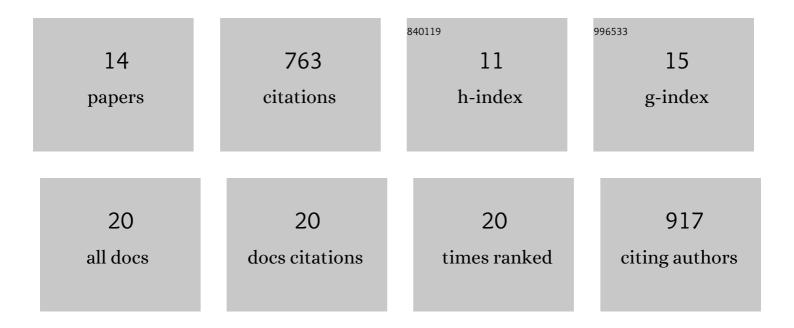
Neda Barghi

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8952391/publications.pdf Version: 2024-02-01



Νέρλ Βλάςμι

#	Article	IF	CITATIONS
1	Neuronal Function and Dopamine Signaling Evolve at High Temperature in Drosophila. Molecular Biology and Evolution, 2020, 37, 2630-2640.	3.5	22
2	Polygenic adaptation: a unifying framework to understand positive selection. Nature Reviews Genetics, 2020, 21, 769-781.	7.7	238
3	Distinct Patterns of Selective Sweep and Polygenic Adaptation in Evolve and Resequence Studies. Genome Biology and Evolution, 2020, 12, 890-904.	1.1	29
4	Rapid sex-specific adaptation to high temperature in Drosophila. ELife, 2020, 9, .	2.8	25
5	A 24 h Age Difference Causes Twice as Much Gene Expression Divergence as 100 Generations of Adaptation to a Novel Environment. Genes, 2019, 10, 89.	1.0	11
6	Shifting the paradigm in Evolve and Resequence studies: From analysis of single nucleotide polymorphisms to selected haplotype blocks. Molecular Ecology, 2019, 28, 521-524.	2.0	18
7	Genetic redundancy fuels polygenic adaptation in Drosophila. PLoS Biology, 2019, 17, e3000128.	2.6	212
8	High-throughput fecundity measurements in Drosophila. Scientific Reports, 2018, 8, 4469.	1.6	16
9	Divergence of the Venom Exogene Repertoire in Two Sister Species of Turriconus. Genome Biology and Evolution, 2017, 9, 2211-2225.	1.1	30
10	<i>Drosophila simulans</i> : A Species with Improved Resolution in Evolve and Resequence Studies. G3: Genes, Genomes, Genetics, 2017, 7, 2337-2343.	0.8	25
11	Structural features of conopeptide genes inferred from partial sequences of the Conus tribblei genome. Molecular Genetics and Genomics, 2016, 291, 411-422.	1.0	27
12	Characterization of the complete mitochondrial genome of Conus tribbleiWalls, 1977. Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis, 2016, 27, 4451-4452.	0.7	7
13	Comparison of the Venom Peptides and Their Expression in Closely RelatedConusSpecies: Insights into Adaptive Post-speciation Evolution ofConusExogenomes. Genome Biology and Evolution, 2015, 7, 1797-1814.	1.1	37
14	High Conopeptide Diversity in Conus tribblei Revealed Through Analysis of Venom Duct Transcriptome Using Two High-Throughput Sequencing Platforms. Marine Biotechnology, 2015, 17, 81-98.	1.1	50