Luis Sanchez-Pulido

List of Publications by Year in descending order

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61 5,178 33 59 g-index
61 61 61 9340

times ranked

citing authors

docs citations

all docs

#	Article	IF	CITATIONS
1	Citrullination Was Introduced into Animals by Horizontal Gene Transfer from Cyanobacteria. Molecular Biology and Evolution, 2022, 39, .	3.5	16
2	Refining the domain architecture model of the replication origin firing factor Treslin/TICRR. Life Science Alliance, 2022, 5, e202101088.	1.3	7
3	<i>PERCC1</i> , a new member of the <i>Yap/TAZ</i> /i>/ <i>FAM181</i> /i>transcriptional co-regulator family. Bioinformatics Advances, 2022, 2, .	0.9	2
4	The Role of MTBP as a Replication Origin Firing Factor. Biology, 2022, 11, 827.	1.3	3
5	Extending the Horizon of Homology Detection with Coevolution-based Structure Prediction. Journal of Molecular Biology, 2021, 433, 167106.	2.0	7
6	Hexa-Longin domain scaffolds for inter-Rab signalling. Bioinformatics, 2020, 36, 990-993.	1.8	5
7	Fam151b, the mouse homologue of C.elegans menorin gene, is essential for retinal function. Scientific Reports, 2020, 10, 437.	1.6	2
8	Missense Mutations in the Human Nanophthalmos Gene <i>TMEM98</i> Cause Retinal Defects in the Mouse., 2019, 60, 2875.		16
9	The Cdk8/19-cyclin C transcription regulator functions in genome replication through metazoan Sld7. PLoS Biology, 2019, 17, e2006767.	2.6	32
10	TMEM132: an ancient architecture of cohesin and immunoglobulin domains define a new family of neural adhesion molecules. Bioinformatics, 2018, 34, 721-724.	1.8	38
11	The phenotypic spectrum of Xiaâ€Gibbs syndrome. American Journal of Medical Genetics, Part A, 2018, 176, 1315-1326.	0.7	34
12	APC/CCdh1 Enables Removal of Shugoshin-2 from the Arms of Bivalent Chromosomes by Moderating Cyclin-Dependent Kinase Activity. Current Biology, 2017, 27, 1462-1476.e5.	1.8	8
13	Mutations in CDC45, Encoding an Essential Component of the Pre-initiation Complex, Cause Meier-Gorlin Syndrome and Craniosynostosis. American Journal of Human Genetics, 2016, 99, 125-138.	2.6	92
14	The role of ADP-ribosylation in regulating DNA interstrand crosslink repair. Journal of Cell Science, 2016, 129, 3845-3858.	1.2	15
15	High incidence of unrecognized visceral/neurological late-onset Niemann-Pick disease, type C1, predicted by analysis of massively parallel sequencing data sets. Genetics in Medicine, 2016, 18, 41-48.	1.1	171
16	The Evolutionarily Conserved Tre2/Bub2/Cdc16 (TBC), Lysin Motif (LysM), Domain Catalytic (TLDc) Domain Is Neuroprotective against Oxidative Stress. Journal of Biological Chemistry, 2016, 291, 2751-2763.	1.6	48
17	RBFOX and PTBP1 proteins regulate the alternative splicing of micro-exons in human brain transcripts. Genome Research, 2015, 25, 1-13.	2.4	187
18	REC-1 and HIM-5 distribute meiotic crossovers and function redundantly in meiotic double-strand break formation in <i>Caenorhabditis elegans</i>	2.7	19

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19	TM6SF2 and MAC30, new enzyme homologs in sterol metabolism and common metabolic disease. Frontiers in Genetics, 2014, 5, 439.	1.1	50
20	The genomic substrate for adaptive radiation in African cichlid fish. Nature, 2014, 513, 375-381.	13.7	874
21	A Code for RanGDP Binding in Ankyrin Repeats Defines a Nuclear Import Pathway. Cell, 2014, 157, 1130-1145.	13.5	67
22	Monoallelic and Biallelic Mutations in MAB21L2 Cause a Spectrum of Major Eye Malformations. American Journal of Human Genetics, 2014, 94, 915-923.	2.6	79
23	The RFTS Domain of Raf2 Is Required for Cul4 Interaction and Heterochromatin Integrity in Fission Yeast. PLoS ONE, 2014, 9, e104161.	1.1	5
24	Ki-67 is a PP1-interacting protein that organises the mitotic chromosome periphery. ELife, 2014, 3, e01641.	2.8	167
25	Homozygous mutations in a predicted endonuclease are a novel cause of congenital dyserythropoietic anemia type I. Haematologica, 2013, 98, 1383-1387.	1.7	71
26	The microcephaly protein Asp regulates neuroepithelium morphogenesis byÂcontrolling theÂspatial distribution of myosin II. Nature Cell Biology, 2013, 15, 1294-1306.	4.6	114
27	MISP is a novel Plk1 substrate required for proper spindle orientation and mitotic progression. Journal of Cell Biology, 2013, 200, 773-787.	2.3	65
28	Family with Sequence Similarity 60A (FAM60A) Protein Is a Cell Cycle-fluctuating Regulator of the SIN3-HDAC1 Histone Deacetylase Complex. Journal of Biological Chemistry, 2012, 287, 32346-32353.	1.6	45
29	Nprl3 is required for normal development of the cardiovascular system. Mammalian Genome, 2012, 23, 404-415.	1.0	38
30	Repo-Man Coordinates Chromosomal Reorganization with Nuclear Envelope Reassembly during Mitotic Exit. Developmental Cell, 2011, 21, 328-342.	3.1	172
31	Regulation of DNA Replication through Sld3-Dpb11 Interaction Is Conserved from Yeast to Humans. Current Biology, 2011, 21, 1152-1157.	1.8	135
32	Structure and evolutionary history of DISC1. Human Molecular Genetics, 2011, 20, R175-R181.	1.4	13
33	Homology explains the functional similarities of Treslin/Ticrr and Sld3. Current Biology, 2010, 20, R509-R510.	1.8	69
34	Drosophila SMN complex proteins Gemin2, Gemin3, and Gemin5 are components of U bodies. Experimental Cell Research, 2010, 316, 2354-2364.	1.2	36
35	Loss of fish actinotrichia proteins and the fin-to-limb transition. Nature, 2010, 466, 234-237.	13.7	144
36	Identification of the MMS22L-TONSL Complex that Promotes Homologous Recombination. Molecular Cell, 2010, 40, 632-644.	4.5	100

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37	Stc1: A Critical Link between RNAi and Chromatin Modification Required for Heterochromatin Integrity. Cell, 2010, 140, 666-677.	13.5	195
38	The Protein Composition of Mitotic Chromosomes Determined Using Multiclassifier Combinatorial Proteomics. Cell, 2010, 142, 810-821.	13.5	266
39	Detection of Alpha-Rod Protein Repeats Using a Neural Network and Application to Huntingtin. PLoS Computational Biology, 2009, 5, e1000304.	1.5	59
40	Mitochondrial vesicles: an ancient process providing new links to peroxisomes. Current Opinion in Cell Biology, 2009, 21, 560-567.	2.6	64
41	Common Ancestry of the CENP-A Chaperones Scm3 and HJURP. Cell, 2009, 137, 1173-1174.	13.5	136
42	Coordination of Structure-Specific Nucleases by Human SLX4/BTBD12 Is Required for DNA Repair. Molecular Cell, 2009, 35, 116-127.	4.5	300
43	Homologues of Arabidopsis Microtubule-Associated AIR9 in Trypanosomatid Parasites. Plant Signaling and Behavior, 2007, 2, 296-299.	1.2	9
44	Are promyelocytic leukaemia protein nuclear bodies a scaffold for caspase-2 programmed cell death?. Trends in Biochemical Sciences, 2007, 32, 400-406.	3.7	20
45	Microtubule-Associated AIR9 Recognizes the Cortical Division Site at Preprophase and Cell-Plate Insertion. Current Biology, 2006, 16, 1938-1943.	1.8	118
46	Gas1 Is Related to the Glial Cell-derived Neurotrophic Factor Family Receptors \hat{l}_{\pm} and Regulates Ret Signaling. Journal of Biological Chemistry, 2006, 281, 14330-14339.	1.6	55
47	Death inducer obliterator protein 1 in the context of DNA regulation. Sequence analyses of distant homologues point to a novel functional role. FEBS Journal, 2005, 272, 3505-3511.	2.2	34
48	The Shwachman-Bodian-Diamond Syndrome Protein Family Is Involved in RNA Metabolism. Journal of Biological Chemistry, 2005, 280, 19213-19220.	1.6	100
49	The variant E233G of theRAD51Dgene could be a low-penetrance allele in high-risk breast cancer families withoutBRCA1/2mutations. International Journal of Cancer, 2004, 110, 845-849.	2.3	26
50	POTRA: a conserved domain in the FtsQ family and a class of \hat{l}^2 -barrel outer membrane proteins. Trends in Biochemical Sciences, 2003, 28, 523-526.	3.7	195
51	No mutations in the XRCC 2 gene in BRCA 1/2-negative high-risk breast cancer families. International Journal of Cancer, 2003, 103, 136-137.	2.3	7
52	Molecular and structural analysis of the panallergen profilin B cell epitopes defined by monoclonal antibodies. International Immunology, 2002, 14, 993-1001.	1.8	26
53	BRICHOS: a conserved domain in proteins associated with dementia, respiratory distress and cancer. Trends in Biochemical Sciences, 2002, 27, 329-332.	3.7	192
54	MARVEL: a conserved domain involved in membrane apposition events. Trends in Biochemical Sciences, 2002, 27, 599-601.	3.7	199

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55	Improvement in affinity and HIV-1 neutralization by somatic mutation in the heavy chain first complementarity-determining region of antibodies triggered by HIV-1 infection. European Journal of Immunology, 2001, 31, 128-137.	1.6	5
56	Molecular analysis of HIV-1 gp120 antibody response using isotype IgM and IgG phage display libraries from a long-term non-progressor HIV-1-infected individual. European Journal of Immunology, 1999, 29, 2666-2675.	1.6	24
57	Shaping of Drosophila Alcohol Dehydrogenase Through Evolution: Relationship with Enzyme Functionality. Journal of Molecular Evolution, 1998, 47, 211-221.	0.8	23
58	Influenza virus epidemiological surveillance in Argentina, 1987-1993, with molecular characterization of 1990 and 1993 isolates. Revista Panamericana De Salud Publica/Pan American Journal of Public Health, 1998, 4, 405-10.	0.6	0
59	A Single Residue Substitution Causes a Switch from the Dual DNA Binding Specificity of Plant Transcription Factor MYB.Ph3 to the Animal c-MYB Specificity. Journal of Biological Chemistry, 1997, 272, 2889-2895.	1.6	44
60	Specific DNA recognition by the Aspergillus nidulans three zinc finger transcription factor PacC. Journal of Molecular Biology, 1997, 274, 466-480.	2.0	107
61	The \hat{I}^2 -tubulin monomer release factor (p14) has homology with a region of the DnaJ protein. FEBS Letters, 1996, 397, 283-289.	1.3	28