Jose Castresana

List of Publications by Year in descending order

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109137 110170 17,797 65 35 64 citations h-index g-index papers 67 67 67 23239 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Divergence time estimation using ddRAD data and an isolation-with-migration model applied to water vole populations of Arvicola. Scientific Reports, 2022, 12, 4065.	1.6	9
2	The genome of the Pyrenean desman and the effects of bottlenecks and inbreeding on the genomic landscape of an endangered species. Evolutionary Applications, 2021, 14, 1898-1913.	1.5	11
3	Size increase without genetic divergence in the Eurasian water shrew Neomys fodiens. Scientific Reports, 2019, 9, 17375.	1.6	3
4	Quantitative analysis of connectivity in populations of a semiâ€aquatic mammal using kinship categories and network assortativity. Molecular Ecology Resources, 2019, 19, 310-326.	2.2	29
5	Evolutionary history of the endemic water shrew <i>Neomys anomalus</i> : Recurrent phylogeographic patterns in semiâ€aquatic mammals of the Iberian Peninsula. Ecology and Evolution, 2018, 8, 10138-10146.	0.8	9
6	A pipeline for metabarcoding and diet analysis from fecal samples developed for a small semi-aquatic mammal. PLoS ONE, 2018, 13, e0201763.	1.1	22
7	Postglacial dispersal patterns and mitochondrial genetic structure of the Pyrenean desman (<i>Galemys pyrenaicus</i>) in the northwestern region of the Iberian Peninsula. Ecology and Evolution, 2017, 7, 4486-4495.	0.8	9
8	Spatial mixing of mitochondrial lineages and greater genetic diversity in some invasive populations of the American mink (Neovison vison) compared to native populations. Biological Invasions, 2017, 19, 2663-2673.	1.2	10
9	Using relatedness networks to infer contemporary dispersal: Application to the endangered mammal <i>Galemys pyrenaicus (i). Molecular Ecology, 2017, 26, 3343-3357.</i>	2.0	30
10	Genomic diversity and geographical structure of the Pyrenean desman. Conservation Genetics, 2016, 17, 1333-1344.	0.8	15
11	Multilocus species trees and species delimitation in a temporal context: application to the water shrews of the genus Neomys. BMC Evolutionary Biology, 2015, 15, 209.	3.2	21
12	Development of Rapidly Evolving Intron Markers to Estimate Multilocus Species Trees of Rodents. PLoS ONE, 2014, 9, e96032.	1.1	8
13	Phylogeography and postglacial expansion of the endangered semi-aquatic mammal Galemys pyrenaicus. BMC Evolutionary Biology, 2013, 13, 115.	3.2	33
14	Diversification rates and the latitudinal gradient of diversity in mammals. Proceedings of the Royal Society B: Biological Sciences, 2012, 279, 4148-4155.	1.2	53
15	Impact of Deep Coalescence on the Reliability of Species Tree Inference from Different Types of DNA Markers in Mammals. PLoS ONE, 2012, 7, e30239.	1.1	35
16	Patterns of mammalian diversification in recent evolutionary times: global tendencies and methodological issues. Journal of Evolutionary Biology, 2011, 24, 2611-2623.	0.8	8
17	Novel intron markers to study the phylogeny of closely related mammalian species. BMC Evolutionary Biology, 2010, 10, 369.	3.2	43

Distribution and evolutionary trends of photoprotective isoprenoids (xanthophylls and) Tj ETQq0.00 rgBT /Overlock 10.00 Tf 10.00 Td (too

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19	Paleogenomics in a Temperate Environment: Shotgun Sequencing from an Extinct Mediterranean Caprine. PLoS ONE, 2009, 4, e5670.	1.1	30
20	Estimation of Phylogenetic Inconsistencies in the Three Domains of Life. Molecular Biology and Evolution, 2008, 25, 2319-2329.	3.5	18
21	Accelerated Evolution of Genes of Recent Origin. , 2008, , 45-59.		4
22	The K tree score: quantification of differences in the relative branch length and topology of phylogenetic trees. Bioinformatics, 2007, 23, 2954-2956.	1.8	93
23	Improvement of Phylogenies after Removing Divergent and Ambiguously Aligned Blocks from Protein Sequence Alignments. Systematic Biology, 2007, 56, 564-577.	2.7	4,438
24	Topological variation in single-gene phylogenetic trees. Genome Biology, 2007, 8, 216.	3.8	71
25	On homology searches by protein Blast and the characterization of the age of genes. BMC Evolutionary Biology, 2007, 7, 53.	3.2	86
26	Phylogenetic position of Salinibacter ruber based on concatenated protein alignments. Systematic and Applied Microbiology, 2007, 30, 171-179.	1.2	29
27	Is mammalian chromosomal evolution driven by regions of genome fragility?. Genome Biology, 2006, 7, R115.	13.9	130
28	Molecular dating of caprines using ancient DNA sequences of Myotragus balearicus, an extinct endemic Balearic mammal. BMC Evolutionary Biology, 2005, 5, 70.	3.2	66
29	Inverse Relationship Between Evolutionary Rate and Age of Mammalian Genes. Molecular Biology and Evolution, 2005, 22, 598-606.	3.5	152
30	Phylogenetic and Ecological Analysis of Novel Marine Stramenopiles. Applied and Environmental Microbiology, 2004, 70, 3528-3534.	1.4	321
31	Clustering of Genes Coding for DNA Binding Proteins in a Regionof Atypical Evolution of the Human Genome. Journal of Molecular Evolution, 2004, 59, 72-9.	0.8	17
32	The cDNA for leucomyosuppressin in Blattella germanica and molecular evolution of insect myosuppressins. Peptides, 2004, 25, 1883-1889.	1.2	16
33	Orcokinins in insects and other invertebrates. Insect Biochemistry and Molecular Biology, 2004, 34, 1141-1146.	1.2	48
34	Genes on human chromosome 19 show extreme divergence from the mouse orthologs and a high GC content. Nucleic Acids Research, 2002, 30, 1751-1756.	6. 5	50
35	Estimation of genetic distances from human and mouse introns. Genome Biology, 2002, 3, research0028.1.	13.9	23
36	Comparative genomics and bioenergetics. Biochimica Et Biophysica Acta - Bioenergetics, 2001, 1506, 147-162.	0.5	51

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37	Cytochrome b Phylogeny and the Taxonomy of Great Apes and Mammals. Molecular Biology and Evolution, 2001, 18, 465-471.	3.5	99
38	A Molecular Phylogeny of Two Extinct Sloths. Molecular Phylogenetics and Evolution, 2001, 18, 94-103.	1.2	63
39	Selection of Conserved Blocks from Multiple Alignments for Their Use in Phylogenetic Analysis. Molecular Biology and Evolution, 2000, 17, 540-552.	3.5	8,843
40	Nitric oxide reductases in bacteria. Biochimica Et Biophysica Acta - Bioenergetics, 2000, 1459, 266-273.	0.5	172
41	Respiratory Chains in the Last Common Ancestor of Living Organisms. Journal of Molecular Evolution, 1999, 49, 453-460.	0.8	66
42	A hominoid-specific nuclear insertion of the mitochondrial D-loop: implications for reconstructing ancestral mitochondrial sequences. Molecular Biology and Evolution, 1998, 15, 463-469.	3.5	40
43	Codon reassignment and amino acid composition in hemichordate mitochondria. Proceedings of the National Academy of Sciences of the United States of America, 1998, 95, 3703-3707.	3.3	80
44	The Mitochondrial Genome of the Hemichordate Balanoglossus carnosus and the Evolution of Deuterostome Mitochondria. Genetics, 1998, 150, 1115-1123.	1.2	90
45	Evolution of the spectrin repeat. BioEssays, 1997, 19, 811-817.	1.2	69
46	Evolution of energetic metabolism: the respiration-early hypothesis. Trends in Biochemical Sciences, 1995, 20, 443-448.	3.7	195
47	New Archaebacterial Genes Coding for Redox Proteins: Implications for the Evolution of Aerobic Metabolism. Journal of Molecular Biology, 1995, 250, 202-210.	2.0	108
48	Does Vav bind to F-actin through a CH domain?. FEBS Letters, 1995, 374, 149-151.	1.3	135
49	Evolution of cytochrome oxidase, an enzyme older than atmospheric oxygen EMBO Journal, 1994, 13, 2516-2525.	3.5	232
50	A Second Terminal Oxidase in Sulfolobus acidocaldarius. FEBS Journal, 1994, 224, 151-159.	0.2	95
51	Cytochrome oxidase evolved by tinkering with denitrification enzymes. FEBS Letters, 1994, 341, 1-4.	1.3	230
52	A Role for Cholesterol as a Structural Effector of the Nicotinic Acetylcholine Receptor. Biochemistry, 1994, 33, 4065-4071.	1.2	72
53	Structure and thermal denaturation of crystalline and noncrystalline cytochrome oxidase as studied by infrared spectroscopy. Biochemistry, 1994, 33, 11650-11655.	1.2	132
54	Role of cholesterol as a structural and functional effector of the nicotinic acetylcholine receptor. Biochemical Society Transactions, 1994, 22, 776-780.	1.6	13

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55	Quantitative studies of the structure of proteins in solution by fourier-transform infrared spectroscopy. Progress in Biophysics and Molecular Biology, 1993, 59, 23-56.	1.4	764
56	Partial dehydration of phosphatidylethanolamine phosphate groups during hexagonal phase formation, as seen by i.r. spectroscopy. Biochemical Journal, 1992, 282, 467-470.	1.7	40
57	Protein stability and interaction of the nicotinic acetylcholine receptor with cholinergic ligands studied by Fourier-transform infrared spectroscopy. Biochemical Journal, 1992, 288, 421-426.	1.7	44
58	Protein structural effects of agonist binding to the nicotinic acetylcholine receptor. FEBS Letters, 1992, 314, 171-175.	1.3	27
59	The physical state of ubiquinone-10, in pure form and incorporated into phospholipid bilayers. A Fourier-transform infrared spectroscopic study. FEBS Journal, 1992, 204, 1125-1130.	0.2	22
60	An infrared spectroscopic study of specifically deuterated fatty-acyl methyl groups in phosphatidylcholine liposomes. Biochimica Et Biophysica Acta - Biomembranes, 1991, 1065, 29-34.	1.4	14
61	Solubilization of sarcoplasmic reticulum membranes by sodium dodecylsulphate. FEBS Letters, 1990, 269, 324-327.	1.3	5
62	The lamellar to hexagonal phase transition in phosphatidylethanolamine liposomes: A fluorescence anisotropy study. Biochemical and Biophysical Research Communications, 1990, 168, 987-992.	1.0	9
63	An infrared spectroscopic study of \hat{l}^2 -galactosidase structure in aqueous solutions. FEBS Letters, 1989, 252, 118-120.	1.3	52
64	Interaction of SDS with \hat{I}^2 -galactosidase. A FT-IR study of the influence of detergent concentration and temperature Journal of Molecular Structure, 1988, 175, 67-72.	1.8	3
65	The structure of proteins in aqueous solutions: An assessment of triose phosphate isomerase structure by fourier-transform infrared spectroscopy. Biochemical and Biophysical Research Communications, 1988, 152, 69-75.	1.0	34