

Chao-Dong Zhu

List of Publications by Year in descending order

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#	ARTICLE	IF	CITATIONS
1	Phylogenetic relatedness, functional traits, and spatial scale determine herbivore co-occurrence in a subtropical forest. <i>Ecological Monographs</i> , 2022, 92, e01492.	5.4	8
2	Chromosome-Level Genome Assembly of <i>Anthidium xuezhongi</i> Niu & Zhu, 2020 (Hymenoptera: Apoidea: Tj ETQq0,0,0 rgBT /Overlock 1	2.5	1
3	Testing the systematic status of <i>Homalictus</i> and <i>Rostrohalictus</i> with weakened crossâ€in groups within Halictini (Hymenoptera: Halictidae) using lowâ€coverage wholeâ€genome sequencing. <i>Insect Science</i> , 2022, 29, 1819-1833.	3.0	7
4	Patterns of endemism of the Chinese Larentiinae (Lepidoptera: Geometridae). <i>Archives of Insect Biochemistry and Physiology</i> , 2022, 111, e21891.	1.5	1
5	Notes on the Genus <i>Aceratoneuromyia</i> Girault (Hymenoptera: Eulophidae). <i>Insects</i> , 2022, 13, 450.	2.2	1
6	Key to <i>Ooctonus</i> Haliday (Hymenoptera: Mymaridae) in China, with one new species and three new country records. <i>Zootaxa</i> , 2022, 5155, 581-588.	0.5	0
7	The First Biological Portrait of Stalk-Eyed Fruit Flies: Life History, Reproductive Biology and Host Use Patterns in <i>Pelmatops</i> spp. (Diptera: Tephritidae). <i>Annals of the Entomological Society of America</i> , 2022, 115, 365-377.	2.5	2
8	Global Patterns and Drivers of Bee Distribution. <i>Current Biology</i> , 2021, 31, 451-458.e4.	3.9	155
9	Taxonomy must engage with new technologies and evolve to face future challenges. <i>Nature Ecology and Evolution</i> , 2021, 5, 3-4.	7.8	25
10	Relationships between wild bee abundance and fruit set of <i>Camellia oleifera</i> Abel. <i>Journal of Applied Entomology</i> , 2021, 145, 277-285.	1.8	4
11	Working landscapes need at least 20% native habitat. <i>Conservation Letters</i> , 2021, 14, e12773.	5.7	116
12	Phylogeny of Adramini (Diptera, Tephritidae) based on integrative evidence. <i>Zoologica Scripta</i> , 2021, 50, 71-83.	1.7	2
13	Comparison of two criteria on the essential number calculation of <i>Andrena camellia</i> . <i>Bulletin of Entomological Research</i> , 2021, 111, 364-370.	1.0	2
14	Ultraconserved element phylogenomics and biogeography of the agriculturally important mason bee subgenus <i>Osmia</i> (<i>Osmia</i>). <i>Systematic Entomology</i> , 2021, 46, 453-472.	3.9	25
15	The complete mitochondrial genome of <i>Metaphycus eriococci</i> (Timberlake) (Hymenoptera: Encyrtidae). <i>Mitochondrial DNA Part B: Resources</i> , 2021, 6, 550-552.	0.4	0
16	Investigating the Parasitoid Community Associated with the Invasive Mealybug <i>Phenacoccus solenopsis</i> in Southern China. <i>Insects</i> , 2021, 12, 290.	2.2	11
17	Tree diversity and functional leaf traits drive herbivoreâ€associated microbiomes in subtropical China. <i>Ecology and Evolution</i> , 2021, 11, 6153-6166.	1.9	1
18	Multi-trophic communities re-establish with canopy cover and microclimate in a subtropical forest biodiversity experiment. <i>Oecologia</i> , 2021, 196, 289-301.	2.0	12

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19	<i>Liomelita mollipalma</i> , a new genus and species of Melitidae (Amphipoda: Hadzioidea) from hydrothermal vents of the Okinawa Trough, North-West Pacific. <i>Journal of Natural History</i> , 2021, 55, 1299-1310.	0.5	0
20	Tree diversity promotes predatory wasps and parasitoids but not pollinator bees in a subtropical experimental forest. <i>Basic and Applied Ecology</i> , 2021, 53, 134-142.	2.7	8
21	Sampling biases shape our view of the natural world. <i>Ecography</i> , 2021, 44, 1259-1269.	4.5	190
22	The complete mitochondrial genome of <i>Cerceris quinquefasciata</i> (Hymenoptera: Crabronidae). <i>Mitochondrial DNA Part B: Resources</i> , 2021, 6, 2044-2045.	0.4	0
23	The complete mitochondrial genome of <i>Cerceris bucculata</i> (Hymenoptera: Crabronidae). <i>Mitochondrial DNA Part B: Resources</i> , 2021, 6, 1959-1960.	0.4	0
24	Review of the bee genus <i>Pseudoanthidium</i> Friese, 1898 (Hymenoptera: Apoidea: Megachilidae: Anthidiini) of China with descriptions of three new species. <i>Zootaxa</i> , 2021, 4996, 133-152.	0.5	1
25	The cellophane bees of <i>Colletes Latreille</i> (Hymenoptera: Colletidae) from Xizang (Tibet), China. <i>Zootaxa</i> , 2021, 5022, 1-72.	0.5	1
26	The significance of tree-tree interactions for forest ecosystem functioning. <i>Basic and Applied Ecology</i> , 2021, 55, 33-52.	2.7	38
27	Reprint of: Tree diversity promotes predatory wasps and parasitoids but not pollinator bees in a subtropical experimental forest. <i>Basic and Applied Ecology</i> , 2021, 55, 124-132.	2.7	0
28	Comparison of the pollination efficiency of <i>Apis cerana</i> with wild bees in oil-seed camellia fields. <i>Basic and Applied Ecology</i> , 2021, 56, 250-258.	2.7	6
29	Morphological and molecular identification of arrhenotokous strain of <i>Diglyphus wani</i> (Hymenoptera, Eulophidae) found in China as a control agent against agromyzid leafminers. <i>ZooKeys</i> , 2021, 1071, 109-126.	1.1	6
30	Global invasion risk of <i>Apocephalus borealis</i> , a honey bee parasitoid. <i>Apidologie</i> , 2021, 52, 1128-1140.	2.0	0
31	A Simulation-Based Evaluation of Tip-Dating Under the Fossilized Birth-Death Process. <i>Systematic Biology</i> , 2020, 69, 325-344.	5.6	39
32	Description of a new species of the genus <i>Lebbeus</i> White, 1847 (Decapoda, Thoridae) from the Bohai Sea, China. <i>Crustaceana</i> , 2020, 93, 1393-1403.	0.3	1
33	Chinese species of <i>Nomia</i> (<i>Gnathonomia</i>) Pauly, 2005 (Hymenoptera: Tj ETQq1 1,0,784314 rgBT /Ove	0.5	2
34	The First Draft Genome of the Plasterer Bee <i>Colletes gigas</i> (Hymenoptera: Colletidae: Colletes). <i>Genome Biology and Evolution</i> , 2020, 12, 860-866.	2.5	12
35	Host functional and phylogenetic composition rather than host diversity structure plant-herbivore networks. <i>Molecular Ecology</i> , 2020, 29, 2747-2762.	3.9	24
36	Nesting and foraging behavior of <i>Xylocopa valga</i> in the Ejina Oasis, China. <i>PLoS ONE</i> , 2020, 15, e0235769.	2.5	2

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37	Three questions: How can taxonomists survive and thrive worldwide?. <i>Megataxa</i> , 2020, 1, .	3.8	26
38	A new species of <i>Oomyzus Rondani</i> (Hymenoptera, Eulophidae) reared from the pupae of <i>Coccinella septempunctata</i> (Coleoptera, Coccinellidae) in China. <i>ZooKeys</i> , 2020, 953, 49-60.	1.1	6
39	Foraging Behavior of Honeybees (<i>Apis Mellifera</i> L.) and Ground Bumblebees (<i>Bombus Terrestris</i> L.) and its Influence on Seed Yield and Oil Quality of Oil Tree Peony Cultivar ‘Fengdan’™ (<i>Paeonia Ostii</i> T. Hong) <i>Tj ETQq1 1 0.784314 rgBT /Overlo</i>	1.1	8
40	Multiple components of plant diversity loss determine herbivore phylogenetic diversity in a subtropical forest experiment. <i>Journal of Ecology</i> , 2019, 107, 2697-2712.	4.0	33
41	The mitochondrial genome of <i>Platencyrtus parkeri</i> Feriere (Hymenoptera: Encyrtidae). <i>Mitochondrial DNA Part B: Resources</i> , 2019, 4, 3479-3481.	0.4	2
42	Overview of the bee genus <i>Trachusa</i> Panzer, 1804 (Hymenoptera: Apoidea: Megachilidae: Anthidiini) from China with description of three new species. <i>Zootaxa</i> , 2019, 4646, 251-270.	0.5	1
43	Revision of the bee genus <i>Bathanthidium</i> Mavromoustakis (Hymenoptera: Apoidea: Megachilidae.) <i>Tj ETQq1 1 0.784314 rgBT /Overlo</i>	0.5	3
44	A High-quality Draft Genome Assembly of <i>Sinella curviseta</i> : A Soil Model Organism (Collembola). <i>Genome Biology and Evolution</i> , 2019, 11, 521-530.	2.5	13
45	Molecular signatures of X chromosome inactivation and associations with clinical outcomes in epithelial ovarian cancer. <i>Human Molecular Genetics</i> , 2019, 28, 1331-1342.	2.9	19
46	Phylogenomics from low-coverage whole-genome sequencing. <i>Methods in Ecology and Evolution</i> , 2019, 10, 507-517.	5.2	59
47	Description of <i>Seba longimera</i> sp. nov. from hydrothermal vents in the Okinawa Trough, Northwest Pacific (Amphipoda, Amphilochoidea, Sebidae). <i>ZooKeys</i> , 2019, 899, 141-149.	1.1	1
48	Life history and biocontrol potential of the first female-producing parthenogenetic species of <i>Diglyphus</i> (Hymenoptera: Eulophidae) against agromyzid leafminers. <i>Scientific Reports</i> , 2018, 8, 3222.	3.3	16
49	Comparison of Methods for Molecular Species Delimitation Across a Range of Speciation Scenarios. <i>Systematic Biology</i> , 2018, 67, 830-846.	5.6	277
50	Intra- and interspecific tree diversity promotes multitrophic plant-ant interactions in a forest diversity experiment. <i>Basic and Applied Ecology</i> , 2018, 29, 89-97.	2.7	9
51	Parasitoid-host associations of the genus <i>Coccophagus</i> (Hymenoptera: Aphelinidae) in China. <i>Zoological Journal of the Linnean Society</i> , 2018, 182, 38-49.	2.3	7
52	Genetic Variability of <i>Melipona subnitida</i> (Hymenoptera: Apidae) in Introduced and Native Populations. <i>Journal of Insect Science</i> , 2018, 18, .	1.5	1
53	New records of bees of the genus <i>Sphecodes</i> Latreille in the Palaearctic part of China (Hymenoptera,) <i>Tj ETQq1 1 0.784314 rgBT /Overlo</i>	1.1	8
54	Nesting Biology of <i>Xylocopa xinjiangensis</i> (Hymenoptera: Apidae: Xylocopinae). <i>Journal of Insect Science</i> , 2018, 18, .	1.5	1

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55	Impacts of species richness on productivity in a large-scale subtropical forest experiment. <i>Science</i> , 2018, 362, 80-83.	12.6	433
56	Host specificity of parasitoids (Encyrtidae) toward armored scale insects (Diaspididae): Untangling the effect of cryptic species on quantitative food webs. <i>Ecology and Evolution</i> , 2018, 8, 7879-7893.	1.9	10
57	A review of the genus <i>Triancyra</i> Baltazar (Ichneumonidae: Rhyssinae) from Vietnam, with descriptions of three new species. <i>Zootaxa</i> , 2018, 4377, 565.	0.5	2
58	Biodiversity across trophic levels drives multifunctionality in highly diverse forests. <i>Nature Communications</i> , 2018, 9, 2989.	12.8	169
59	Sequencing and characterization of the <i>Megachile sculpturalis</i> (Hymenoptera: Megachilidae) mitochondrial genome. <i>Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis</i> , 2017, 28, 344-346.	0.7	8
60	Chinese species of <i>Pediobius</i> Walker (Hymenoptera: Eulophidae). <i>Zootaxa</i> , 2017, 4240, 1-71.	0.5	12
61	Toward a methodical framework for comprehensively assessing forest multifunctionality. <i>Ecology and Evolution</i> , 2017, 7, 10652-10674.	1.9	41
62	Construction, implementation and testing of an image identification system using computer vision methods for fruit flies with economic importance (Diptera: Tephritidae). <i>Pest Management Science</i> , 2017, 73, 1511-1528.	3.4	14
63	Notes on <i>Kocourekia Bouček</i> (Hymenoptera: Eulophidae: Tetrastichinae) with description of a new species from China. <i>Zootaxa</i> , 2017, 4317, 391.	0.5	3
64	Two new genera and three new species of <i>Epipaschiinae</i> Meyrick from China (Lepidoptera, Pyralidae). <i>ZooKeys</i> , 2017, 722, 95-107.	1.1	1
65	The complete mitochondrial genome of <i>Acleris fimbriana</i> (Lepidoptera: Tortricidae). <i>Mitochondrial DNA</i> , 2016, 27, 1-3.	0.6	5
66	The complete mitochondrial genome of <i>Choristoneura longicellana</i> (Lepidoptera: Tortricidae) and phylogenetic analysis of Lepidoptera. <i>Gene</i> , 2016, 591, 161-176.	2.2	40
67	Tree phylogenetic diversity promotes host-parasitoid interactions. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2016, 283, 20160275.	2.6	41
68	Sequencing and characterization of the <i>Megachile strupigera</i> (Hymenoptera: Megachilidae) mitochondrial genome. <i>Mitochondrial DNA Part B: Resources</i> , 2016, 1, 282-284.	0.4	6
69	Revision of the <i>Anthidiellum</i> Cockerell, 1904 of China (Hymenoptera, Apoidea). <i>Tj ETQq1</i> 1 0.784314 rgBT / Qv 0.5	0.5	6
70	Descriptions of three new species of <i>Dzhanokmenia</i> (Hymenoptera: Eulophidae) from China. <i>Zootaxa</i> , 2016, 4121, 447.	0.5	4
71	Formal nomenclature and description of cryptic species of the <i>Encyrtus sasakii</i> complex (Hymenoptera: Encyrtidae). <i>Scientific Reports</i> , 2016, 6, 34372.	3.3	13
72	China: Change tack to boost basic research. <i>Nature</i> , 2016, 536, 30-30.	27.8	0

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73	The complete mitochondrial genome of the <i>Colletes gigas</i> (Hymenoptera: Colletidae). <i>Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf</i>	0.7	12
74	The complete mitochondrial genome of <i>Carposina sasakii</i> (Lepidoptera: Carposinidae). <i>Mitochondrial DNA</i> , 2016, 27, 1432-1434.	0.6	7
75	A simulation study of sample size for DNA barcoding. <i>Ecology and Evolution</i> , 2015, 5, 5869-5879.	1.9	28
76	Molecular phylogeny and the underestimated species diversity of the endemic white-bellied rat (Rodentia: Muridae: <i>Niviventer</i>) in Southeast Asia and China. <i>Zoologica Scripta</i> , 2015, 44, 475-494.	1.7	22
77	A systematic study of <i>Ichneumonosoma</i> de Meijere, <i>Pelmatops</i> Enderlein, <i>Pseudopelmatops</i> Shiraki and <i>Soita</i> Walker (Diptera: Tephritidae). <i>Zootaxa</i> , 2015, 4013, 301-47.	0.5	4
78	Resolving the phylogeny of a speciose spider group, the family Linyphiidae (Araneae). <i>Molecular Phylogenetics and Evolution</i> , 2015, 91, 135-149.	2.7	23
79	Observational natural history and morphological taxonomy are indispensable for future challenges in biodiversity and conservation. <i>Communicative and Integrative Biology</i> , 2015, 8, e992745.	1.4	3
80	A DNA Barcoding system integrating multigene sequence data. <i>Methods in Ecology and Evolution</i> , 2015, 6, 930-937.	5.2	15
81	gPGA: GPU Accelerated Population Genetics Analyses. <i>PLoS ONE</i> , 2015, 10, e0135028.	2.5	5
82	A Unique Nest-Protection Strategy in a New Species of Spider Wasp. <i>PLoS ONE</i> , 2014, 9, e101592.	2.5	15
83	Two new <i>Aprostocetus</i> species (Hymenoptera: Eulophidae: Tetrastichinae), fortuitous parasitoids of invasive eulophid gall inducers (Tetrastichinae) on <i>Eucalyptus</i> and <i>Erythrina</i> . <i>Zootaxa</i> , 2014, 3846, 261-72.	0.5	17
84	A new phytophagous eulophid wasp (Hymenoptera: Chalcidoidea: Eulophidae) that feeds within leaf buds and cones of <i>Pinus massoniana</i> . <i>Zootaxa</i> , 2014, 3753, 391.	0.5	2
85	Bees of the <i>Colletes flavicornis</i> -group from China with description of one new species (Hymenoptera: Apoidea: Colletidae). <i>Zootaxa</i> , 2014, 3780, 534.	0.5	7
86	A new species of <i>Zagrammosoma</i> Ashmead (Hymenoptera, Eulophidae) from Qinghai Province, China. <i>ZooKeys</i> , 2014, 417, 45-55.	1.1	1
87	Global and Local Persistence of Influenza A(H5N1) Virus. <i>Emerging Infectious Diseases</i> , 2014, 20, 1287-1295.	4.3	49
88	Preliminary phylogeny of the genus <i>Copidosoma</i> (Hymenoptera). <i>Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 147 Td</i> (2014, 39, 325-334).	3.9	6
89	Cryptic diversity, diversification and vicariance in two species complexes of <i>Tomocerus</i> (<i>Tomocerus ollembola</i> , <i>Tomocerus</i>) from China. <i>Zoologica Scripta</i> , 2014, 43, 393-404.	1.7	36
90	A Protocol for Species Delineation of Public DNA Databases, Applied to the Insecta. <i>Systematic Biology</i> , 2014, 63, 712-725.	5.6	8

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91	The complete mitochondrial genome of the endangered Apollo butterfly, <i>Parnassius apollo</i> (Lepidoptera: Papilionidae) and its comparison to other Papilionidae species. <i>Journal of Asia-Pacific Entomology</i> , 2014, 17, 663-671.	0.9	26
92	Application of DNA barcoding to the identification of Hymenoptera parasitoids from the soybean aphid (<i>Aphis glycines</i>) in China. <i>Insect Science</i> , 2014, 21, 363-373.	3.0	13
93	Molecular phylogeny reveals independent origins of body scales in Entomobryidae (Hexapoda: Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 22	2.7	47
94	The Bees of the Genus <i>Colletes</i> (Hymenoptera: Apoidea: Colletidae) from China. <i>Zootaxa</i> , 2014, 3856, 451-83.	0.5	7
95	Heuristic optimization for global species clustering of <sc>DNA</sc> sequence data from multiple loci. <i>Methods in Ecology and Evolution</i> , 2013, 4, 961-970.	5.2	2
96	Hepatitis B virus subgenotyping: History, effects of recombination, misclassifications, and corrections. <i>Infection, Genetics and Evolution</i> , 2013, 16, 355-361.	2.3	89
97	Description of <i>Synergus castaneus</i> n. sp. (Hymenoptera: Cynipidae: Synergini) Associated with an Unknown Gall on <i>Castanea</i> spp. (Fagaceae) in China. <i>Annals of the Entomological Society of America</i> , 2013, 106, 437-446.	2.5	17
98	A review of the genus <i>Monema</i> Walker in China (Lepidoptera, Limacodidae). <i>ZooKeys</i> , 2013, 306, 23-36.	1.1	4
99	<p>Bees of the Colletes clypearis-group (Hymenoptera: Apoidea: Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 22	0.5	7
100	Quantifying Species Diversity with a DNA Barcoding-Based Method: Tibetan Moth Species (Noctuidae) on the Qinghai-Tibetan Plateau. <i>PLoS ONE</i> , 2013, 8, e64428.	2.5	15
101	Characterization of the Complete Mitochondrion Genome of Diurnal Moth <i>Amata emma</i> (Butler) (Lepidoptera: Erebidae) and Its Phylogenetic Implications. <i>PLoS ONE</i> , 2013, 8, e72410.	2.5	62
102	<p class="HeadingRunIn">New species and records of Trypetinae (Diptera: Tephritidae) from China</p>. <i>Zootaxa</i> , 2013, 3710, 333.	0.5	2
103	<p class="HeadingRunIn">First record of the bee genus Homalictus Cockerell for China with description of a new species (Hymenoptera: Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 22	0.5	7
104	A review of the Colletes succinctus-group (Hymenoptera: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 22	0.5	11
105	<p class="HeadingRunIn">XIAO-LIN CHEN, XIN-JIAN WANG & CHAO-DONG ZHU (2013) New species and records of Trypetinae (Diptera: Tephritidae) from China. Zootaxa, 3710(4), 333&hat{=}353.</p>. <i>Zootaxa</i> , 2013, 3718, 500.	0.5	0
106	The Mitochondrial Genome of <i>Elodia flavipalpis</i> Aldrich (Diptera: Tachinidae) and the Evolutionary Timescale of Tachinid Flies. <i>PLoS ONE</i> , 2013, 8, e61814.	2.5	51
107	Parallel Metropolis Coupled Markov Chain Monte Carlo for Isolation with Migration Model. <i>Applied Mathematics and Information Sciences</i> , 2013, 7, 219-224.	0.5	2
108	The Complete Mitochondrial Genome of <i>Leucoptera malifoliella</i> Costa (Lepidoptera: Lyonetiidae). <i>DNA and Cell Biology</i> , 2012, 31, 1508-1522.	1.9	28

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109	The Complete Mitochondrial Genome of the Rice Moth, <i>Corcyra cephalonica</i> . Journal of Insect Science, 2012, 12, 1-14.	1.5	23
110	Subgenotype reclassification of genotype B hepatitis B virus. BMC Gastroenterology, 2012, 12, 116.	2.0	20
111	A new species of Bathanthidium Mavromoustakis (Hymenoptera: Megachilidae: Anthidiini) from China, with a key to the species. Zootaxa, 2012, 3218, 59.	0.5	5
112	A review of Megachile (Chelostomoda) Michener (Megachilidae: Megachilini) known from China with the description of a new species. Zootaxa, 2012, 3267, 55.	0.5	0
113	Complete sequence of the mitochondrial genome of the Japanese buff-tip moth, <i>Phalera flavescens</i> (Lepidoptera: Notodontidae). Genetics and Molecular Research, 2012, 11, 4213-4225.	0.2	30
114	Complete mitochondrial genomes of two cockroaches, <i>Blattella germanica</i> and <i>Periplaneta americana</i> , and the phylogenetic position of termites. Current Genetics, 2012, 58, 65-77.	1.7	36
115	DNA barcoding of six <i>Ceroplastes</i> species (Hemiptera: Coccoidea: Coccidae) from China. Molecular Ecology Resources, 2012, 12, 791-796.	4.8	47
116	Identification of novel inter-genotypic recombinants of human hepatitis B viruses by large-scale phylogenetic analysis. Virology, 2012, 427, 51-59.	2.4	44
117	A fuzzy set theory based approach to analyse species membership in DNA barcoding. Molecular Ecology, 2012, 21, 1848-1863.	3.9	73
118	Phylogenetic Reconstruction and DNA Barcoding for Closely Related Pine Moth Species (<i>Dendrolimus</i>) in China with Multiple Gene Markers. PLoS ONE, 2012, 7, e32544.	2.5	48
119	The Integrative Taxonomic Approach Reveals Host Specific Species in an Encyrtid Parasitoid Species Complex. PLoS ONE, 2012, 7, e37655.	2.5	32
120	Recombination in Hepatitis C Virus: Identification of Four Novel Naturally Occurring Inter-Subtype Recombinants. PLoS ONE, 2012, 7, e41997.	2.5	27
121	Small Mammal Investigation in Spotted Fever Focus with DNA-Barcoding and Taxonomic Implications on Rodents Species from Hainan of China. PLoS ONE, 2012, 7, e43479.	2.5	16
122	Subgenotyping of Genotype C Hepatitis B Virus: Correcting Misclassifications and Identifying a Novel Subgenotype. PLoS ONE, 2012, 7, e47271.	2.5	26
123	Phylogeography of the mud crab (<i>Scylla serrata</i>) in the Indo-West Pacific reappraised from mitochondrial molecular and oceanographic clues: transoceanic dispersal and coastal sequential colonization. Marine Ecology, 2011, 32, 52-64.	1.1	11
124	DNA barcoding of endoparasitoid wasps in the genus <i>Anicetus</i> reveals high levels of host specificity (Hymenoptera: Encyrtidae). Biological Control, 2011, 58, 182-191.	3.0	26
125	The complete mitochondrial genome of <i>Spilonota lechriaspis</i> Meyrick (Lepidoptera: Tortricidae). Molecular Biology Reports, 2011, 38, 3757-3764.	2.3	46
126	Positive selection on hemagglutinin and neuraminidase genes of H1N1 influenza viruses. Virology Journal, 2011, 8, 183.	3.4	48

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127	Potential efficacy of mitochondrial genes for animal DNA barcoding: a case study using eutherian mammals. <i>BMC Genomics</i> , 2011, 12, 84.	2.8	83
128	Scanning electron microscopy studies of antennal sensilla of <i>Ooencyrtus phongi</i> (Hymenoptera: Encyrtidae). <i>Microscopy Research and Technique</i> , 2011, 74, 936-945.	2.2	18
129	A newly recorded Subgenus <i>Sudila</i> from China with description of two new species (Hymenoptera: Tj ETQq1 1 0.784314 rgBT /Overl 0.5	2.3	79
130	Genetic imprints of paleo-oceanographic conditions in the Chinese seas: Population bottlenecks of <i>Scylla paramamosain</i> and <i>Periophthalmus modestus</i> inferred from mitochondrial genes. <i>Journal of Earth Science (Wuhan, China)</i> , 2010, 21, 237-240.	3.2	2
131	The complete mitochondrial genome of the yellow coaster, <i>AcraeaÂïssoria</i> (Lepidoptera: Nymphalidae: Tj ETQq1 1 0.784314 rgBT /Over 2.3	2.3	79
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