

# Norah P Saarman

## List of Publications by Year in descending order

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Version: 2024-02-01

23  
papers

480  
citations

686830

13  
h-index

752256

20  
g-index

25  
all docs

25  
docs citations

25  
times ranked

710  
citing authors

#	ARTICLE	IF	CITATIONS
1	Phylogenomic analysis of Syngnathidae reveals novel relationships, origins of endemic diversity and variable diversification rates. <i>BMC Biology</i> , 2022, 20, 75.	1.7	19
2	A machine-learning approach to map landscape connectivity in <i>Aedes aegypti</i> with genetic and environmental data. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	3.3	27
3	Big Data in Conservation Genomics: Boosting Skills, Hedging Bets, and Staying Current in the Field. <i>Journal of Heredity</i> , 2021, 112, 313-327.	1.0	10
4	A machine learning approach to integrating genetic and ecological data in tsetse flies ( <i>Glossina</i> ) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 1762-1777.	1.5	6
5	Infection with endosymbiotic <i>Spiroplasma</i> disrupts tsetse ( <i>Glossina fuscipes fuscipes</i> ) metabolic and reproductive homeostasis. <i>PLoS Pathogens</i> , 2021, 17, e1009539.	2.1	9
6	Fungal spore diversity, community structure, and traits across a vegetation mosaic. <i>Fungal Ecology</i> , 2020, 45, 100920.	0.7	11
7	Phylogeography and population structure of the tsetse fly <i>Glossina pallidipes</i> in Kenya and the Serengeti ecosystem. <i>PLoS Neglected Tropical Diseases</i> , 2020, 14, e0007855.	1.3	6
8	Spatio-temporal distribution of <i>Spiroplasma</i> infections in the tsetse fly ( <i>Glossina fuscipes fuscipes</i> ) in northern Uganda. <i>PLoS Neglected Tropical Diseases</i> , 2019, 13, e0007340.	1.3	22
9	The population genomics of multiple tsetse fly ( <i>Glossina fuscipes fuscipes</i> ) admixture zones in Uganda. <i>Molecular Ecology</i> , 2019, 28, 66-85.	2.0	11
10	Ectomycorrhizas and tree seedling establishment are strongly influenced by forest edge proximity but not soil inoculum. <i>Ecological Applications</i> , 2019, 29, e01867.	1.8	19
11	How Population Decline Can Impact Genetic Diversity: a Case Study of Eelgrass ( <i>Zostera marina</i> ) in Morro Bay, California. <i>Estuaries and Coasts</i> , 2018, 41, 2356-2367.	1.0	7
12	A spatial genetics approach to inform vector control of tsetse flies ( <i>Glossina fuscipes</i> ) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 302 Td (f	0.8	8
13	Genetic Differentiation of <i>Glossina pallidipes</i> Tsetse Flies in Southern Kenya. <i>American Journal of Tropical Medicine and Hygiene</i> , 2018, 99, 945-953.	0.6	8
14	Effective population sizes of a major vector of human diseases, <i>Aedes aegypti</i> . <i>Evolutionary Applications</i> , 2017, 10, 1031-1039.	1.5	47
15	Molecular phylogeny and patterns of diversification in syngnathid fishes. <i>Molecular Phylogenetics and Evolution</i> , 2017, 107, 388-403.	1.2	54
16	Sequence-Based Analysis of Thermal Adaptation and Protein Energy Landscapes in an Invasive Blue Mussel ( <i>Mytilus galloprovincialis</i> ). <i>Genome Biology and Evolution</i> , 2017, 9, 2739-2751.	1.1	20
17	Multiple evolutionary origins of <i>Trypanosoma evansi</i> in Kenya. <i>PLoS Neglected Tropical Diseases</i> , 2017, 11, e0005895.	1.3	27
18	Temporal genetic differentiation in <i>Glossina pallidipes</i> tsetse fly populations in Kenya. <i>Parasites and Vectors</i> , 2017, 10, 471.	1.0	14

#	ARTICLE	IF	CITATIONS
19	Genetic diversity and population structure of the tsetse fly <i>Glossina fuscipes fuscipes</i> (Diptera: Tj ETQq1 1 0.784314 rgBT /Overlock 10 2017, 11, e0005485.	1.3	26
20	Evidence of temporal stability in allelic and mitochondrial haplotype diversity in populations of <i>Glossina fuscipes fuscipes</i> (Diptera: Glossinidae) in northern Uganda. <i>Parasites and Vectors</i> , 2016, 9, 258.	1.0	13
21	Introgression between invasive and native blue mussels (genus <i>M</i> ) in the central California hybrid zone. <i>Molecular Ecology</i> , 2015, 24, 4723-4738.	2.0	60
22	Genetic differentiation across eastern Pacific oceanographic barriers in the threatened seahorse <i>Hippocampus ingens</i> . <i>Conservation Genetics</i> , 2010, 11, 1989-2000.	0.8	30
23	The evolution of conspicuous facultative mimicry in octopuses: an example of secondary adaptation?. <i>Biological Journal of the Linnean Society</i> , 2010, 101, 68-77.	0.7	24