Stéphanie Sherpa

List of Publications by Year in descending order

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#	Article	lF	CITATIONS
1	Demographic inferences and climatic niche modelling shed light on the evolutionary history of the emblematic coldâ€adapted Apollo butterfly at regional scale. Molecular Ecology, 2022, 31, 448-466.	3.9	8
2	Population decline at distribution margins: Assessing extinction risk in the last glacial relictual but still functional metapopulation of a European butterfly. Diversity and Distributions, 2022, 28, 271-290.	4.1	11
3	Genomic Shifts, Phenotypic Clines, and Fitness Costs Associated With Cold Tolerance in the Asian Tiger Mosquito. Molecular Biology and Evolution, 2022, 39, .	8.9	5
4	The evolutionary dynamics of biological invasions: A multiâ€approach perspective. Evolutionary Applications, 2021, 14, 1463-1484.	3.1	48
5	Combining genetic crosses and pool targeted DNAâ€seq for untangling genomic variations associated with resistance to multiple insecticides in the mosquito <i>Aedes aegypti</i> . Evolutionary Applications, 2020, 13, 303-317.	3.1	22
6	Landscape does matter: Disentangling founder effects from natural and humanâ€aided postâ€introduction dispersal during an ongoing biological invasion. Journal of Animal Ecology, 2020, 89, 2027-2042.	2.8	14
7	Cold adaptation in the Asian tiger mosquito's native range precedes its invasion success in temperate regions. Evolution; International Journal of Organic Evolution, 2019, 73, 1793-1808.	2.3	28
8	Predicting the success of an invader: Niche shift versus niche conservatism. Ecology and Evolution, 2019, 9, 12658-12675.	1.9	20
9	Unravelling the invasion history of the Asian tiger mosquito in Europe. Molecular Ecology, 2019, 28, 2360-2377.	3.9	82
10	Genetic diversity and distribution differ between long-established and recently introduced populations in the invasive mosquito Aedes albopictus. Infection, Genetics and Evolution, 2018, 58, 145-156.	2.3	29
11	Refining the biogeographical scenario of the land snail Cornu aspersum aspersum: Natural spatial expansion and human-mediated dispersal in the Mediterranean basin. Molecular Phylogenetics and Evolution, 2018, 120, 218-232.	2.7	23
12	At the Origin of a Worldwide Invasion: Unraveling the Genetic Makeup of the Caribbean Bridgehead Populations of the Dengue Vector Aedes aegypti. Genome Biology and Evolution, 2018, 10, 56-71.	2.5	24