinida, Arhynchobdellida, Orobdellidae) from western Honshu, Japan. <i>ZooKeys</i> , 2016, 553, 33-51.	0.5	8
3108	Taxonomic reassessment of two subspecies of Chinese skink in Taiwan based on morphological and molecular investigations (Squamata, Scincidae). <i>ZooKeys</i> , 2017, 687, 131-148.	0.5	7
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3110	Evolutionary relationships and population genetics of the Afrotropical leaf-nosed bats (Chiroptera, Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50).	0.5	18
3111	Integrative taxonomy reveals three new taxa within the <i>Tylotriton asperrimus</i> complex (Caudata, Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50).	0.5	12
3112	A revised taxonomy of Asian snail-eating snakes <i>Pareas</i> (Squamata, Pareidae): evidence from morphological comparison and molecular phylogeny. <i>ZooKeys</i> , 2020, 939, 45-64.	0.5	15
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3115	Four new species of the genus <i>Orobdella</i> from Shikoku and Awajishima island, Japan (Hirudinida, Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50).	0.4	10
3116	Molecular phylogenetic analysis of a taxonomically unstable ranid from Sumatra, Indonesia, reveals a new genus with gastromyzophorous tadpoles and two new species. <i>Zoosystematics and Evolution</i> , 2018, 94, 163-193.	0.4	14
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3118	Outcomes Assessment Methodology for a Course in Probability and Random Variables. <i>Journal of Assessment and Institutional Effectiveness</i> , 2018, 8, 71-85.	0.4	1
3119	Genetic Characterization of Porcine Circovirus Type 2 from Pigs with Porcine Circovirus Associated Diseases in Argentina. <i>ISRN Veterinary Science</i> , 2011, 2011, 1-6.	1.1	9
3120	Ascospore Infection and <i>Colletotrichum</i> Species Causing Glomerella Leaf Spot of Apple in Uruguay. <i>Plant Pathology Journal</i> , 2019, 35, 100-111.	0.7	14
3121	First Record of <i>Orobdella tsushimensis</i> (Hirudinida: Arhynchobdellida: Gastrostomobdellidae) from the Korean Peninsula and Molecular Phylogenetic Relationships of the Specimens. <i>Animal Systematics, Evolution and Diversity</i> , 2014, 30, 87-94.	0.2	10
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3123	Next-generation sequencing of mixed genomic DNA allows efficient assembly of rearranged mitochondrial genomes in <i>Amolops chunganensis</i> and <i>Quasipaa boulengeri</i> . <i>PeerJ</i> , 2016, 4, e2786.	0.9	27

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3132	Characterizing gene tree conflict in plastome-inferred phylogenies. PeerJ, 2019, 7, e7747.	0.9	91
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3134	Novel <i>Neoacanthostigma</i> Species from Aquatic Habitats. Cryptogamie, Mycologie, 2017, 38, 169-190.	0.2	12
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3136	Phylogeny of a new ciliate family Clampidae fam. nov. (Protista: Ciliophora), with notes on morphology and morphogenesis. Zoological Journal of the Linnean Society, 2022, 196, 88-104.	1.0	4
3137	Establishment of <i>Alloptilella splendida</i> gen. et sp. nov. and resurrection of <i>Scytalium veneris</i> (Thomson & Henderson, 1906), two sea pens (Cnidaria: Pennatulacea) from seamounts in the tropical Western Pacific. Journal of Oceanology and Limnology, 2021, 39, 1790-1804.	0.6	6
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3140	Pleistocene origin and colonization history of <i>Lobelia columnaris</i> Hook. f. (Campanulaceae:) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5 2019, 7, e7747.	0.8	1
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3172	Morpho-molecular Characterization of the Litostomatean Predatory Ciliate Phialina pupula (Müller), Tj ETQq1 1 0,784314 rgBT /Overlo	0.5	5
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3186	A novel clade of bat-associated Bartonella detected in the bat fly Leptocyclopodia ferrari (Diptera: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 Microbiology, 2022, 264, 109284.	0.8	5
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3208	DNA barcoding identification of Greek freshwater fishes. <i>PLoS ONE</i> , 2022, 17, e0263118.	1.1	13
3210	New species of <i>Colletotrichum</i> from wild Poaceae and Cyperaceae plants in Iran. <i>Mycologia</i> , 2022, 114, 89-113.	0.8	6
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3215	Survey of <i>Volvariella</i> (Agaricales, Basidiomycota) including Two New Species, <i>V. neovolvacea</i> and <i>V. thailandensis</i> , from Northern Thailand. <i>Diversity</i> , 2022, 14, 161.	0.7	0
3216	Distribution and characterization of <i>Colletotrichum</i> species associated with Citrus anthracnose in eastern Mediterranean region of Turkey. <i>European Journal of Plant Pathology</i> , 2022, 163, 125-141.	0.8	7
3217	<i>Paraconiothyrium fuckelii</i> , <i>Diaporthe eres</i> and <i>Neocosmospora parceramosa</i> causing cane blight of red raspberry in Northern Italy. <i>Journal of Plant Pathology</i> , 2022, 104, 683-698.	0.6	7
3218	Redescription of the giant Southeast Asian millipede <i>Spirobolus macrurus</i> Pocock, 1893 and its assignment to the new genus <i>Macrurobolus</i> gen. nov. (Diplopoda, Spirobolida, Pachybolidae). <i>ZooKeys</i> , 2022, 1087, 1-18.	0.5	1
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3227	Mitogenomic phylogenetics and population genetics of several taxa of agouties (Dasyprocta sp.) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 1.	0.6	1
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3381	Evaluation of Statistical Reliability and Model Assumptions. <i>SpringerBriefs in Statistics</i> , 2022, , 49-57.	0.3	0
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3387	An Evolutionary Framework of Acanthaceae Based on Transcriptomes and Genome Skims. <i>Systematic Botany</i> , 2022, 47, 716-728.	0.2	0
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3389	Data Integration in Bayesian Phylogenetics. <i>Annual Review of Statistics and Its Application</i> , 2023, 10, 353-377.	4.1	2
3390	Characterization of <i>Clinostomum</i> (Digenea: Clinostomidae) spp. in India. <i>Parasitology Research</i> , 2022, 121, 3083-3089.	0.6	2
3391	A new species of land planarian split off from <i>Luteostriata ernesti</i> (Leal-Zanchet & ) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 1-15.	0.5	0
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3399	Morphology and morphogenesis of the hypotrich ciliate <i>Parentocirrus hortualis</i> Vol <sup>2</sup> , 1997, with notes on the phylogeny of <i>Parentocirrus</i> (Ciliophora, Hypotrichia). <i>European Journal of Protistology</i> , 2022, 86, 125936.	0.5	2
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3401	Insight into the Taxonomic Resolution of <i>Apiospora</i> : Introducing Novel Species and Records from Bamboo in China and Thailand. <i>Diversity</i> , 2022, 14, 918.	0.7	5
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3404	A New Species of the Genus <i>Microhyla</i> (Amphibia: Anura: Microhylidae) from the Dabie Mountains, China. <i>Animals</i> , 2022, 12, 2894.	1.0	1
3405	<i>Diophrys pauciciliata</i> n. sp. (Ciliophora, Euplotida), a new marine spirotrich ciliate with highly static morphometrics. <i>European Journal of Protistology</i> , 2022, 86, 125937.	0.5	1
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3408	Avian surfactant protein (SP)-A2 first arose in an early tetrapod before the divergence of amphibians and gradually lost the collagen domain. <i>Developmental and Comparative Immunology</i> , 2023, 139, 104582.	1.0	2
3409	The taxonomic revision of <i>Melanogaster</i> (Paxillaceae, Boletales) in China based on molecular and morphological evidence. <i>Mycological Progress</i> , 2022, 21, .	0.5	1
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3419	Employing tree bisection and reconnection rearrangement for parsimony inference in MPBoot. , 2022, , .		0
3420	<i>Hypotrichidium tisiae</i> (Gelei, 1929) Gelei, 1954: a unique hypotrichid ciliate having a highly specialized developmental pattern during binary division. <i>Marine Life Science and Technology</i> , 2022, 4, 536-550.	1.8	8



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3424	Phylogenomic analysis of <i>Citrobacter</i> sp. strain AAK_AS5 and its metabolic capabilities to support nitrogen removal behavior. <i>Journal of Basic Microbiology</i> , 0, , .	1.8	1
3425	Description of a New Cobra ( <i>Naja Laurenti</i> , 1768; Squamata, Elapidae) from China with Designation of a Neotype for <i>Naja atra</i> . <i>Animals</i> , 2022, 12, 3481.	1.0	4
3426	Genetic Analysis of Torque Teno Canis Virus Identified in Republic of Korea. <i>Veterinary Sciences</i> , 2022, 9, 693.	0.6	1
3427	Morphological Characteristics and Phylogeny Reveal Six New Species in <i>Russula</i> Subgenus <i>Russula</i> ( <i>Russulaceae</i> , <i>Russulales</i> ) from Yanshan Mountains, North China. <i>Journal of Fungi (Basel)</i> , 2022, 8, 1010.	1.0	50
3428	High-Density Genetic Linkage Map of the Southern Blue-ringed Octopus ( <i>Octopodidae</i> : <i>Hapalochlaena</i> )	0.7	2
3429	Morphological characterization and pathogenicity of <i>Colletotrichum aenigma</i> and <i>C. siamense</i> causing anthracnose on <i>Euonymus japonicus</i> in Beijing, China. <i>Plant Pathology</i> , 2023, 72, 430-441.	1.2	2
3430	To explore strange new worlds – The diversification in <i>Tremella caloplacae</i> was linked to the adaptive radiation of the <i>Teloschistaceae</i> . <i>Molecular Phylogenetics and Evolution</i> , 2023, 180, 107680.	1.2	2
3431	Not all bad: Gyromitrin has a limited distribution in the false morels as determined by a new ultra high-performance liquid chromatography method. <i>Mycologia</i> , 2023, 115, 1-15.	0.8	3
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3434	Molecular Detection of <i>Rickettsia</i> and Other Bacteria in Ticks and Birds in an Urban Fragment of Tropical Dry Forest in Magdalena, Colombia. <i>Life</i> , 2023, 13, 145.	1.1	2
3435	Identification of Small GTPases That Phosphorylate IRF3 through TBK1 Activation Using an Active Mutant Library Screen. <i>Biomolecules and Therapeutics</i> , 2023, 31, 48-58.	1.1	3
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3438	Integrative taxonomy reveals that not all European reddish runcinids are the same: the case of the <i>Runcina ferruginea</i> Kress, 1977 ( <i>Gastropoda</i> , <i>Heterobranchia</i> , <i>Runcinida</i> ) species-complex, with the description of a new genus. <i>Invertebrate Systematics</i> , 2023, 37, 61.	0.5	0

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3441	Review of the Mimetic Subgroup of <i>Culex</i> ( <i>Culex</i> ) (Diptera: Culicidae), With an Assessment of the Specific Status of Three Nominal Species Described From India. <i>Journal of Medical Entomology</i> , 0, , .	0.9	0
3442	<i>Cytospora parapleurivora</i> sp. nov. isolated from orchards with fruit tree decline syndrome in Ontario, Canada. <i>PLoS ONE</i> , 2023, 18, e0279490.	1.1	3
3443	Taxonomic review of <i>Saguinus mystax</i> (Spix, 1823) (Primates, Callitrichidae), and description of a new species. <i>PeerJ</i> , 0, 11, e14526.	0.9	3
3444	Re-discovery and novel contributions to morphology and multigene phylogeny of <i>Protospirella mazurica</i> (Raabe, 1968) Aesch, 2001 (Ciliophora: Pleuronematida), an obligate symbiont of the river nerite <i>Theodoxus fluviatilis</i> Linnaeus, 1758 (Mollusca: Gastropoda). <i>European Journal of Protistology</i> , 2023, 88, 125956.	0.5	1
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3446	<i>Villoboletus persicus</i> , gen. et sp. nov. (Boletaceae), a bolete with flocculent-covered stipe from northern China. <i>Mycologia</i> , 2023, 115, 255-262.	0.8	1
3447	Taxonomic and phylogenetic approach to some Antarctic lichenicolous fungi. <i>Mycological Progress</i> , 2023, 22, .	0.5	0
3448	Traditional taxonomy underestimates the number of species of <i>Bokermannohyla</i> (Amphibia: Tj ETQq1 1 0.784314 rgBT /Overl Biodiversity, 2023, 21, .	0.5	0
3449	Functional genome annotation and transcriptome analysis of <i>Pseudozyma hubeiensis</i> BOT-O, an oleaginous yeast that utilizes glucose and xylose at equal rates. <i>Fungal Genetics and Biology</i> , 2023, 166, 103783.	0.9	1
3450	Diversification of the African legless skinks in the subfamily Acontinae (Family Scincidae). <i>Molecular Phylogenetics and Evolution</i> , 2023, 182, 107747.	1.2	1
3451	The systematics and evolution of the Sri Lankan rainforest land snail <i>Corilla</i> : New insights from RADseq-based phylogenetics. <i>Molecular Phylogenetics and Evolution</i> , 2023, 182, 107731.	1.2	0
3452	Metabolic regulation mechanism of <i>Trametes gibbosa</i> CB1 on lignin. <i>International Journal of Biological Macromolecules</i> , 2023, 240, 124189.	3.6	2
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3459	A new species of <i>Dipsas</i> (Serpentes, Dipsadidae) from central Panama. <i>ZooKeys</i> , 0, 1145, 131-167.	0.5	0
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3469	A Case Report on a Human Bite Contact with a Rabid Honey Badger <i>Mellivora capensis</i> (Kromdraai Area, Tj ETQq1 1 0.784314 rgBT /Ov	0.9	1
3470	Addition of three new lineages in Mycosphaerellaceae: <i>Neoacervuloseptoria</i> gen. nov., <i>Neocercospora</i> gen. nov. and <i>Neoramulariopsis</i> gen. nov.. <i>Mycological Progress</i> , 2023, 22, .	0.5	0
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3478	Morphologic and molecular characterization of <i>Apertospathula pilata</i> n. sp., a novel freshwater spathidiid (Ciliophora, Litostomatea) from Idaho, USA. <i>European Journal of Protistology</i> , 2023, 89, 125990.	0.5	1
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