

Jianhua Huang

List of Publications by Year in descending order

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91
papers

1,519
citations

394421

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395702

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95
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times ranked

1338
citing authors

#	ARTICLE	IF	CITATIONS
1	Virus and endogenous viral element-derived small non-coding RNAs and their roles in insect-virus interaction. <i>Current Opinion in Insect Science</i> , 2022, 49, 85-92.	4.4	2
2	Hidden RNA pairings counteract the “first-come, first-served” splicing principle to drive stochastic choice in <i>Dscam1</i> splice variants. <i>Science Advances</i> , 2022, 8, eabm1763.	10.3	6
3	The mitochondrial genome of <i>Chelonus formosanus</i> (Hymenoptera: Braconidae) with novel gene orders and phylogenetic implications. <i>Archives of Insect Biochemistry and Physiology</i> , 2022, , e21870.	1.5	4
4	Identification and Functional Analysis of Glutathione S-Transferases from <i>Sitophilus zeamais</i> in Olfactory Organ. <i>Insects</i> , 2022, 13, 259.	2.2	11
5	Characterization of Molting Process during the Different Developmental Stages of the Diamondback Moth <i>Plutella xylostella</i> . <i>Insects</i> , 2022, 13, 289.	2.2	0
6	A new subgenus of <i>Chelonus</i> Panzer, 1806 (Braconidae: Cheloninae) from China. <i>Zootaxa</i> , 2022, 5115, 288-294.	0.5	6
7	The Dual Functions of a Bracovirus C-Type Lectin in Caterpillar Immune Response Manipulation. <i>Frontiers in Immunology</i> , 2022, 13, .	4.8	3
8	Expression and functional characterization of odorant-binding protein genes in the endoparasitic wasp <i>Cotesia vestalis</i> . <i>Insect Science</i> , 2021, 28, 1354-1368.	3.0	16
9	<i>CLP</i> gene family, a new gene family of <i>Cotesia vestalis</i> bracovirus inhibits melanization of <i>Plutella xylostella</i> hemolymph. <i>Insect Science</i> , 2021, 28, 1567-1581.	3.0	6
10	Migration trajectories of the diamondback moth <i>Plutella xylostella</i> in China inferred from population genomic variation. <i>Pest Management Science</i> , 2021, 77, 1683-1693.	3.4	18
11	Two novel venom proteins underlie divergent parasitic strategies between a generalist and a specialist parasite. <i>Nature Communications</i> , 2021, 12, 234.	12.8	25
12	Symbiotic bracovirus of a parasite manipulates host lipid metabolism via tachykinin signaling. <i>PLoS Pathogens</i> , 2021, 17, e1009365.	4.7	17
13	The mitochondrial genome of <i>Telenomus remus</i> (Hymenoptera: Platygasteridae). <i>Mitochondrial DNA Part B: Resources</i> , 2021, 6, 844-845.	0.4	1
14	Illustrated keys to Scoliidae (Insecta, Hymenoptera, Scoliioidea) from China. <i>ZooKeys</i> , 2021, 1025, 139-175.	1.1	6
15	Large-Scale Annotation and Evolution Analysis of MiRNA in Insects. <i>Genome Biology and Evolution</i> , 2021, 13, .	2.5	15
16	Taxonomy of <i>Bethylus</i> (Hymenoptera, Bethylinidae) from China with description of nine new species. <i>Zootaxa</i> , 2021, 4974, 361-382.	0.5	0
17	A checklist of Scoliidae (Insecta: Hymenoptera) from China. <i>Zootaxa</i> , 2021, 4966, 101126.	0.5	5
18	The genus <i>Casinaria</i> Holmgren, 1859 (Hymenoptera: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 67 Td (Ichneumonidae). <i>Zootaxa</i> , 2021, 4966, 101126.	0.5	0

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19	Comparative Transcriptome Analysis Reveals Sex-Based Differences during the Development of the Adult Parasitic Wasp <i>Cotesia vestalis</i> (Hymenoptera: Braconidae). <i>Genes</i> , 2021, 12, 896.	2.4	4
20	Comparative mitogenomics and phylogenetics of the stinging wasps (Hymenoptera: Aculeata). <i>Molecular Phylogenetics and Evolution</i> , 2021, 159, 107119.	2.7	13
21	Intron-targeted mutagenesis reveals roles for Dscam1 RNA pairing architecture-driven splicing bias in neuronal wiring. <i>Cell Reports</i> , 2021, 36, 109373.	6.4	11
22	Bracoviruses recruit host integrases for their integration into caterpillar's genome. <i>PLoS Genetics</i> , 2021, 17, e1009751.	3.5	15
23	Comparative transcriptome analysis reveals a potential mechanism for host nutritional manipulation after parasitization by <i>Leptopilina boulardi</i> . <i>Comparative Biochemistry and Physiology Part D: Genomics and Proteomics</i> , 2021, 39, 100862.	1.0	2
24	Neofunctionalization of an ancient domain allows parasites to avoid intraspecific competition by manipulating host behaviour. <i>Nature Communications</i> , 2021, 12, 5489.	12.8	15
25	Juvenile hormone signaling promotes ovulation and maintains egg shape by inducing expression of extracellular matrix genes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	7.1	37
26	The complete mitochondrial genome of <i>Leptopilina syphax</i> (Hymenoptera: Figitidae). <i>Mitochondrial DNA Part B: Resources</i> , 2021, 6, 17-18.	0.4	0
27	Five new species of the genus <i>Sinophorus</i> Förster (Hymenoptera, Ichneumonidae, Campopleginae) from China. <i>Zootaxa</i> , 2021, 5061, 115-133.	0.5	0
28	The genus <i>Campoplex</i> Gravenhorst, 1829 (Hymenoptera, Ichneumonidae, Campopleginae) from China. <i>Zootaxa</i> , 2021, 5066, 1-121.	0.5	0
29	An investigation of irreproducibility in maximum likelihood phylogenetic inference. <i>Nature Communications</i> , 2020, 11, 6096.	12.8	32
30	The complete mitochondrial genome of <i>Asobara japonica</i> (Hymenoptera: Braconidae). <i>Mitochondrial DNA Part B: Resources</i> , 2020, 5, 1279-1281.	0.4	5
31	The complete mitochondrial genome of <i>Trichopria drosophilae</i> (Hymenoptera: Diapriidae). <i>Mitochondrial DNA Part B: Resources</i> , 2020, 5, 2391-2393.	0.4	3
32	The first mitochondrial genome of the living-fossil sawfly <i>Macroxyela ferruginea</i> (Hymenoptera: Tj ETQq0 0 0 rgBT /Oyerlock 10 Tf 50 22	0.4	1
33	Genome-Wide Profiling of <i>Diadegma semiclausum</i> Ichnovirus Integration in Parasitized <i>Plutella xylostella</i> Hemocytes Identifies Host Integration Motifs and Insertion Sites. <i>Frontiers in Microbiology</i> , 2020, 11, 608346.	3.5	7
34	The mitochondrial genome of <i>Aenasius arizonensis</i> (Hymenoptera: Encyrtidae) with novel gene order. <i>Mitochondrial DNA Part B: Resources</i> , 2019, 4, 2023-2024.	0.4	4
35	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.	2.0	17
36	The genomes of two parasitic wasps that parasitize the diamondback moth. <i>BMC Genomics</i> , 2019, 20, 893.	2.8	17

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37	The developmental transcriptome of <i>Trichopria drosophilae</i> (Hymenoptera: Diapriidae) and insights into cuticular protein genes. <i>Comparative Biochemistry and Physiology Part D: Genomics and Proteomics</i> , 2019, 29, 245-254.	1.0	5
38	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.	7.5	17
39	Mitochondrial phylogenomics of the Hymenoptera. <i>Molecular Phylogenetics and Evolution</i> , 2019, 131, 8-18.	2.7	104
40	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.	2.7	49
41	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.	2.0	15
42	Parasitoid polydnnaviruses and immune interaction with secondary hosts. <i>Developmental and Comparative Immunology</i> , 2018, 83, 124-129.	2.3	46
43	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.	2.3	7
44	Biocontrol characteristics of the fruit fly pupal parasitoid <i>Trichopria drosophilae</i> (Hymenoptera: Diapriidae). <i>Journal of Applied Entomology</i> , 2018, 10, 22-27.	3.3	22
45	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.	7.5	23
46	Parasitic insect-derived miRNAs modulate host development. <i>Nature Communications</i> , 2018, 9, 2205.	12.8	77
47	The genera <i>Areopraon</i> Mackauer, 1959 and <i>Pseudopraon</i> Starbäck, 1975 (Hymenoptera, Braconidae). <i>Journal of Natural History</i> , 2017, 51, 2745-2758.	1.1	5
48	Yorkie and Hedgehog independently restrict BMP production in Escort cells to permit germline differentiation in the <i>Drosophila</i> ovary. <i>Development (Cambridge)</i> , 2017, 144, 2584-2594.	2.5	32
49	The genus <i>Bassus</i> Fabricius, 1804 (Hymenoptera: Braconidae: Agathidinae) in China, with description of three new species. <i>Journal of Natural History</i> , 2017, 51, 2745-2758.	0.5	0
50	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.	3.3	17
51	RNA interference of an antimicrobial peptide, Btdef, reduces <i>Tomato yellow leaf curl China virus</i> accumulation in the whitefly <i>Bemisia tabaci</i> . <i>Pest Management Science</i> , 2017, 73, 1421-1427.	3.4	16
52	An illustrated key to the genera and subgenera of the Alysini (Hymenoptera, Braconidae, Alysiniinae), with three genera new for China. <i>ZooKeys</i> , 2017, 722, 37-79.	1.1	20
53	Effects of Transgenic Bt Rice on Nontarget <i>Rhopalosiphum maidis</i> (Homoptera: Aphididae). <i>Environmental Entomology</i> , 2016, 45, 1090-1096.	1.4	6
54	General morphology and ultrastructure of the female reproductive apparatus of <i>Trichomalopsis shirakii</i> crawford (Hymenoptera, Pteromalidae). <i>Microscopy Research and Technique</i> , 2016, 79, 625-636.	2.2	4

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55	A peptidoglycan recognition protein acts in whitefly (<i>Bemisia tabaci</i>) immunity and involves in Begomovirus acquisition. <i>Scientific Reports</i> , 2016, 6, 37806.	3.3	31
56	Multiple Lines of Evidence from Mitochondrial Genomes Resolve Phylogenetic Relationships of Parasitic Wasps in Braconidae. <i>Genome Biology and Evolution</i> , 2016, 8, 2651-2662.	2.5	57
57	The genus <i>Euurobracon</i> Ashmead (Hymenoptera, Braconidae, Braconinae) in China, with description of three new species. <i>Zootaxa</i> , 2016, 4132, 383.	0.5	2
58	The genus <i>Pholetesor</i> Mason, 1981 (Hymenoptera, Braconidae, Microgastrinae) from China, with descriptions of eleven new species. <i>Zootaxa</i> , 2016, 4150, 351.	0.5	6
59	Comparative and phylogenetic analysis of the mitochondrial genomes in basal hymenopterans. <i>Scientific Reports</i> , 2016, 6, 20972.	3.3	56
60	<i>Cotesia vestalis</i> teratocytes express a diversity of genes and exhibit novel immune functions in parasitism. <i>Scientific Reports</i> , 2016, 6, 26967.	3.3	20
61	The autophagy pathway participates in resistance to tomato yellow leaf curl virus infection in whiteflies. <i>Autophagy</i> , 2016, 12, 1560-1574.	9.1	108
62	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, 265-276.	3.0	22
63	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, 2860-2861.	0.7	12
64	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, 2783-2784.	0.7	12
65	The genus <i>Spathius</i> Nees, 1818 (Hymenoptera: Braconidae, Doryctinae) from China with a key to species. <i>Zootaxa</i> , 2015, 3960, 1.	0.5	13
66	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	6
67	The discovery of the genus <i>Protodacnusa</i> Griffiths, 1964 (Hymenoptera: Braconidae, Alysiniinae) in China, with descriptions of six new species. <i>Zootaxa</i> , 2015, 3990, 355.	0.5	0
68	The <i>lacteus</i> -, <i>laspeyresiella</i> - and <i>mycetophilus</i> -groups of <i>Apanteles</i> Foerster, 1862 (Hymenoptera, Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 8	0.5	8
69	The Endoparasitoid, <i>Cotesia vestalis</i> , Regulates Host Physiology by Reprogramming the Neuropeptide Transcriptional Network. <i>Scientific Reports</i> , 2015, 5, 8173.	3.3	22
70	The <i>grandiculus</i> - and <i>metacarpalis</i> -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.	0.5	10
71	The genus <i>Ademon</i> Haliday (Hymenoptera: Braconidae: Opiinae) from China, with descriptions of two new species. <i>Zootaxa</i> , 2014, 3794, 294.	0.5	2
72	Review of the genus <i>Cystomutilla</i> Andr�, 1896 (Hymenoptera: Mutillidae: Sphaerophthalminae:) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 6	0.5	5

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73	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.	1.5	13
74	The subgenus <i>Choeras</i> Mason, 1981 of genus <i>Apanteles</i> Foerster, 1862 (Hymenoptera, Braconidae,) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5	0.5	14
75	The genus <i>Polystenus</i> (Hymenoptera: Braconidae: Doryctinae)in China, with descriptions of two new species. <i>Journal of Insect Science</i> , 2014, 14, 66.	1.5	3
76	Redescription of Aquatic Grass Inhabiting <i>Frankliniella zizaniophila</i> (Thripidae: Thripinae) With Remarks on Its Systematic Position Within the Genus <i>Frankliniella</i> (Thysanoptera). <i>Journal of Insect Science</i> , 2014, 14, 154.	1.5	0
77	Two mitochondrial genomes from the families Bethyidae 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.	2.7	57
78	Flower-visiting insects and their potential impact on transgene flow in rice. <i>Journal of Applied Ecology</i> , 2014, 51, 1357-1365.	4.0	27
79	Deep sequencing of <i>Cotesia vestalis</i> bracovirus reveals the complexity of a polydnavirus genome. <i>Virology</i> , 2011, 414, 42-50.	2.4	70
80	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	13
81	The Genus <i>Minanga</i> 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	7
82	The genus <i>Asiabregma</i> Belokobylskij, Zaldivar & Maeto (Hymenoptera: Braconidae) from China, with description of a new species. <i>Entomological Science</i> , 2009, 12, 411-415.	0.6	1
83	Two new species of genus <i>Chablisea</i> Gauld et Dubois, 2006 (Hymenoptera: Ichneumonidae: Pimplinae) from China. <i>Biologia (Poland)</i> , 2009, 64, 1165-1169.	1.5	3
84	Characterization of an Î ⁹ Bâ€like gene in <i>Cotesia vestalis</i> polydnavirus. <i>Archives of Insect Biochemistry and Physiology</i> , 2008, 68, 71-78.	1.5	8
85	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.	3.0	8
86	Aphid dispersal flight disseminates fungal pathogens and parasitoids as natural control agents of aphids. <i>Ecological Entomology</i> , 2007, 32, 97-104.	2.2	49
87	The discovery of the genus <i>Spinadesha</i> (Hymenoptera, Braconidae, Braconinae) in China, with description of a new species. <i>Biologia (Poland)</i> , 2006, 61, 145-147.	1.5	7
88	A Review of <i>Bracon</i> (Rostrobracon) (Hymenoptera: Braconidae: Braconinae) from China, With Description of One New Species. <i>Oriental Insects</i> , 2004, 38, 341-345.	0.3	5
89	Title is missing!. <i>BioControl</i> , 2003, 48, 515-527.	2.0	50
90	THE DISCOVERY OF THE GENUS <i>SHELFORDIA</i> CAMERON (HYMENOPTERA: BRACONTOAE: BRACONINAE) IN CHINA, WITH DESCRIPTION OF ONE NEW SPECIES. <i>Insect Science</i> , 2003, 10, 215-220.	3.0	1

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91	Books Received. <i>Journal of Islamic Studies</i> , 2003, 14, 119-125.	0.0	0