

# Mauro Simonato

## List of Publications by Year in descending order

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Version: 2024-02-01

16  
papers

643  
citations

840776

11  
h-index

940533

16  
g-index

16  
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16  
docs citations

16  
times ranked

719  
citing authors

#	ARTICLE	IF	CITATIONS
1	A population genetic study of the egg parasitoid <i>Baryscapus servadeii</i> reveals large scale automictic parthenogenesis and almost fixed homozygosity. <i>Biological Control</i> , 2019, 139, 104097.	3.0	14
2	Evidence of potential hybridization in the <i>Thaumetopoea pityocampa-wilkinsoni</i> complex. <i>Agricultural and Forest Entomology</i> , 2018, 20, 9-17.	1.3	11
3	Metagenomic analysis reveals changes of the <i>Drosophila suzukii</i> microbiota in the newly colonized regions. <i>Insect Science</i> , 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. <i>Ecology and Evolution</i> , 2016, 6, 4274-4288.	1.9	13
5	High genetic diversity in the <i>Culex pipiens</i> complex from a West Nile Virus epidemic area in Southern Europe. <i>Parasites and Vectors</i> , 2016, 9, 150.	2.5	7
6	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. <i>BMC Microbiology</i> , 2013, 13, 129.	3.3	11
7	Host and Phenology Shifts in the Evolution of the Social Moth Genus <i>Thaumetopoea</i> . <i>PLoS ONE</i> , 2013, 8, e57192.	2.5	24
8	Testing for host-associated differentiation in two egg parasitoids of a forest herbivore. <i>Entomologia Experimentalis Et Applicata</i> , 2012, 145, 124-133.	1.4	10
9	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. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 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). <i>Journal of Biogeography</i> , 2010, 37, 1478-1490.	3.0	38
11	Phylogenetic relationships between flies of the Tephritinae subfamily (Diptera, Tephritidae) and their symbiotic bacteria. <i>Molecular Phylogenetics and Evolution</i> , 2010, 56, 312-326.	2.7	26
12	Quaternary history and contemporary patterns in a currently expanding species. <i>BMC Evolutionary Biology</i> , 2009, 9, 220.	3.2	83
13	Characterization and evolution of two bacteriome-inhabiting symbionts in cixiid planthoppers (Hemiptera: Fulgoromorpha: Pentastirini). <i>Environmental Microbiology</i> , 2009, 11, 3265-3279.	3.8	53
14	Identification and biological traits of a planthopper from the genus <i>Pentastiridius</i> (Hemiptera: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 222	1.2	30
15	The complete mitochondrial genome of the bag-shelter moth <i>Ochrogaster lunifer</i> (Lepidoptera,) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 222	2.8	208
16	Presence of specific symbiotic bacteria in flies of the subfamily Tephritinae (Diptera Tephritidae) and their phylogenetic relationships: proposal of 'Candidatus <i>Stammerula tephritidis</i> '. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2008, 58, 1277-1287.	1.7	59