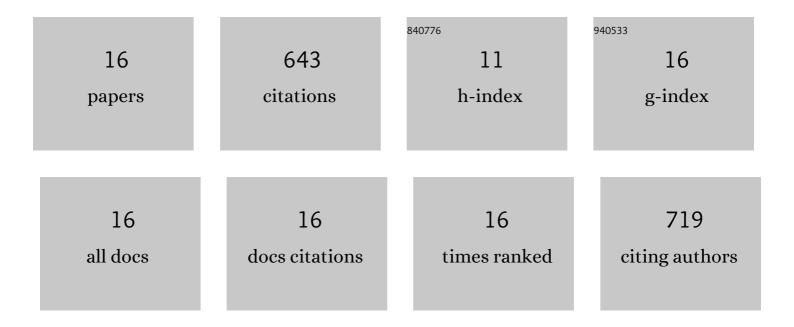
Mauro Simonato

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/11172827/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	A population genetic study of the egg parasitoid Baryscapus servadeii reveals large scale automictic parthenogenesis and almost fixed homozygosity. Biological Control, 2019, 139, 104097.	3.0	14
2	Evidence of potential hybridization in the <i><scp>T</scp>haumetopoea pityocampaâ€wilkinsoni</i> complex. Agricultural and Forest Entomology, 2018, 20, 9-17.	1.3	11
3	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
4	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
5	High genetic diversity in the Culex pipiens complex from a West Nile Virus epidemic area in Southern Europe. Parasites and Vectors, 2016, 9, 150.	2.5	7
6	A unique midgut-associated bacterial community hosted by the cave beetle Cansiliella servadeii(Coleoptera: Leptodirini) reveals parallel phylogenetic divergences from universal gut-specific ancestors. BMC Microbiology, 2013, 13, 129.	3.3	11
7	Host and Phenology Shifts in the Evolution of the Social Moth Genus Thaumetopoea. PLoS ONE, 2013, 8, e57192.	2.5	24
8	Testing for hostâ€associated differentiation in two egg parasitoids of a forest herbivore. Entomologia Experimentalis Et Applicata, 2012, 145, 124-133.	1.4	10
9	Evidence of two lineages of the symbiont â€~Candidatus Erwinia dacicola' in Italian populations of Bactrocera oleae (Rossi) based on 16S rRNA gene sequences. International Journal of Systematic and Evolutionary Microbiology, 2012, 62, 179-187.	1.7	31
10	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
11	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
12	Quaternary history and contemporary patterns in a currently expanding species. BMC Evolutionary Biology, 2009, 9, 220.	3.2	83
13	Characterization and evolution of two bacteriomeâ€inhabiting symbionts in cixiid planthoppers (Hemiptera: Fulgoromorpha: Pentastirini). Environmental Microbiology, 2009, 11, 3265-3279.	3.8	53
14	Identification and biological traits of a planthopper from the genus Pentastiridius (Hemiptera:) Tj ETQq0 0 0 rgB	[/Qverlock	10 Tf 50 222

 15
 The complete mitochondrial genome of the bag-shelter moth Ochrogaster lunifer (Lepidoptera,) Tj ETQq1 1 0.784314 rgBT /Qyerlock 10

 16
 Presence of specific symbiotic bacteria in flies of the subfamily Tephritinae (Diptera Tephritidae) and their phylogenetic relationships: proposal of 'Candidatus Stammerula tephritidis'. International

 16
 Image: Symbol of Systematic and Evolutionary Microbiology, 2008, 58, 1277-1287.