

Armando Moreno Geraldés

List of Publications by Year in descending order

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Version: 2024-02-01

26
papers

2,251
citations

430442

18
h-index

610482

24
g-index

28
all docs

28
docs citations

28
times ranked

3045
citing authors

#	ARTICLE	IF	CITATIONS
1	The tip of the iceberg: Genome wide marker analysis reveals hidden hybridization during invasion. <i>Molecular Ecology</i> , 2021, 30, 810-825.	2.0	3
2	Editorial 2020. <i>Molecular Ecology</i> , 2020, 29, 1-19.	2.0	3
3	Population genomic analyses reveal a highly differentiated and endangered genetic cluster of northern goshawks (<i>Accipiter gentilis laingi</i>) in Haida Gwaii. <i>Evolutionary Applications</i> , 2019, 12, 757-772.	1.5	14
4	Some perspective on <i>Molecular Ecology</i> perspectives: Are women being left out?. <i>Molecular Ecology</i> , 2019, 28, 2451-2455.	2.0	5
5	Editorial 2017. <i>Molecular Ecology</i> , 2017, 26, 383-412.	2.0	2
6	Editorial 2016. <i>Molecular Ecology</i> , 2016, 25, 433-449.	2.0	0
7	Recent Y chromosome divergence despite ancient origin of dioecy in poplars (<i>Populus</i>). <i>Molecular Ecology</i> , 2015, 24, 3243-3256.	2.0	121
8	LANDSCAPE GENOMICS OF <i>POPULUS TRICHOCARPA</i> : THE ROLE OF HYBRIDIZATION, LIMITED GENE FLOW, AND NATURAL SELECTION IN SHAPING PATTERNS OF POPULATION STRUCTURE. <i>Evolution; International Journal of Organic Evolution</i> , 2014, 68, 3260-3280.	1.1	88
9	Testing a "genes-to-ecosystems" approach to understanding aquatic-terrestrial linkages. <i>Molecular Ecology</i> , 2014, 23, 5888-5903.	2.0	28
10	Geographical and environmental gradients shape phenotypic trait variation and genetic structure in <i>Populus trichocarpa</i> . <i>New Phytologist</i> , 2014, 201, 1263-1276.	3.5	185
11	Genome-wide association implicates numerous genes underlying ecological trait variation in natural populations of <i>Populus trichocarpa</i> . <i>New Phytologist</i> , 2014, 203, 535-553.	3.5	171
12	A 34K SNP genotyping array for <i>Populus trichocarpa</i> : Design, application to the study of natural populations and transferability to other <i>Populus</i> species. <i>Molecular Ecology Resources</i> , 2013, 13, 306-323.	2.2	92
13	<i>Populus trichocarpa</i> cell wall chemistry and ultrastructure trait variation, genetic control and genetic correlations. <i>New Phytologist</i> , 2013, 197, 777-790.	3.5	100
14	The genome of flax (<i>Linum usitatissimum</i>) assembled <i>de novo</i> from short shotgun sequence reads. <i>Plant Journal</i> , 2012, 72, 461-473.	2.8	415
15	Genome resequencing reveals multiscale geographic structure and extensive linkage disequilibrium in the forest tree <i>Populus trichocarpa</i> . <i>New Phytologist</i> , 2012, 196, 713-725.	3.5	173
16	The Genetic Structure of Domestic Rabbits. <i>Molecular Biology and Evolution</i> , 2011, 28, 1801-1816.	3.5	101
17	SNP discovery in black cottonwood (<i>Populus trichocarpa</i>) by population transcriptome resequencing. <i>Molecular Ecology Resources</i> , 2011, 11, 81-92.	2.2	104
18	Higher differentiation among subspecies of the house mouse (<i>Mus musculus</i>) in genomic regions with low recombination. <i>Molecular Ecology</i> , 2011, 20, 4722-4736.	2.0	124

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19	Pushing north one bottleneck at a time: site frequency spectra tell the history of Sitka spruce. <i>Molecular Ecology</i> , 2010, 19, 3837-3839.	2.0	0
20	Extensive Gene Conversion Drives the Concerted Evolution of Paralogous Copies of the SRY Gene in European Rabbits. <i>Molecular Biology and Evolution</i> , 2010, 27, 2437-2440.	3.5	26
21	Reduced introgression of the Y chromosome between subspecies of the European rabbit (<i>Oryctolagus cuniculus</i>) in the Iberian Peninsula. <i>Molecular Ecology</i> , 2008, 17, 4489-4499.	2.0	45
22	Inferring the history of speciation in house mice from autosomal, X-linked, Y-linked and mitochondrial genes. <i>Molecular Ecology</i> , 2008, 17, 5349-5363.	2.0	219
23	Nucleotide Variation in Wild and Inbred Mice. <i>Genetics</i> , 2007, 177, 2277-2291.	1.2	90
24	Contrasting Patterns of Introgression at X-Linked Loci Across the Hybrid Zone Between Subspecies of the European Rabbit (<i>Oryctolagus cuniculus</i>). <i>Genetics</i> , 2006, 173, 919-933.	1.2	89
25	A 7-bp insertion in the 3' untranslated region suggests the duplication and concerted evolution of the rabbit SRY gene. <i>Genetics Selection Evolution</i> , 2006, 38, 313.	1.2	17
26	High levels of nucleotide diversity in the European rabbit (<i>Oryctolagus cuniculus</i>) SRY gene. <i>Animal Genetics</i> , 2005, 36, 349-351.	0.6	34