Horacio F Naveira

List of Publications by Year
in descending order

40 papers	2,269 citations	471509 17 h-index	289244 40 g-index
40 all docs	40 docs citations	40 times ranked	2663 citing authors

#	Article	IF	Citations
1	Evidence of doubly uniparental inheritance of the mitochondrial<scp>DNA</scp>inPolititapes rhomboides(Bivalvia, Veneridae): Evolutionary and population genetic analysis of F and M mitotypes. Journal of Zoological Systematics and Evolutionary Research, 2020, 58, 541-560.	1.4	7
2	Fully automatic multi-temporal land cover classification using Sentinel-2 image data. Procedia Computer Science, 2019, 159, 650-657.	2.0	16
3	Census and contemporary effective population size of two populations of the protected Spanish Moon Moth (Graellsia isabellae). Insect Conservation and Diversity, 2019, 12, 147-160.	3.0	2
4	Phylogeography of the Spanish Moon Moth Graellsia isabellae (Lepidoptera, Saturniidae). BMC Evolutionary Biology, 2016, 16, 139.	3.2	11
5	Evolutionary dynamics of two satellite DNA families in rock lizards of the genus Iberolacerta (Squamata, Lacertidae): different histories but common traits. Chromosome Research, 2015, 23, 441-461.	2.2	20
6	Isolation and characterization of two satellite DNAs in some Iberian rock lizards (Squamata,) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 547 T 322, 13-26.	1.3	11
7	Karyological Characterization of the Endemic Iberian Rock Lizard, <i>Iberolacerta monticola</i> (Squamata, Lacertidae): Insights into Sex Chromosome Evolution. Cytogenetic and Genome Research, 2014, 142, 28-39.	1.1	16
8	Evidence for the persistence of an active endogenous retrovirus (ERVE) in humans. Genetica, 2014, 142, 451-460.	1.1	10
9	A new mountain lizard from Montes de LeÃ³n (NW Iberian Peninsula):ÂIberolacerta monticola astur ssp. nov. (Squamata: Lacertidae) . Zootaxa, 2014, 3796, 201.	0.5	7
10	Causes and Evolutionary Consequences of Population Subdivision of an Iberian Mountain Lizard, Iberolacerta monticola. PLoS ONE, 2013, 8, e66034.	2.5	11
11	Chronicle of an extinction foretold: genetic properties of an extremely small population of Iberolacerta monticola. Conservation Genetics, 2012, 13, 131-142.	1.5	6
12	Evolutionary Dynamics of the Ty3/Gypsy LTR Retrotransposons in the Genome of Anopheles gambiae. PLoS ONE, 2011, 6, e16328.	2.5	15
13	Sequencing of Culex quinquefasciatus Establishes a Platform for Mosquito Comparative Genomics. Science, 2010, 330, 86-88.	12.6	424
14	Isolation and characterization of polymorphic microsatellite markers in Iberolacerta monticola, and crossâ€species amplification in Iberolacerta galani and Zootoca vivipara. Molecular Ecology Resources, 2008, 8, 1351-1353.	4.8	8
15	Genome Sequence of Aedes aegypti, a Major Arbovirus Vector. Science, 2007, 316, 1718-1723.	12.6	1,025
16	Structural and Evolutionary Analyses of the Ty3/gypsy Group of LTR Retrotransposons in the Genome of Anopheles gambiae. Molecular Biology and Evolution, 2005, 22, 29-39.	8.9	26
17	Paleogenomic Record of the Extinction of Human Endogenous Retrovirus ERV9. Journal of Virology, 2005, 79, 6997-7004.	3.4	15
18	Evolution of the mdg1 lineage of the Ty3/gypsy group of LTR retrotransposons in Anopheles gambiae. Gene, 2004, 330, 123-131.	2.2	9

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19	On the relative roles of faster-X evolution and dominance in the establishment of intrinsic postzygotic isolating barriers. Genetica, 2003, 118, 41-50.	1.1	9
20	Amplification and Phylogenetic Relationships of a Subfamily of blood, a Retrotransposable Element of Drosophila. Journal of Molecular Evolution, 2001, 52, 342-350.	1.8	11
21	Structural Features of the mdg1 Lineage of the Ty3/gypsy Group of LTR Retrotransposons Inferred from the Phylogenetic Analyses of Its Open Reading Frames. Journal of Molecular Evolution, 2001, 53, 165-171.	1.8	8
22	Evolutionary History of the Human Endogenous Retrovirus Family ERV9. Molecular Biology and Evolution, 2000, 17, 320-330.	8.9	44
23	Hidden Effects of X Chromosome Introgressions on Spermatogenesis in Drosophila simulans Ã— D. mauritiana Hybrids Unveiled by Interactions Among Minor Genetic Factors. Genetics, 1998, 150, 745-754.	2.9	26
24	A three-locus system of interspecific incompatibility underlies male inviability in hybrids between Drosophila buzzatii and D. koepferae. Genetica, 1996, 98, 1-19.	1.1	32
25	On the difficulties of discriminating between major and minor hybrid male sterility factors in Drosophila by examining the segregation ratio of sterile and fertile sons in backcrossing experiments. Heredity, 1996, 77, 433-438.	2.6	6
26	A polygenic basis of hybrid sterility may give rise to spurious localizations of major sterility factors. Heredity, 1996, 77, 488-492.	2.6	15
27	The evolutionary history of Drosophila buzzatti. XXVI. Macrogeographic patterns of inversion polymorphism in New World populations. Journal of Evolutionary Biology, 1995, 8, 369-384.	1.7	63
28	The estimation of genotypic probabilities in an adult population by the analysis of descendants. Genetical Research, 1992, 59, 131-137.	0.9	3
29	Location of X-linked polygenic effects causing sterility in male hybrids of Drosophila simulans and D. mauritiana. Heredity, 1992, 68, 211-217.	2.6	46
30	The evolutionary history of D. buzzatii. XXII. Chromosomal and genic sterility in male hybrids of Drosophila buzzatii and Drosophila koepferae. Heredity, 1991, 66, 233-239.	2.6	26
31	The evolutionary history of Drosophila buzzatii. XXI. Cumulative action of multiple sterility factors on spermatogenesis in hybrids of D. buzzatii and D. koepferae. Heredity, 1991, 67, 57-72.	2.6	33
32	The evolutionary history of Drosophila buzzatii. XVI. Fitness component analysis in an original natural population from Argentina. Journal of Evolutionary Biology, 1991, 4, 209-225.	1.7	38
33	Genetic Mapping of the Adh Locus in the Repleta Group of Drosophila by in situ Hybridization. Journal of Heredity, 1990, 81, 83-86.	2.4	28
34	Drosophila koepferae: a New Member of the Drosophila serido (Diptera: Drosophilidae) Superspecies Taxon1. Annals of the Entomological Society of America, 1988, 81, 380-385.	2.5	63
35	The evolutionary history of Drosophila buzzatii XI. A new method for cytogenetic localization based on asynapsis of polytene chromosomes in interspecific hybrids of Drosophila. Genetica, 1986, 71, 199-212.	1.1	18
36	THE EVOLUTIONARY HISTORY OF DROSOPHILA BUZZATII. XII. THE GENETIC BASIS OF STERILITY IN HYBRIDS BETWEEN D. BUZZATII AND ITS SIBLING D. SERIDO FROM ARGENTINA. Genetics, 1986, 114, 841-857.	2.9	76

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37	The evolutionary history of Drosophila buzzatii. Chromosoma, 1985, 91, 87-94.	2.2	71
38	Chromosomal localization of the locus PGM (phosphoglucomutase) inDrosophila buzzatii. Experientia, 1985, 41, 507-508.	1.2	2
39	The evolutionary history ofDrosophila buzzatii. V. Differential survivorship onOpuntia betweenD. buzzatii andD. serido. Experientia, 1985, 41, 129-131.	1.2	3
40	Spermiogenesis of inversion heterozygotes in backcross hybrids between Drosophila buzzatii and D. serido. Genetica, 1984, 65, 205-214.	1.1	9