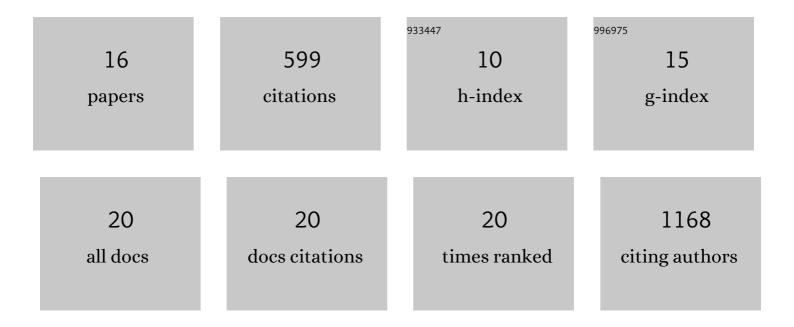
## **Stefan Laurent**

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/9104042/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Demographic Inference Reveals African and European Admixture in the North American <i>Drosophila melanogaster</i> Population. Genetics, 2013, 193, 291-301.	2.9	139
2	Linking a mutation to survival in wild mice. Science, 2019, 363, 499-504.	12.6	126
3	The Evolutionary History of Nebraska Deer Mice: Local Adaptation in the Face of Strong Gene Flow. Molecular Biology and Evolution, 2018, 35, 792-806.	8.9	76
4	The population genomics of rapid adaptation: disentangling signatures of selection and demography in white sands lizards. Molecular Ecology, 2016, 25, 306-323.	3.9	56
5	Mitochondrial DNA Analyses Indicate High Diversity, Expansive Population Growth and High Genetic Connectivity of Vent Copepods (Dirivultidae) across Different Oceans. PLoS ONE, 2016, 11, e0163776.	2.5	29
6	Detecting and Quantifying Changing Selection Intensities from Time-Sampled Polymorphism Data. G3: Genes, Genomes, Genetics, 2016, 6, 893-904.	1.8	27
7	Gene expression evolution in pattern-triggered immunity within <i>Arabidopsis thaliana</i> and across Brassicaceae species. Plant Cell, 2021, 33, 1863-1887.	6.6	27
8	The Demographic History of African Drosophila melanogaster. Genome Biology and Evolution, 2018, 10, 2338-2342.	2.5	26
9	The population genetics of crypsis in vertebrates: recent insights from mice, hares, and lizards. Heredity, 2020, 124, 1-14.	2.6	24
10	Autoregulation of RCO by Low-Affinity Binding Modulates Cytokinin Action and Shapes Leaf Diversity. Current Biology, 2019, 29, 4183-4192.e6.	3.9	21
11	Population Genomics on the Fly: Recent Advances in Drosophila. Methods in Molecular Biology, 2020, 2090, 357-396.	0.9	17
12	Demographic analyses of a new sample of haploid genomes from a Swedish population of Drosophila melanogaster. Scientific Reports, 2020, 10, 22415.	3.3	12
13	Genetic diversity of two Daphnia-infecting microsporidian parasites, based on sequence variation in the internal transcribed spacer region. Parasites and Vectors, 2016, 9, 293.	2.5	8
14	Impact of Genetic Variation in Gene Regulatory Sequences: A Population Genomics Perspective. Frontiers in Genetics, 2021, 12, 660899.	2.3	2
15	CRISPR/Cas9-Mediated Mutagenesis of RCO in Cardamine hirsuta. Plants, 2020, 9, 268.	3.5	1
16	<scp>TEMPLE</scp> : analysing population genetic variation at transcription factor binding sites. Molecular Ecology Resources, 2016, 16, 1428-1434.	4.8	0