Jos Pereira-Leal

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

59	4,945	31	65
papers	citations	h-index	g-index
65 ext. papers	5,623 ext. citations	7.2 avg, IF	5.54 L-index

#	Paper	IF	Citations
59	A Pilot Study on the Metabolic Impact of Mediterranean Diet in Type 2 Diabetes: Is Gut Microbiota the Key?. <i>Nutrients</i> , 2021 , 13,	6.7	5
58	A novel cyanobacterial geosmin producer, revising GeoA distribution and dispersion patterns in Bacteria. <i>Scientific Reports</i> , 2020 , 10, 8679	4.9	11
57	Disulfiram, an alcohol dependence therapy, can inhibit the in vitro growth of Francisella tularensis. <i>International Journal of Antimicrobial Agents</i> , 2019 , 54, 85-88	14.3	6
56	Pericentrin-mediated SAS-6 recruitment promotes centriole assembly. <i>ELife</i> , 2019 , 8,	8.9	15
55	Diversity and Composition of Pelagic Prokaryotic and Protist Communities in a Thin Arctic Sea-Ice Regime. <i>Microbial Ecology</i> , 2019 , 78, 388-408	4.4	11
54	Genetic Competence Drives Genome Diversity in Bacillus subtilis. <i>Genome Biology and Evolution</i> , 2018 , 10, 108-124	3.9	29
53	Collective electrical oscillations of a diatom population induced by dark stress. <i>Scientific Reports</i> , 2018 , 8, 5484	4.9	6
52	Microbial Diversity and Toxin Risk in Tropical Freshwater Reservoirs of Cape Verde. <i>Toxins</i> , 2018 , 10,	4.9	5
51	Centrosome amplification arises before neoplasia and increases upon p53 loss in tumorigenesis. <i>Journal of Cell Biology</i> , 2018 , 217, 2353-2363	7.3	36
50	Rabifier2: an improved bioinformatic classifier of Rab GTPases. <i>Bioinformatics</i> , 2017 , 33, 568-570	7.2	3
49	Does Hypoxic Response Mediate Primary Resistance to Sunitinib in Untreated Locally Advanced Breast Cancer?. <i>Current Cancer Drug Targets</i> , 2017 , 17, 62-73	2.8	3
48	RAG Recombinase as a Selective Pressure for Genome Evolution. <i>Genome Biology and Evolution</i> , 2016 , 8, 3364-3376	3.9	4
47	Are There Rab GTPases in Archaea?. Molecular Biology and Evolution, 2016, 33, 1833-42	8.3	18
46	CYR61 and TAZ Upregulation and Focal Epithelial to Mesenchymal Transition May Be Early Predictors of Barrettu Esophagus Malignant Progression. <i>PLoS ONE</i> , 2016 , 11, e0161967	3.7	5
45	Evolutionary patterns in coiled-coils. <i>Genome Biology and Evolution</i> , 2015 , 7, 545-56	3.9	15
44	Coiled-coil length: Size does matter. <i>Proteins: Structure, Function and Bioinformatics</i> , 2015 , 83, 2162-9	4.2	9
43	Staphylococcus aureus Survives with a Minimal Peptidoglycan Synthesis Machine but Sacrifices Virulence and Antibiotic Resistance. <i>PLoS Pathogens</i> , 2015 , 11, e1004891	7.6	55

(2011-2015)

	42	Bioinformatic approaches to identifying and classifying Rab proteins. <i>Methods in Molecular Biology</i> , 2015 , 1298, 17-28	1.4	1	
	41	Evolutionary cell biology: two origins, one objective. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014 , 111, 16990-4	11.5	75	
,	40	A comprehensive assessment of the transcriptome of cork oak (Quercus suber) through EST sequencing. <i>BMC Genomics</i> , 2014 , 15, 371	4.5	31	
	39	Hope for GWAS: relevant risk genes uncovered from GWAS statistical noise. <i>International Journal of Molecular Sciences</i> , 2014 , 15, 17601-21	6.3	2	
	38	Bioinformatics projects supporting life-sciences learning in high schools. <i>PLoS Computational Biology</i> , 2014 , 10, e1003404	5	8	
	37	inTB - a data integration platform for molecular and clinical epidemiological analysis of tuberculosis. <i>BMC Bioinformatics</i> , 2013 , 14, 264	3.6	7	
	36	SNP typing reveals similarity in Mycobacterium tuberculosis genetic diversity between Portugal and Northeast Brazil. <i>Infection, Genetics and Evolution</i> , 2013 , 18, 238-46	4.5	17	
	35	Evolution of intracellular compartmentalization. <i>Biochemical Journal</i> , 2013 , 449, 319-31	3.8	98	
	34	Genome of a Gut Strain of Bacillus subtilis. <i>Genome Announcements</i> , 2013 , 1,		15	
	33	A genomic signature and the identification of new sporulation genes. <i>Journal of Bacteriology</i> , 2013 , 195, 2101-15	3.5	83	
	32	Aurora at the pole and equator: overlapping functions of Aurora kinases in the mitotic spindle. <i>Open Biology</i> , 2013 , 3, 120185	7	77	
	31	The superfamily of heme-copper oxygen reductases: types and evolutionary considerations. <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2012 , 1817, 629-37	4.6	121	
	30	Mechanisms underlying the dual-mode regulation of microtubule dynamics by Kip3/kinesin-8. <i>Molecular Cell</i> , 2011 , 43, 751-63	17.6	91	
	29	A bioinformatics classifier and database for heme-copper oxygen reductases. <i>PLoS ONE</i> , 2011 , 6, e1911	7 3.7	40	
;	28	Evolution: Tracing the origins of centrioles, cilia, and flagella. <i>Journal of Cell Biology</i> , 2011 , 194, 165-75	7.3	257	
	27	Thousands of rab GTPases for the cell biologist. <i>PLoS Computational Biology</i> , 2011 , 7, e1002217	5	136	
;	26	Loss of genetic redundancy in reductive genome evolution. <i>PLoS Computational Biology</i> , 2011 , 7, e1001	082	47	
:	25	MAIS-TB: An Integrated Web Tool for Molecular Epidemiology Analysis. <i>Lecture Notes in Computer Science</i> , 2011 , 183-185	0.9		

24	Stepwise evolution of the centriole-assembly pathway. <i>Journal of Cell Science</i> , 2010 , 123, 1414-26	5.3	164
23	Single choroideremia gene in nonmammalian vertebrates explains early embryonic lethality of the zebrafish model of choroideremia 2009 , 50, 3009-16		28
22	Evolution and dynamics of protein interactions and networks. <i>Current Opinion in Structural Biology</i> , 2008 , 18, 349-57	8.1	93
21	The Ypt/Rab family and the evolution of trafficking in fungi. <i>Traffic</i> , 2008 , 9, 27-38	5.7	56
20	Multiple domain insertions and losses in the evolution of the Rab prenylation complex. <i>BMC Evolutionary Biology</i> , 2007 , 7, 140	3	17
19	Evolution of protein complexes by duplication of homomeric interactions. <i>Genome Biology</i> , 2007 , 8, R51	18.3	139
18	3D complex: a structural classification of protein complexes. <i>PLoS Computational Biology</i> , 2006 , 2, e155	5	256
17	The origins and evolution of functional modules: lessons from protein complexes. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2006 , 361, 507-17	5.8	113
16	Comparative genomics of trypanosomatid parasitic protozoa. <i>Science</i> , 2005 , 309, 404-9	33.3	614
15	The relationship between domain duplication and recombination. <i>Journal of Molecular Biology</i> , 2005 , 346, 355-65	6.5	131
14	The vertebrate Hef ortholog is a component of the Fanconi anemia tumor-suppressor pathway. <i>Nature Structural and Molecular Biology</i> , 2005 , 12, 763-71	17.6	163
13	Novel specificities emerge by stepwise duplication of functional modules. <i>Genome Research</i> , 2005 , 15, 552-9	9.7	70
12	An exponential core in the heart of the yeast protein interaction network. <i>Molecular Biology and Evolution</i> , 2005 , 22, 421-5	8.3	44
11	Functional evolution of the yeast protein interaction network. <i>Molecular Biology and Evolution</i> , 2004 , 21, 1171-6	8.3	56
10	Detection of functional modules from protein interaction networks. <i>Proteins: Structure, Function and Bioinformatics</i> , 2004 , 54, 49-57	4.2	288
9	Classification schemes for protein structure and function. <i>Nature Reviews Genetics</i> , 2003 , 4, 508-19	30.1	74
8	Structural determinants of Rab and Rab Escort Protein interaction: Rab family motifs define a conserved binding surface. <i>Biochemical and Biophysical Research Communications</i> , 2003 , 301, 92-7	3.4	17
7	Beyond 100 genomes. <i>Genome Biology</i> , 2003 , 4, 402	18.3	19

LIST OF PUBLICATIONS

6	Genome evolution reveals biochemical networks and functional modules. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2003 , 100, 15428-33	11.5	123
5	Analysis and preparation of stable complexes between Rab GTPases, Rab escort protein, and Rab geranylgeranyl transferase. <i>Methods in Molecular Biology</i> , 2002 , 189, 157-65	1.4	1
4	Evolution of the Rab family of small GTP-binding proteins. <i>Journal of Molecular Biology</i> , 2001 , 313, 889-	-9 6 .5	626
3	Prenylation of Rab GTPases: molecular mechanisms and involvement in genetic disease. <i>FEBS Letters</i> , 2001 , 498, 197-200	3.8	132
2	The mammalian Rab family of small GTPases: definition of family and subfamily sequence motifs suggests a mechanism for functional specificity in the Ras superfamily. <i>Journal of Molecular Biology</i> , 2000 , 301, 1077-87	6.5	375
1	An ancestral role of pericentrin in centriole formation through SAS-6 recruitment		2