

Teng-Fei Shi

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2560739/publications.pdf>

Version: 2024-02-01

9
papers

161
citations

1478505

6
h-index

1474206

9
g-index

9
all docs

9
docs citations

9
times ranked

191
citing authors

#	ARTICLE	IF	CITATIONS
1	Sublethal Effects of the Neonicotinoid Insecticide Thiamethoxam on the Transcriptome of the Honey Bees (Hymenoptera: Apidae). <i>Journal of Economic Entomology</i> , 2017, 110, 2283-2289.	1.8	57
2	Metabolomic analysis of honey bee, <i>Apis mellifera</i> L. response to thiacloprid. <i>Pesticide Biochemistry and Physiology</i> , 2018, 152, 17-23.	3.6	34
3	Influence of the Neonicotinoid Insecticide Thiamethoxam on miRNA Expression in the Honey Bee (Hymenoptera: Apidae). <i>Journal of Insect Science</i> , 2017, 17, .	1.5	21
4	The microRNA ame-miR-279a regulates sucrose responsiveness of forager honey bees (<i>Apis mellifera</i>). <i>Insect Biochemistry and Molecular Biology</i> , 2017, 90, 34-42.	2.7	15
5	Effects of Field-Realistic Concentrations of Carbendazim on Survival and Physiology in Forager Honey Bees (Hymenoptera: Apidae). <i>Journal of Insect Science</i> , 2018, 18, .	1.5	11
6	Genetic structure of Mount Huang honey bee (<i>Apis cerana</i>) populations: evidence from microsatellite polymorphism. <i>Hereditas</i> , 2016, 153, 8.	1.4	9
7	Age and Behavior-Dependent Differential miRNAs Expression in the Hypopharyngeal Glands of Honeybees (<i>Apis mellifera</i> L.). <i>Insects</i> , 2021, 12, 764.	2.2	7
8	Transcriptomic analysis to elucidate the response of honeybees (Hymenoptera: Apidae) to amitraz treatment. <i>PLoS ONE</i> , 2020, 15, e0228933.	2.5	6
9	Proteome analysis reveals the molecular basis of honeybee brain and midgut response to sulfoxaflor. <i>Pesticide Biochemistry and Physiology</i> , 2022, 186, 105168.	3.6	1