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List of Publications by Year in descending order

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Version: 2024-02-01

16
papers

643
citations

840776

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#	ARTICLE	IF	CITATIONS
1	The complete mitochondrial genome of the bag-shelter moth <i>Ochrogaster lunifer</i> (Lepidoptera.) Tj ETQq1 1 0.784314 rgBT /Overlock 10	2.8	208
2	Quaternary history and contemporary patterns in a currently expanding species. BMC Evolutionary Biology, 2009, 9, 220.	3.2	83
3	Presence of specific symbiotic bacteria in flies of the subfamily Tephritinae (Diptera Tephritidae) and their phylogenetic relationships: proposal of 'Candidatus <i>Stammerula tephritidis</i> '. International Journal of Systematic and Evolutionary Microbiology, 2008, 58, 1277-1287.	1.7	59
4	Characterization and evolution of two bacteriome-inhabiting symbionts in cixiid planthoppers (Hemiptera: Fulgoromorpha: Pentastirini). Environmental Microbiology, 2009, 11, 3265-3279.	3.8	53
5	The role of topography in structuring the demographic history of the pine processionary moth, <i>Thaumetopoea pityocampa</i> (Lepidoptera: Notodontidae). Journal of Biogeography, 2010, 37, 1478-1490.	3.0	38
6	Evidence of two lineages of the symbiont <i>Candidatus Erwinia dacicola</i> ™ in Italian populations of <i>Bactrocera oleae</i> (Rossi) based on 16S rRNA gene sequences. International Journal of Systematic and Evolutionary Microbiology, 2012, 62, 179-187.	1.7	31
7	Identification and biological traits of a planthopper from the genus <i>Pentastiridius</i> (Hemiptera:) Tj ETQq1 1 0.784314 rgBT /Overlock 10	1.25	30
8	Phylogenetic relationships between flies of the Tephritinae subfamily (Diptera, Tephritidae) and their symbiotic bacteria. Molecular Phylogenetics and Evolution, 2010, 56, 312-326.	2.7	26
9	Metagenomic analysis reveals changes of the <i>Drosophila suzukii</i> microbiota in the newly colonized regions. Insect Science, 2018, 25, 833-846.	3.0	25
10	Host and Phenology Shifts in the Evolution of the Social Moth Genus <i>Thaumetopoea</i> . PLoS ONE, 2013, 8, e57192.	2.5	24
11	A population genetic study of the egg parasitoid <i>Baryscapus servadeii</i> reveals large scale automictic parthenogenesis and almost fixed homozygosity. Biological Control, 2019, 139, 104097.	3.0	14
12	Genetic differentiation of the pine processionary moth at the southern edge of its range: contrasting patterns between mitochondrial and nuclear markers. Ecology and Evolution, 2016, 6, 4274-4288.	1.9	13
13	A unique midgut-associated bacterial community hosted by the cave beetle <i>Cansiliella servadeii</i> (Coleoptera: Leptodirini) reveals parallel phylogenetic divergences from universal gut-specific ancestors. BMC Microbiology, 2013, 13, 129.	3.3	11
14	Evidence of potential hybridization in the <i>Thaumetopoea pityocampa</i> - <i>Wilkisoni</i> complex. Agricultural and Forest Entomology, 2018, 20, 9-17.	1.3	11
15	Testing for host-associated differentiation in two egg parasitoids of a forest herbivore. Entomologia Experimentalis Et Applicata, 2012, 145, 124-133.	1.4	10
16	High genetic diversity in the <i>Culex pipiens</i> complex from a West Nile Virus epidemic area in Southern Europe. Parasites and Vectors, 2016, 9, 150.	2.5	7