## Joel Atallah

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8342071/publications.pdf

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17 papers	816	11 h-index	996975 15 g-index
papero	Citations	II-IIIQCA	g-mucx
19 all docs	19 docs citations	19 times ranked	1418 citing authors

#	Article	IF	CITATIONS
1	The early embryonic transcriptome of a Hawaiian <i>Drosophila</i> pictureâ€wing fly shows evidence of altered gene expression and novel gene evolution. Journal of Experimental Zoology Part B: Molecular and Developmental Evolution, 2022, 338, 277-291.	1.3	О
2	ClassifyTE: a stacking-based prediction of hierarchical classification of transposable elements. Bioinformatics, 2021, 37, 2529-2536.	4.1	14
3	Evolution of larval segment position across 12 <i>Drosophila</i> species*. Evolution; International Journal of Organic Evolution, 2020, 74, 1409-1422.	2.3	5
4	An <i>in silico</i> model of LINE-1-mediated neoplastic evolution. Bioinformatics, 2020, 36, 4144-4153.	4.1	O
5	Evolution of maternal and zygotic mRNA complements in the early Drosophila embryo. PLoS Genetics, 2018, 14, e1007838.	3.5	23
6	Large-Scale Coding Sequence Change Underlies the Evolution of Postdevelopmental Novelty in Honey Bees. Molecular Biology and Evolution, 2015, 32, 334-346.	8.9	75
7	Evolution ofDrosophilasex comb length illustrates the inextricable interplay between selection and variation. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, E4103-E4109.	7.1	17
8	Comparative validation of the <i>D. melanogaster</i> modENCODE transcriptome annotation. Genome Research, 2014, 24, 1209-1223.	5.5	147
9	Sex-specific repression of dachshund is required for Drosophila sex comb development. Developmental Biology, 2014, 386, 440-447.	2.0	20
10	The making of a pest: the evolution of a fruit-penetrating ovipositor in <i>Drosophila suzukii </i> nd related species. Proceedings of the Royal Society B: Biological Sciences, 2014, 281, 20132840.	2.6	277
11	Phylogenomics Resolves Evolutionary Relationships among Ants, Bees, and Wasps. Current Biology, 2013, 23, 2058-2062.	3.9	143
12	The Utility of Shallow RNA-Seq for Documenting Differential Gene Expression in Genes with High and Low Levels of Expression. PLoS ONE, 2013, 8, e84160.	2.5	11
13	Many ways to make a novel structure: a new mode of sex comb development in Drosophilidae. Evolution & Development, 2012, 14, 476-483.	2.0	18
14	Cell dynamics and developmental bias in the ontogeny of a complex sexually dimorphic trait in <i>Drosophila melanogaster</i> . Evolution & Development, 2009, 11, 191-204.	2.0	20
15	Developmental constraints and convergent evolution in <i>Drosophila </i> sex comb formation. Evolution & Development, 2009, 11, 205-218.	2.0	24
16	Chapter 3 Genotype–Phenotype Mapping. International Review of Cell and Molecular Biology, 2009, 278, 119-148.	3.2	7
17	The environmental and genetic regulation of obake expressivity: morphogenetic fields as evolvable systems. Evolution & Development, 2004, 6, 114-122.	2.0	15