## Sara Guirao-Rico

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/2344971/publications.pdf

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20 papers

4,875 citations

840585 11 h-index 752573 20 g-index

23 all docs 23 docs citations

 $\begin{array}{c} 23 \\ times \ ranked \end{array}$ 

7587 citing authors

#	Article	IF	CITATIONS
1	DnaSP 6: DNA Sequence Polymorphism Analysis of Large Data Sets. Molecular Biology and Evolution, 2017, 34, 3299-3302.	3.5	4,056
2	Genome sequences of the human body louse and its primary endosymbiont provide insights into the permanent parasitic lifestyle. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 12168-12173.	3.3	482
3	Population-scale long-read sequencing uncovers transposable elements associated with gene expression variation and adaptive signatures in Drosophila. Nature Communications, 2022, 13, 1948.	5.8	53
4	Populationâ€specific dynamics and selection patterns of transposable element insertions in European natural populations. Molecular Ecology, 2019, 28, 1506-1522.	2.0	45
5	<i>Drosophila</i> Evolution over Space and Time (DEST): A New Population Genomics Resource. Molecular Biology and Evolution, 2021, 38, 5782-5805.	3.5	37
6	<scp>RAD</scp> mapping reveals an evolving, polymorphic and fuzzy boundary of a plant pseudoautosomal region. Molecular Ecology, 2016, 25, 414-430.	2.0	29
7	Recombination changes at the boundaries of fully and partially sex-linked regions between closely related Silene species pairs. Heredity, 2017, 118, 395-403.	1.2	19
8	Benchmarking the performance of Poolâ€seq SNP callers using simulated and real sequencing data. Molecular Ecology Resources, 2021, 21, 1216-1229.	2.2	19
9	Molecular Population Genetics of the Insulin/TOR Signal Transduction Pathway: A Network-Level Analysis in Drosophila melanogaster. Molecular Biology and Evolution, 2012, 29, 123-132.	3 <b>.</b> 5	17
10	Sequence diversity patterns suggesting balancing selection in partially sexâ€linked genes of the plant <i>Silene latifolia</i> are not generated by demographic history or gene flow. Molecular Ecology, 2017, 26, 1357-1370.	2.0	17
11	Temperature, rainfall and wind variables underlie environmental adaptation in natural populations of <i>Drosophila melanogaster</i> Molecular Ecology, 2021, 30, 938-954.	2.0	15
12	Porcine Y-chromosome variation is consistent with the occurrence of paternal gene flow from non-Asian to Asian populations. Heredity, 2018, 120, 63-76.	1.2	14
13	Comparative analysis of tissue-specific transcriptomes in the funnel-web spider (i) Macrothele calpeiana (i) (Araneae, Hexathelidae). PeerJ, 2015, 3, e1064.	0.9	14
14	Positive Selection Has Driven the Evolution of the Drosophila Insulin-Like Receptor (InR) at Different Timescales. Molecular Biology and Evolution, 2009, 26, 1723-1732.	3.5	11
15	Evolutionary History of Major Chemosensory Gene Families across Panarthropoda. Molecular Biology and Evolution, 2020, 37, 3601-3615.	3.5	10
16	DOMINO: development of informative molecular markers for phylogenetic and genome-wide population genetic studies in non-model organisms. Bioinformatics, 2016, 32, 3753-3759.	1,8	8
17	Evolutionary insights from large scale resequencing datasets in Drosophila melanogaster. Current Opinion in Insect Science, 2019, 31, 70-76.	2.2	8
18	Molecular Evolution of the Ligands of the Insulin-Signaling Pathway: dilp Genes in the Genus Drosophila. Molecular Biology and Evolution, 2011, 28, 1557-1560.	3 <b>.</b> 5	6

#	Article	lF	CITATIONS
19	Computational prediction of the phenotypic effects of genetic variants: basic concepts and some application examples in Drosophila nervous system genes. Journal of Neurogenetics, 2017, 31, 307-319.	0.6	2
20	Patterns of Nucleotide Diversity at the Regions Encompassing the Drosophila Insulin-Like Peptide (dilp) Genes: Demography vs. Positive Selection in Drosophila melanogaster. PLoS ONE, 2013, 8, e53593.	1.1	1