

Xue-xin Chen

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219
papers

2,664
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242
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3,369
ext. citations

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avg, IF

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L-index

#	Paper	IF	Citations
219	New views on strand asymmetry in insect mitochondrial genomes. <i>PLoS ONE</i> , 2010 , 5, e12708	3.7	166
218	Comparative mitogenomics of Braconidae (Insecta: Hymenoptera) and the phylogenetic utility of mitochondrial genomes with special reference to Holometabolous insects. <i>BMC Genomics</i> , 2010 , 11, 3714-5	4.5	120
217	The complete mitochondrial genome of <i>Diadegma semiclausum</i> (hymenoptera: ichneumonidae) indicates extensive independent evolutionary events. <i>Genome</i> , 2009 , 52, 308-19	2.4	83
216	Differences in induced volatile emissions among rice varieties result in differential attraction and parasitism of <i>Nilaparvata lugens</i> eggs by the parasitoid <i>Anagrus nilaparvatae</i> in the field. <i>Journal of Chemical Ecology</i> , 2006 , 32, 2375-87	2.7	81
215	The complete mitochondrial genome of <i>Evania appendigaster</i> (Hymenoptera: Evaniidae) has low A+T content and a long intergenic spacer between <i>atp8</i> and <i>atp6</i> . <i>Molecular Biology Reports</i> , 2010 , 37, 1931-42	2.8	74
214	The autophagy pathway participates in resistance to tomato yellow leaf curl virus infection in whiteflies. <i>Autophagy</i> , 2016 , 12, 1560-74	10.2	73
213	Antimicrobial peptide evolution in the Asiatic honey bee <i>Apis cerana</i> . <i>PLoS ONE</i> , 2009 , 4, e4239	3.7	68
212	Tea: Biological control of insect and mite pests in China. <i>Biological Control</i> , 2014 , 68, 73-91	3.8	67
211	Deep sequencing of <i>Cotesia vestalis</i> bracovirus reveals the complexity of a polydnavirus genome. <i>Virology</i> , 2011 , 414, 42-50	3.6	63
210	Genetic structure and demographic history reveal migration of the diamondback moth <i>Plutella xylostella</i> (Lepidoptera: Plutellidae) from the southern to northern regions of China. <i>PLoS ONE</i> , 2013 , 8, e59654	3.7	63
209	Population genetic structure of <i>Chilo suppressalis</i> (Walker) (Lepidoptera: Crambidae): strong subdivision in China inferred from microsatellite markers and mtDNA gene sequences. <i>Molecular Ecology</i> , 2008 , 17, 2880-97	5.7	60
208	Systematics, Phylogeny, and Evolution of Braconid Wasps: 30 Years of Progress. <i>Annual Review of Entomology</i> , 2019 , 64, 335-358	21.8	54
207	Mitochondrial phylogenomics of the Hymenoptera. <i>Molecular Phylogenetics and Evolution</i> , 2019 , 131, 8-18	4.1	51
206	Two mitochondrial genomes from the families Bethyridae and Mutillidae: independent rearrangement of protein-coding genes and higher-level phylogeny of the Hymenoptera. <i>Molecular Phylogenetics and Evolution</i> , 2014 , 77, 1-10	4.1	46
205	Aphid dispersal flight disseminates fungal pathogens and parasitoids as natural control agents of aphids. <i>Ecological Entomology</i> , 2007 , 32, 97-104	2.1	46
204	Population genetic structure and approximate Bayesian computation analyses reveal the southern origin and northward dispersal of the oriental fruit moth <i>Grapholita molesta</i> (Lepidoptera: Tortricidae) in its native range. <i>Molecular Ecology</i> , 2015 , 24, 4094-111	5.7	42
203	Multiple Lines of Evidence from Mitochondrial Genomes Resolve Phylogenetic Relationships of Parasitic Wasps in Braconidae. <i>Genome Biology and Evolution</i> , 2016 , 8, 2651-62	3.9	42

202	Comparative and phylogenetic analysis of the mitochondrial genomes in basal hymenopterans. <i>Scientific Reports</i> , 2016 , 6, 20972	4.9	38
201	Effects of venom/calyx fluid from the endoparasitic wasp <i>Cotesia plutellae</i> on the hemocytes of its host <i>Plutella xylostella</i> in vitro. <i>Journal of Insect Physiology</i> , 2007 , 53, 22-9	2.4	38
200	Parasitic insect-derived miRNAs modulate host development. <i>Nature Communications</i> , 2018 , 9, 2205	17.4	37
199	Isolation and characterization of an immunosuppressive protein from venom of the pupa-specific endoparasitoid <i>Pteromalus puparum</i> . <i>Journal of Invertebrate Pathology</i> , 2008 , 99, 186-91	2.6	37
198	Venom of parasitoid, <i>Pteromalus puparum</i> , suppresses host, <i>Pieris rapae</i> , immune promotion by decreasing host C-type lectin gene expression. <i>PLoS ONE</i> , 2011 , 6, e26888	3.7	36
197	Arthropod abundance and diversity in Bt and non-Bt rice fields. <i>Environmental Entomology</i> , 2007 , 36, 646-54	2.1	34
196	Parasitoid wasps as effective biological control agents. <i>Journal of Integrative Agriculture</i> , 2019 , 18, 705-715	3.5	31
195	Utility of multi-gene loci for forensic species diagnosis of blowflies. <i>Journal of Insect Science</i> , 2011 , 11, 59	2	31
194	Gene arrangement and sequence of mitochondrial genomes yield insights into the phylogeny and evolution of bees and sphecid wasps (Hymenoptera: Apoidea). <i>Molecular Phylogenetics and Evolution</i> , 2018 , 124, 1-9	4.1	30
193	The occurrence of leafminers and their parasitoids on vegetables and weeds in Hangzhou area, Southeast China. <i>BioControl</i> , 2003 , 48, 515-527	2.3	30
192	The complete mitochondrial genome of <i>Taeniogonalos taihorina</i> (Bischoff) (Hymenoptera: Trigonalidae) reveals a novel gene rearrangement pattern in the Hymenoptera. <i>Gene</i> , 2014 , 543, 76-84	3.8	28
191	Thrips (Insecta: Thysanoptera) of China. <i>Check List</i> , 2011 , 7, 720	1	25
190	The genomic features of parasitism, Polyembryony and immune evasion in the endoparasitic wasp <i>Macrocentrus cingulum</i> . <i>BMC Genomics</i> , 2018 , 19, 420	4.5	23
189	Flower-visiting insects and their potential impact on transgene flow in rice. <i>Journal of Applied Ecology</i> , 2014 , 51, 1357-1365	5.8	23
188	Four heat shock protein genes of the endoparasitoid wasp, <i>Cotesia vestalis</i> , and their transcriptional profiles in relation to developmental stages and temperature. <i>PLoS ONE</i> , 2013 , 8, e59723	3.7	23
187	Complete mitochondrial genome of <i>Empoasca vitis</i> (Hemiptera: Cicadellidae). <i>Mitochondrial DNA</i> , 2016 , 27, 1052-3		22
186	Molecular phylogeny of the Aphidiinae (Hymenoptera: Braconidae) based on DNA sequences of 16S rRNA, 18S rDNA and ATPase 6 genes. <i>European Journal of Entomology</i> , 2005 , 102, 133-138		22
185	Parasitic castration of <i>Plutella xylostella</i> larvae induced by polydnaviruses and venom of <i>Cotesia vestalis</i> and <i>Diadegma semiclausum</i> . <i>Archives of Insect Biochemistry and Physiology</i> , 2009 , 70, 30-43	2.3	21

184	Parasitoid polydnviruses and immune interaction with secondary hosts. <i>Developmental and Comparative Immunology</i> , 2018 , 83, 124-129	3.2	20
183	Pteromalus puparum venom impairs host cellular immune responses by decreasing expression of its scavenger receptor gene. <i>Insect Biochemistry and Molecular Biology</i> , 2011 , 41, 852-62	4.5	20
182	A peptidoglycan recognition protein acts in whitefly (<i>Bemisia tabaci</i>) immunity and involves in Begomovirus acquisition. <i>Scientific Reports</i> , 2016 , 6, 37806	4.9	20
181	The endoparasitoid, <i>Cotesia vestalis</i> , regulates host physiology by reprogramming the neuropeptide transcriptional network. <i>Scientific Reports</i> , 2015 , 5, 8173	4.9	19
180	Multi-generation effects of Bt rice on <i>Anagrus nilaparvatae</i> , a parasitoid of the nontarget pest <i>Nilaparvata lugens</i> . <i>Environmental Entomology</i> , 2010 , 39, 2039-44	2.1	18
179	Immature Development of <i>Cotesia vestalis</i> (Hymenoptera: Braconidae), an Endoparasitoid of <i>Plutella xylostella</i> (Lepidoptera: Plutellidae). <i>Annals of the Entomological Society of America</i> , 2008 , 101, 189-196	2	18
178	The complete mitochondrial genome of <i>Tambocerus</i> sp. (Hemiptera: Cicadellidae). <i>Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis</i> , 2017 , 28, 133-134	1.3	16
177	Transgenic plants expressing the AaIT/GNA fusion protein show increased resistance and toxicity to both chewing and sucking pests. <i>Insect Science</i> , 2016 , 23, 265-76	3.6	16
176	Interspecific competition between two endoparasitoids <i>Cotesia vestalis</i> (Hymenoptera: Braconidae) and <i>Oomyzus sokolowskii</i> (Hymenoptera: Eulophidae). <i>Archives of Insect Biochemistry and Physiology</i> , 2011 , 76, 156-67	2.3	16
175	Characterization of a novel gene encoding ankyrin repeat domain from <i>Cotesia vestalis</i> polydnvirus (CvBV). <i>Virology</i> , 2008 , 375, 374-82	3.6	16
174	Characterization of the mitochondrial genome of the diamondback moth <i>Plutella xylostella</i> (Lepidoptera: Plutellidae) and phylogenetic analysis of advanced moths and butterflies. <i>DNA and Cell Biology</i> , 2013 , 32, 173-87	3.6	15
173	Parasitism-induced effects on host growth and metabolic efficiency in <i>Plutella xylostella</i> larvae parasitized by <i>Cotesia vestalis</i> or <i>Diadegma semiclausum</i> . <i>Insect Science</i> , 2008 , 15, 237-243	3.6	15
172	<i>Cotesia vestalis</i> teratocytes express a diversity of genes and exhibit novel immune functions in parasitism. <i>Scientific Reports</i> , 2016 , 6, 26967	4.9	14
171	Ras1CA-upregulated BCPI inhibits cathepsin activity to prevent tissue destruction of the Bombyx posterior silk gland. <i>Journal of Proteome Research</i> , 2013 , 12, 1924-34	5.6	14
170	Complete mitochondrial genome of the tea looper caterpillar, <i>Ectropis obliqua</i> (Lepidoptera: Geometridae) with a phylogenetic analysis of Geometridae. <i>International Journal of Biological Macromolecules</i> , 2018 , 114, 491-496	7.9	13
169	The subgenus <i>Choeras</i> Mason, 1981 of genus <i>Apanteles</i> Foerster, 1862 (Hymenoptera, Braconidae, Microgastrinae) from China, with descriptions of eighteen new species. <i>Zootaxa</i> , 2014 , 3754, 501-54	0.5	13
168	Ultrastructural and functional characterization of circulating hemocytes from <i>Plutella xylostella</i> larva: cell types and their role in phagocytosis. <i>Tissue and Cell</i> , 2010 , 42, 360-4	2.7	13
167	Changes in hemocytes of <i>Plutella xylostella</i> after parasitism by <i>Diadegma semiclausum</i> . <i>Archives of Insect Biochemistry and Physiology</i> , 2009 , 70, 177-87	2.3	13

166	The first two mitochondrial genomes of the family Aphelinidae with novel gene orders and phylogenetic implications. <i>International Journal of Biological Macromolecules</i> , 2018 , 118, 386-396	7.9	13
165	Four serine protease cDNAs from the midgut of <i>Plutella xylostella</i> and their proteinase activity are influenced by the endoparasitoid, <i>Cotesia vestalis</i> . <i>Archives of Insect Biochemistry and Physiology</i> , 2013 , 83, 101-14	2.3	12
164	RNA interference of an antimicrobial peptide, <i>Btdef</i> , reduces Tomato yellow leaf curl China virus accumulation in the whitefly <i>Bemisia tabaci</i> . <i>Pest Management Science</i> , 2017 , 73, 1421-1427	4.6	12
163	The genomes of two parasitic wasps that parasitize the diamondback moth. <i>BMC Genomics</i> , 2019 , 20, 893	4.5	12
162	Rearrangement of the <i>nad1</i> gene in <i>Pristaulacus compressus</i> (Spinola) (Hymenoptera: Evanioidea: Aulacidae) mitochondrial genome. <i>Mitochondrial DNA</i> , 2015 , 26, 629-30		11
161	An illustrated key to the genera of Thripinae (Thysanoptera, Thripidae) from Iran. <i>ZooKeys</i> , 2013 , 27-52	1.2	11
160	The genus <i>Diolcogaster</i> Ashmead, 1900 (Hymenoptera, Braconidae, Microgastrinae) from China. <i>ZooKeys</i> , 2011 , 49-87	1.2	11
159	Venom apparatus of the endoparasitoid wasp <i>Opius caricivora</i> Fischer (Hymenoptera: Braconidae): morphology and ultrastructure. <i>Microscopy Research and Technique</i> , 2006 , 69, 820-5	2.8	11
158	The first two mitochondrial genomes of wood wasps (Hymenoptera: Symphyta): Novel gene rearrangements and higher-level phylogeny of the basal hymenopterans. <i>International Journal of Biological Macromolecules</i> , 2019 , 123, 1189-1196	7.9	11
157	Biocontrol characteristics of the fruit fly pupal parasitoid <i>Trichopria drosophilae</i> (Hymenoptera: Diapriidae) emerging from different hosts. <i>Scientific Reports</i> , 2018 , 8, 13323	4.9	11
156	Laccase 1 gene from <i>Plutella xylostella</i> (PxLac1) and its functions in humoral immune response. <i>Journal of Insect Physiology</i> , 2018 , 107, 197-203	2.4	10
155	Comparative transcriptome analysis of venom glands from <i>Cotesia vestalis</i> and <i>Diadromus collaris</i> , two endoparasitoids of the host <i>Plutella xylostella</i> . <i>Scientific Reports</i> , 2017 , 7, 1298	4.9	10
154	First description of the male of the wheat thrips, <i>Anaphothrips obscurus</i> (Thysanoptera: Thripidae). <i>Zootaxa</i> , 2010 , 2540, 65	0.5	10
153	The mitochondrial genome of <i>Tenthredo tienmushana</i> (Takeuchi) and a related phylogenetic analysis of the sawflies (Insecta: Hymenoptera). <i>Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis</i> , 2016 , 27, 2860-1	1.3	9
152	The mitochondrial genome of <i>Polistes jokahamae</i> and a phylogenetic analysis of the Vespoidea (Insecta: Hymenoptera). <i>Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis</i> , 2016 , 27, 2783-4	1.3	9
151	Identification and characterization of defensin genes from the endoparasitoid wasp <i>Cotesia vestalis</i> (Hymenoptera: Braconidae). <i>Journal of Insect Physiology</i> , 2013 , 59, 1095-103	2.4	9
150	Molecular identification of two prophenoloxidase-activating proteases from the hemocytes of <i>Plutella xylostella</i> (Lepidoptera: Plutellidae) and their transcript abundance changes in response to microbial challenges. <i>Journal of Insect Science</i> , 2014 , 14, 179	2	9
149	The genus <i>Doryctes</i> Haliday, 1836 (Hymenoptera: Braconidae, Doryctinae) in China. <i>Zootaxa</i> , 2012 , 3226, 46	0.5	9

148	A new record and description of a new species of the genus Thrips, with an updated key to species from Iran. <i>Journal of Insect Science</i> , 2012 , 12, 90		9
147	Oogenesis of <i>Diadegma semiclausum</i> (Hymenoptera: Ichneumonidae) and its associated polydnavirus. <i>Microscopy Research and Technique</i> , 2008 , 71, 676-83	2.8	9
146	An investigation of irreproducibility in maximum likelihood phylogenetic inference. <i>Nature Communications</i> , 2020 , 11, 6096	17.4	9
145	The grandiculus- and metacarpalis-group of the genus <i>Apanteles</i> Foerster, 1862 (Hymenoptera, Braconidae, Microgastrinae) from China, with descriptions of eight new species. <i>Zootaxa</i> , 2014 , 3765, 435-57	0.5	8
144	Effects of Transgenic Bt+CpTI Cotton on Field Abundance of Non-Target Pests and Predators in Xinjiang, China. <i>Journal of Integrative Agriculture</i> , 2012 , 11, 1493-1499	3.2	8
143	The genera <i>Deuterixys</i> Mason, 1981 and <i>Wilkinsonellus</i> Mason, 1981 (Hymenoptera, Braconidae, Microgastrinae) from China, with description of two new species. <i>ZooKeys</i> , 2011 , 27-40	1.2	8
142	The genus <i>Saphonecrus</i> Dalla Torre et Kieffer, 1910 (Hymenoptera: Cynipidae) in China, with descriptions of two new species. <i>Biologia (Poland)</i> , 2010 , 65, 1034-1039	1.5	8
141	Characterization of two genes of <i>Cotesia vestalis</i> polydnavirus and their expression patterns in the host <i>Plutella xylostella</i> . <i>Journal of General Virology</i> , 2007 , 88, 3317-3322	4.9	8
140	Four new species of Gasteruption Latreille from NW China, with an illustrated key to the species from Palaeartic China (Hymenoptera, Gasteruptionidae). <i>ZooKeys</i> , 2016 , 51-112	1.2	8
139	Mitochondrial DNA and their nuclear copies in the parasitic wasp <i>Pteromalus puparum</i> : A comparative analysis in Chalcidoidea. <i>International Journal of Biological Macromolecules</i> , 2019 , 121, 572-579	7.9	8
138	A trypsin inhibitor-like protein secreted by <i>Cotesia vestalis</i> teratocytes inhibits hemolymph prophenoloxidase activation of <i>Plutella xylostella</i> . <i>Journal of Insect Physiology</i> , 2019 , 116, 41-48	2.4	7
137	The lacteus-, laspeyresiella- and mycetophilus-groups of <i>Apanteles</i> Foerster, 1862 (Hymenoptera, Braconidae, Microgastrinae) in China, with descriptions of eight new species. <i>Zootaxa</i> , 2015 , 3949, 370-92	9.5	7
136	<i>Spathius</i> Nees, 1818 (Hymenoptera: Braconidae, Doryctinae) from China with a key to species. <i>Zootaxa</i> , 2015 , 3960, 1-132	0.5	7
135	A preliminary survey of carrion breeding insects associated with the Eid ul Azha festival in remote Pakistan. <i>Forensic Science International</i> , 2011 , 209, 186-94	2.6	7
134	Effects of starvation on the vitellogenesis, ovarian development and fecundity in the ectoparasitoid, <i>Nasonia vitripennis</i> (Hymenoptera: Pteromalidae). <i>Insect Science</i> , 2008 , 15, 429-440	3.6	7
133	Immature Morphology and Development of <i>Opius caricivora</i> (Hymenoptera: Braconidae), an Endoparasitoid of the Leafminer <i>Liriomyza sativae</i> (Diptera: Agromyzidae). <i>Annals of the Entomological Society of America</i> , 2007 , 100, 425-432	2	7
132	An illustrated key to the genera and subgenera of the Alysini (Hymenoptera, Braconidae, Alysiniinae), with three genera new for China. <i>ZooKeys</i> , 2017 , 37-79	1.2	7
131	Different genetic structures revealed resident populations of a specialist parasitoid wasp in contrast to its migratory host. <i>Ecology and Evolution</i> , 2017 , 7, 5400-5409	2.8	6

130	Molecular Characterization of Two Fatty Acyl-CoA Reductase Genes From <i>Phenacoccus solenopsis</i> (Hemiptera: Pseudococcidae). <i>Journal of Insect Science</i> , 2016 , 16,	2	6
129	A new species of <i>Andricus</i> oak gallwasp from China (Hymenoptera: Cynipidae: Cynipini). <i>Biologia (Poland)</i> , 2013 , 68, 974-978	1.5	6
128	The mitochondrial genome of <i>Diadromus collaris</i> (Hymenoptera: Ichneumonidae). <i>Mitochondrial DNA</i> , 2015 , 26, 303-4		6
127	<i>Heterospilus</i> Haliday, 1836 (Hymenoptera: Braconidae, Doryctinae) from China with a key to species. <i>Zootaxa</i> , 2013 , 3683, 201-46	0.5	6
126	Characterization of midgut trypsinogen-like cDNA and enzymatic activity in <i>Plutella xylostella</i> parasitized by <i>Cotesia vestalis</i> or <i>Diadegma semiclausum</i> . <i>Archives of Insect Biochemistry and Physiology</i> , 2009 , 70, 3-17	2.3	6
125	Revision of the genus <i>Eodendrus</i> Belokobylskij (Hymenoptera: Braconidae, Doryctinae). <i>Journal of Natural History</i> , 2005 , 39, 2715-2743	0.5	6
124	Venom gland of the ichneumonid <i>Diadromus collaris</i> : morphology, ultrastructure and age-related changes. <i>Insect Science</i> , 2006 , 13, 137-143	3.6	6
123	<i>Cornutorogas</i> , a new genus with four new species of the subfamily Rogadinae (Hymenoptera: Braconidae) from the Oriental region. <i>Journal of Natural History</i> , 2004 , 38, 2211-2223	0.5	6
122	The genus <i>Atanycolus</i> Foerster (Hymenoptera, Braconidae, Braconinae) in China, with description of one new species. <i>ZooKeys</i> , 27, 31-41	1.2	6
121	Two novel venom proteins underlie divergent parasitic strategies between a generalist and a specialist parasite. <i>Nature Communications</i> , 2021 , 12, 234	17.4	6
120	Two types of lysozymes from the whitefly <i>Bemisia tabaci</i> : Molecular characterization and functional diversification. <i>Developmental and Comparative Immunology</i> , 2018 , 81, 252-261	3.2	6
119	The laevigata-group of the genus <i>Dolichogenidea</i> Mason, 1981 from China, with descriptions of 26 new species. <i>Zootaxa</i> , 2018 , 4436, 1-74	0.5	6
118	Bracoviruses recruit host integrases for their integration into caterpillar genome. <i>PLoS Genetics</i> , 2021 , 17, e1009751	6	6
117	A Cuboid Spider Silk: Structure-Function Relationship and Polypeptide Signature. <i>Macromolecular Rapid Communications</i> , 2020 , 41, e1900583	4.8	5
116	The genus <i>Pholetesor</i> Mason, 1981 (Hymenoptera, Braconidae, Microgastrinae) from China, with descriptions of eleven new species. <i>Zootaxa</i> , 2016 , 4150, 351-87	0.5	5
115	Review of the tribe Helconini Foerster s.s. from China, with the description of 18 new species. <i>Zootaxa</i> , 2017 , 4291, 401	0.5	5
114	Review of the genus <i>Taiwanomyrme</i> Tsuneki, 1993 (Hymenoptera, Mutillidae, Mutillinae), with description of two new species from China. <i>Zootaxa</i> , 2015 , 4020, 588-600	0.5	5
113	The Chinese species of the genus <i>Ontsira</i> Cameron (Hymenoptera, Braconidae, Doryctinae). <i>ZooKeys</i> , 2013 , 73-96	1.2	5

112	A new species of Sericothripinae from China (Thysanoptera: Thripidae), with two new synonyms and one new record. <i>Zootaxa</i> , 2011 , 3009, 55	0.5	5
111	Two new species of the genus Mycterothrips from Western Iran (Thysanoptera: Thripidae). <i>Zootaxa</i> , 2011 , 3130, 57	0.5	5
110	The Genus Minanga Cameron (Hymenoptera: Braconidae) in China, With Description of a New Subgenus and Species. <i>Annals of the Entomological Society of America</i> , 2010 , 103, 360-365	2	5
109	External Morphology and Development of Immature Stages of Diadegma semiclausum (Hymenoptera: Ichneumonidae), an Important Endoparasitoid of Plutella xylostella (Lepidoptera: Plutellidae). <i>Annals of the Entomological Society of America</i> , 2009 , 102, 532-538	2	5
108	Ultrastructure of the male reproductive system of Cotesia vestalis (Hymenoptera: Braconidae) with preliminary characterization of the secretions. <i>Microscopy Research and Technique</i> , 2007 , 70, 563-71	2.8	5
107	Characterization of an IkappaB-like gene in Cotesia vestalis polydnavirus. <i>Archives of Insect Biochemistry and Physiology</i> , 2008 , 68, 71-8	2.3	5
106	Hecabolomorpha n. gen., a new Asian genus from the tribe Hecabolini (Hymenoptera: Braconidae: Doryctinae). <i>Annales De La Societe Entomologique De France</i> , 2006 , 42, 107-111	0.5	5
105	The genus Testudobracon in eastern China and the description of three new species (Hymenoptera: Braconidae: Braconinae). <i>European Journal of Entomology</i> , 2003 , 100, 411-416		5
104	Pathogenic fungi and parasitoids of aphids present in air captures of migratory alates in the low-latitude plateau of Yunnan, China. <i>Environmental Entomology</i> , 2008 , 37, 1264-71	2.1	5
103	A new species of Schlettererius Ashmead from China, with a key to the species (Hymenoptera, Stephanidae). <i>Journal of Hymenoptera Research</i> , 45 , 75-86	0	5
102	Expression and functional characterization of odorant-binding protein genes in the endoparasitic wasp Cotesia vestalis. <i>Insect Science</i> , 2021 , 28, 1354-1368	3.6	5
101	Effects of Transgenic Bt Rice on Nontarget Rhopalosiphum maidis (Homoptera: Aphididae). <i>Environmental Entomology</i> , 2016 , 45, 1090-6	2.1	5
100	Taxonomy of parasitoid wasps in China: An overview. <i>Biological Control</i> , 2014 , 68, 57-72	3.8	4
99	Halycaea Cameron, 1903 (Hymenoptera: Braconidae, Doryctinae) from China with a key to world species. <i>Zootaxa</i> , 2012 , 3218, 18	0.5	4
98	The genus Arhaconotus Belokobylskij (Hymenoptera, Braconidae, Doryctinae) from China, with description of a new species. <i>ZooKeys</i> , 2010 , 63-8	1.2	4
97	Two genera of Braconinae (Hymenoptera, Braconidae) in China, with descriptions of four new species. <i>ZooKeys</i> , 2010 , 47-62	1.2	4
96	Characterization of a protein tyrosine phosphatase gene CvBV202 from Cotesia vestalis polydnavirus (CvBV). <i>Virus Genes</i> , 2008 , 36, 595-601	2.3	4
95	A Review of Bracon (Rostrobracon) (Hymenoptera: Braconidae: Braconinae) from China, With Description of One New Species. <i>Oriental Insects</i> , 2004 , 38, 341-345	0.3	4

94	Two new species of Aivalykus (Hymenoptera: Braconidae: Doryctinae) from China and Indonesia, with a key to species. <i>European Journal of Entomology</i> , 2002 , 99, 73-78		4
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90	Bioinspired Conical Micropattern Modulates Cell Behaviors.. <i>ACS Applied Bio Materials</i> , 2018 , 1, 1416-1423	1	4
89	Complete mitochondrial genome of Neochauliodes bowringi (MacLachlan) (Megaloptera: Corydalidae). <i>Mitochondrial DNA</i> , 2015 , 26, 112-3		3
88	An illustrated key to the species of subgenus Gyrostoma Kirby, 1828 (Hymenoptera, Vespidae, Polistinae) from China, with discovery of Polistes (Gyrostoma) tenuispunctia Kim, 2001. <i>Zootaxa</i> , 2014 , 3785, 377-99	0.5	3
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85	Biological characteristics of Cionus latefasciatus (Coleoptera: Curculionidae) and effects of temperature on its growth and development. <i>Journal of Economic Entomology</i> , 2009 , 102, 1039-43	2.2	3
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79	The genus Coeloides Wesmael of subfamily Braconinae (Hymenoptera: Braconidae) in China. <i>Zootaxa</i> , 2006 , 1239, 1	0.5	3
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66	The genus <i>Ipodoryctes</i> Granger, 1949 (Hymenoptera: Braconidae, Doryctinae) from China. <i>Zootaxa</i> , 2011 , 2784, 1	0.5	2
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28	The genus <i>Eurobracon</i> Ashmead (Hymenoptera, Braconidae, Braconinae) in China, with description of three new species. <i>Zootaxa</i> , 2016 , 4132, 383-92	0.5	0
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26	A new <i>Odontothrips</i> species (Thysanoptera: Thripidae) from Iran. <i>Zootaxa</i> , 2013 , 3736, 598-600	0.5	0
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