

# Panagiota Koskinioti

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4464597/publications.pdf>

Version: 2024-02-01

10  
papers

464  
citations

1040056

9  
h-index

1281871

11  
g-index

12  
all docs

12  
docs citations

12  
times ranked

797  
citing authors

#	ARTICLE	IF	CITATIONS
1	Genetic sexing strains for the population suppression of the mosquito vector <i>Aedes aegypti</i> . Philosophical Transactions of the Royal Society B: Biological Sciences, 2021, 376, 20190808.	4.0	24
2	Next-generation biological control: the need for integrating genetics and genomics. Biological Reviews, 2020, 95, 1838-1854.	10.4	67
3	The impact of fruit fly gut bacteria on the rearing of the parasitic wasp <i>Diachasmimorpha longicaudata</i> . Entomologia Experimentalis Et Applicata, 2020, 168, 541-559.	1.4	9
4	Manipulation of insect gut microbiota towards the improvement of <i>Bactrocera oleae</i> artificial rearing. Entomologia Experimentalis Et Applicata, 2020, 168, 523-540.	1.4	18
5	<i>Aedes aegypti</i> lines for combined sterile insect technique and incompatible insect technique applications: the importance of host genomic background. Entomologia Experimentalis Et Applicata, 2020, 168, 560-572.	1.4	24
6	Irradiation induced inversions suppress recombination between the M locus and morphological markers in <i>Aedes aegypti</i> . BMC Genetics, 2020, 21, 142.	2.7	15
7	<i>Maleness-on-the-Y</i> ( <i>MoY</i> ) orchestrates male sex determination in major agricultural fruit fly pests. Science, 2019, 365, 1457-1460.	12.6	88
8	The effects of geographic origin and antibiotic treatment on the gut symbiotic communities of <i>Bactrocera oleae</i> populations. Entomologia Experimentalis Et Applicata, 2019, 167, 197-208.	1.4	56
9	Housekeeping in Tephritid insects: the best gene choice for expression analyses in the medfly and the olive fly. Scientific Reports, 2017, 7, 45634.	3.3	30
10	The whole genome sequence of the Mediterranean fruit fly, <i>Ceratitis capitata</i> (Wiedemann), reveals insights into the biology and adaptive evolution of a highly invasive pest species. Genome Biology, 2016, 17, 192.	8.8	130