

Hao-Liang Chen

List of Publications by Year in descending order

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Version: 2024-02-01

20
papers

248
citations

933447

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996975

15
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21
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docs citations

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189
citing authors

#	ARTICLE	IF	CITATIONS
1	Parasitism and Host-Location Preference in <i>Habrobracon hebetor</i> (Hymenoptera: Braconidae) (Hymenoptera: Tj ETQq1 1 0.784314 rgBT /Overlock 610-615.	1.8	35
2	Maternal and progeny quality of <i>Habrobracon hebetor</i> Say (Hymenoptera: Braconidae) after cold storage. <i>Biological Control</i> , 2011, 58, 255-261.	3.0	27
3	Transcriptome Analysis and Identification of Insecticide Tolerance-Related Genes after Exposure to Insecticide in <i>Sitobion avenae</i> . <i>Genes</i> , 2019, 10, 951.	2.4	24
4	De novo Sequencing, Assembly and Characterization of Antennal Transcriptome of <i>Anomala corpulenta</i> Motschulsky (Coleoptera: Rutelidae). <i>PLoS ONE</i> , 2014, 9, e114238.	2.5	22
5	Performance of diapausing parasitoid wasps, <i>Habrobracon hebetor</i> , after cold storage. <i>Biological Control</i> , 2013, 64, 186-194.	3.0	21
6	Induction of Reproductive Diapause in <i>Habrobracon hebetor</i> (Hymenoptera: Braconidae) When Reared at Different Photoperiods at Low Temperatures. <i>Environmental Entomology</i> , 2012, 41, 697-705.	1.4	15
7	De novo sequencing and characterization of the <i>Bradysia odoriphaga</i> (Diptera: Sciaridae) larval transcriptome. <i>Comparative Biochemistry and Physiology Part D: Genomics and Proteomics</i> , 2015, 16, 20-27.	1.0	13
8	Using Next-Generation Sequencing to Detect Differential Expression Genes in <i>Bradysia odoriphaga</i> after Exposure to Insecticides. <i>International Journal of Molecular Sciences</i> , 2017, 18, 2445.	4.1	12
9	Evaluation of reference genes for quantitative real-time PCR normalization in the scarab beetle <i>Holotrichia oblita</i> . <i>PLoS ONE</i> , 2020, 15, e0240972.	2.5	12
10	Transcriptome Analysis of Detoxification-Related Genes in <i>Spodoptera frugiperda</i> (Lepidoptera: Tj ETQq0 0 0 rgBT /Overlock 10 T 1.58 11	1.5	11
11	Survey of the <i>Bradysia odoriphaga</i> Transcriptome Using PacBio Single-Molecule Long-Read Sequencing. <i>Genes</i> , 2019, 10, 481.	2.4	8
12	PBP1 plays key roles in sex pheromone reception of the fall armyworm. <i>International Journal of Biological Macromolecules</i> , 2022, 214, 162-169.	7.5	8
13	Transcript analysis and expression profiling of three heat shock protein 70 genes in the ectoparasitoid <i>Habrobracon hebetor</i> (Hymenoptera: Braconidae). <i>Insect Science</i> , 2014, 21, 415-428.	3.0	6
14	Influence of Constant Temperature on Reproductive Parameters of <i>Holotrichia oblita</i> (Coleoptera: Scarabaeidae). <i>Journal of Insect Science</i> , 2015, 15, 93.	1.5	4
15	Genome-wide analysis of developmental stage-specific transcriptome in <i>Bradysia odoriphaga</i> . <i>Comparative Biochemistry and Physiology Part D: Genomics and Proteomics</i> , 2019, 30, 45-54.	1.0	4
16	Effect of Photoperiod on Longevity, Food Consumption, and Reproduction of <i>Holotrichia oblita</i> (Coleoptera: Scarabaeidae). <i>Environmental Entomology</i> , 2021, 50, 1151-1157.	1.4	3
17	Expression Analysis of Odorant-Binding Protein Genes and Chemosensory Protein Genes in <i>Anomala corpulenta</i> Motschulsky (Coleoptera: Scarabaeidae). <i>Journal of the Kansas Entomological Society</i> , 2019, 92, 376.	0.2	3
18	Characterizations of botanical attractant of <i>Halyomorpha halys</i> and selection of relevant deorphanization candidates via computational approach. <i>Scientific Reports</i> , 2022, 12, 4170.	3.3	3

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19	Comparative analyses of six cytochrome P450 genes and their roles in differential insecticide susceptibilities between the red flour beetle and the confused flour beetle. <i>Journal of Stored Products Research</i> , 2022, 96, 101951.	2.6	1
20	Transcriptome analysis of <i>Holotrichia oblita</i> reveals differentially expressed unigenes related to reproduction and development under different photoperiods. <i>Comparative Biochemistry and Physiology Part D: Genomics and Proteomics</i> , 2022, 42, 100959.	1.0	0