## Olle Terenius

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7137682/publications.pdf

Version: 2024-02-01

218677 175258 2,941 61 26 52 citations h-index g-index papers 62 62 62 3203 all docs docs citations times ranked citing authors

#	Article	IF	CITATIONS
1	RNA interference in Lepidoptera: An overview of successful and unsuccessful studies and implications for experimental design. Journal of Insect Physiology, 2011, 57, 231-245.	2.0	729
2	16S rRNA Gene-Based Identification of Midgut Bacteria from Field-Caught Anopheles gambiae Sensu Lato and A. funestus Mosquitoes Reveals New Species Related to Known Insect Symbionts. Applied and Environmental Microbiology, 2005, 71, 7217-7223.	3.1	177
3	Hemolingene silencing by ds-RNA injected into Cecropia pupae is lethal to next generation embryos. Insect Molecular Biology, 2002, 11, 267-271.	2.0	148
4	Parasiteâ€specific immune response in adult Drosophila melanogaster: a genomic study. EMBO Reports, 2004, 5, 207-212.	4.5	120
5	Male Fertility of Malaria Parasites Is Determined by GCS1, a Plant-Type Reproduction Factor. Current Biology, 2008, 18, 607-613.	3.9	118
6	Developmental succession of the microbiome of Culex mosquitoes. BMC Microbiology, 2015, 15, 140.	3.3	118
7	Midgut bacterial dynamics in Aedes aegypti. FEMS Microbiology Ecology, 2012, 80, 556-565.	2.7	117
8	Bacterial associations reveal spatial population dynamics in Anopheles gambiae mosquitoes. Scientific Reports, 2016, 6, 22806.	3.3	114
9	Exposure of rat brain to 915 MHz GSM microwaves induces changes in gene expression but not double stranded DNA breaks or effects on chromatin conformation. Bioelectromagnetics, 2006, 27, 295-306.	1.6	108
10	Malpighian tubules are important determinants of Pseudomonas transstadial transmission and longtime persistence in Anopheles stephensi. Parasites and Vectors, 2015, 8, 36.	2.5	78
11	Isolation and identification of culturable bacteria from wild Anopheles culicifacies, a first step in a paratransgenesis approach. Parasites and Vectors, 2014, 7, 419.	2.5	75
12	The gut microbiota of the pine weevil is similar across Europe and resembles that of other coniferâ€feeding beetles. Molecular Ecology, 2016, 25, 4014-4031.	3.9	75
13	Baculovirus and dsRNA induce Hemolin, but no antibacterial activity, in Antheraea pernyi. Insect Molecular Biology, 2004, 13, 399-405.	2.0	70
14	Molecular Genetic Manipulation of Vector Mosquitoes. Cell Host and Microbe, 2008, 4, 417-423.	11.0	63
15	RNA interference of Hemolin causes depletion of phenoloxidase activity in Hyalophora cecropia.  Developmental and Comparative Immunology, 2007, 31, 571-575.	2.3	56
16	Janibacter anophelis sp. nov., isolated from the midgut of Anopheles arabiensis. International Journal of Systematic and Evolutionary Microbiology, 2006, 56, 389-392.	1.7	52
17	16S rRNA Gene Sequences from Bacteria Associated with Adult Anopheles darlingi (Diptera: Culicidae) Mosquitoes. Journal of Medical Entomology, 2008, 45, 172-175.	1.8	52
18	West Nile Virus in Mosquitoes of Iranian Wetlands. Vector-Borne and Zoonotic Diseases, 2015, 15, 750-754.	1.5	49

#	Article	IF	Citations
19	The environment and species affect gut bacteria composition in laboratory co-cultured Anopheles gambiae and Aedes albopictus mosquitoes. Scientific Reports, 2020, 10, 3352.	3.3	47
20	16S rRNA Gene Sequences from Bacteria Associated with Adult <l>Anopheles darlingi</l> (Diptera: Culicidae) Mosquitoes. Journal of Medical Entomology, 2008, 45, 172-175.	1.8	42
21	Antheraea pernyi (Lepidoptera: Saturniidae) and Its Importance in Sericulture, Food Consumption, and Traditional Chinese Medicine. Journal of Economic Entomology, 2017, 110, 1404-1411.	1.8	42
22	Hemolin—A lepidopteran anti-viral defense factor?. Developmental and Comparative Immunology, 2008, 32, 311-316.	2.3	39
23	Presence of Aedes and Anopheles mosquito larvae is correlated to bacteria found in domestic water-storage containers. FEMS Microbiology Ecology, 2018, 94, .	2.7	32
24	Persistence of sociality in group dynamics of dairy cattle. Applied Animal Behaviour Science, 2020, 223, 104921.	1.9	30
25	Thorsellia anophelis gen. nov., sp. nov., a new member of the Gammaproteobacteria. International Journal of Systematic and Evolutionary Microbiology, 2006, 56, 335-338.	1.7	28
26	Cloning, expression and phylogenetic analysis of , from the Chinese oak silkmoth,. Developmental and Comparative Immunology, 2005, 29, 853-864.	2.3	27
27	Re-introducing bacteria in mosquitoesâ€"A method for determination of mosquito feeding preferences based on coloured sugar solutions. Acta Tropica, 2006, 99, 173-183.	2.0	27
28	Bacterial, but not baculoviral infections stimulate Hemolin expression in noctuid moths. Developmental and Comparative Immunology, 2009, 33, 1176-1185.	2.3	23
29	A fungal metabolite masks the host plant odor for the pine weevil ( Hylobius abietis ). Fungal Ecology, 2015, 13, 103-111.	1.6	23
30	Genetic Characterization of a Novel Iflavirus Associated with Vomiting Disease in the Chinese Oak Silkmoth Antheraea pernyi. PLoS ONE, 2014, 9, e92107.	2.5	23
31	Proposal of Thorsellia kenyensis sp. nov. and Thorsellia kandunguensis sp. nov., isolated from larvae of Anopheles arabiensis, as members of the family Thorselliaceae fam. nov International Journal of Systematic and Evolutionary Microbiology, 2015, 65, 444-451.	1.7	22
32	Characterization of Bacterial Communities in Breeding Waters of Anopheles darlingi in Manaus in the Amazon Basin Malaria-Endemic Area. Microbial Ecology, 2019, 78, 781-791.	2.8	21
33	Molecular Characterization of Mosquitoes (Diptera: Culicidae) in Northwestern Iran by Using rDNA-ITS2. Japanese Journal of Infectious Diseases, 2016, 69, 319-322.	1.2	20
34	Single nucleotide polymorphism analysis of the ITS2 region of two sympatric malaria mosquito species in Sweden: <i>Anopheles daciae</i> and <i>Anopheles messeae</i> Medical and Veterinary Entomology, 2020, 34, 364-368.	1.5	15
35	Culturable bacteria associated with Anopheles darlingi and their paratransgenesis potential. Malaria Journal, 2021, 20, 40.	2.3	14
36	Studies on the transmission and tissue distribution of Antheraea pernyi iflavirus in the Chinese oak silkmoth Antheraea pernyi. Virology, 2017, 502, 171-175.	2.4	13

#	Article	IF	CITATIONS
37	Characterization of a Novel RNA Virus Discovered in the Autumnal Moth Epirrita autumnata in Sweden. Viruses, 2017, 9, 214.	3.3	13
38	Chemodiversity and biodiversity of fungi associated with the pine weevil Hylobius abietis. Fungal Biology, 2015, 119, 738-746.	2.5	12
39	New and Earlier Records of Ticks (Acari: Ixodidae, Argasidae) from Guinea-Bissau. Journal of Medical Entomology, 2000, 37, 973-976.	1.8	10
40	Coetzeea brasiliensis gen. nov., sp. nov. isolated from larvae of Anopheles darlingi. International Journal of Systematic and Evolutionary Microbiology, 2016, 66, 5211-5217.	1.7	10
41	Wikipedia in Health Professional Schools: from an Opponent to an Ally. Medical Science Educator, 2021, 31, 2209-2216.	1.5	9
42	Development of an instrument for early detection of dementia in people with Down syndrome. Journal of Intellectual and Developmental Disability, 2002, 27, 325-345.	1.6	8
43	Injection of An. stephensi Embryos to Generate Malaria-resistant Mosquitoes. Journal of Visualized Experiments, 2007, , 216.	0.3	8
44	Antifeedants Produced by Bacteria Associated with the Gut of the Pine Weevil Hylobius abietis. Microbial Ecology, 2017, 74, 177-184.	2.8	8
45	Heterobasidion-growth inhibiting Bacillus subtilis A18 exhibits medium- and age-dependent production of lipopeptides. Microbiological Research, 2019, 223-225, 129-136.	5.3	8
46	Larvicidal Activities against Aedes aegypti of Supernatant and Pellet Fractions from Cultured Bacillus spp. Isolated from Amazonian Microenvironments. Tropical Medicine and Infectious Disease, 2021, 6, 104.	2.3	6
47	Supplementation of Lactobacillus casei reduces the mortality of Bombyx mori larvae challenged by Nosema bombycis. BMC Research Notes, 2021, 14, 398.	1.4	6
48	Global similarity, and some key differences, in the metagenomes of Swedish varroa-surviving and varroa-susceptible honeybees. Scientific Reports, 2021, 11, 23214.	3.3	5
49	Genetic characterisation of an Iflavirus associated with a vomiting disease in the Indian tropical tasar silkworm, Antheraea mylitta. Virus Research, 2022, 311, 198703.	2.2	5
50	cDNA Cloning and Expression Analysis of Pattern Recognition Proteins from the Chinese Oak Silkmoth, Antheraea pernyi. Insects, 2012, 3, 1093-1104.	2.2	4
51	The phytopathogen powdery mildew affects food-searching behavior and survival of Coccinella septempunctata. Arthropod-Plant Interactions, 2018, 12, 685-690.	1.1	4
52	Molecular analysis of the mitochondrial markers COI, 12S rDNA and 16S rDNA for six species of Iranian scorpions. BMC Research Notes, 2021, 14, 40.	1.4	4
53	Evidence for the presence of <i>Ctenocephalides orientis</i> in livestock dwellings in northwest Iran. Medical and Veterinary Entomology, 2018, 32, 383-387.	1.5	3
54	Development and optimization of a TaqMan assay for Nosema bombycis, causative agent of p $\tilde{A}$ ©brine disease in Bombyx mori silkworm, based on the $\hat{I}^2$ -tubulin gene. Journal of Microbiological Methods, 2021, 186, 106238.	1.6	3

## OLLE TERENIUS

#	Article	IF	CITATION
55	Extracts of Amazonian Fungi With Larvicidal Activities Against Aedes aegypti. Frontiers in Microbiology, 2021, 12, 743246.	3.5	3
56	An investigation into the effects of infection and ORF expression patterns of the Indian bidensovirus isolate (BmBDV) infecting the silkworm Bombyx mori. VirusDisease, 2022, 33, 76-83.	2.0	2
57	Evaporated Extracts of Samples for Pesticide Residue Analysis Simplifies Transport from Remote Places. Bulletin of Environmental Contamination and Toxicology, 1997, 58, 341-347.	2.7	1
58	Premature Proposal of the Pine Weevil as a Vector of a Human Pathogen. Journal of Clinical Microbiology, 2014, 52, 4115-4115.	3.9	1
59	Sensitivity of Polyvoltine Thai Strains of Bombyx mori to a BmNPV Isolate From Mahasarakham. Journal of Insect Science, 2020, 20, .	1.5	1
60	Windsurfing in Mute Swans (Cygnus olor). Wilson Journal of Ornithology, 2016, 128, 628-631.	0.2	0
61	Molecular characterization of Ribosomal DNA (ITS2) of hard ticks in Iran: understanding the conspecificity of Dermacentor marginatus and D. niveus. BMC Research Notes, 2020, 13, 478.	1.4	0