

Jordi Paps

List of Publications by Citations

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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.

38
papers

3,409
citations

23
h-index

41
g-index

41
ext. papers

4,201
ext. citations

8.5
avg, IF

5.03
L-index

#	Paper	IF	Citations
38	The oyster genome reveals stress adaptation and complexity of shell formation. <i>Nature</i> , 2012 , 490, 49-54	30.4	1464
37	The genomes of four tapeworm species reveal adaptations to parasitism. <i>Nature</i> , 2013 , 496, 57-63	50.4	483
36	A phylogenetic analysis of myosin heavy chain type II sequences corroborates that Acoela and Nemertodermatida are basal bilaterians. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2002 , 99, 11246-51	11.5	180
35	Phylogenetic relationships within the Opisthokonta based on phylogenomic analyses of conserved single-copy protein domains. <i>Molecular Biology and Evolution</i> , 2012 , 29, 531-44	8.3	133
34	The phylogenetic position of ctenophores and the origin(s) of nervous systems. <i>EvoDevo</i> , 2015 , 6, 1	3.2	97
33	Lophotrochozoa internal phylogeny: new insights from an up-to-date analysis of nuclear ribosomal genes. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2009 , 276, 1245-54	4.4	96
32	Bilateria phylogeny: a broad sampling of 13 nuclear genes provides a new Lophotrochozoa phylogeny and supports a paraphyletic basal acoelomorpha. <i>Molecular Biology and Evolution</i> , 2009 , 26, 2397-406	8.3	80
31	Molecular phylogeny of unikonts: new insights into the position of apusomonads and ancyromonads and the internal relationships of opisthokonts. <i>Protist</i> , 2013 , 164, 2-12	2.5	76
30	Evolutionary origins of sensation in metazoans: functional evidence for a new sensory organ in sponges. <i>BMC Evolutionary Biology</i> , 2014 , 14, 3	3	76
29	SMG-1 and mTORC1 act antagonistically to regulate response to injury and growth in planarians. <i>PLoS Genetics</i> , 2012 , 8, e1002619	6	64
28	Back in time: a new systematic proposal for the Bilateria. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2008 , 363, 1481-91	5.8	58
27	New genes from old: asymmetric divergence of gene duplicates and the evolution of development. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2017 , 372,	5.8	55
26	Reconstruction of the ancestral metazoan genome reveals an increase in genomic novelty. <i>Nature Communications</i> , 2018 , 9, 1730	17.4	55
25	Human oxygen sensing may have origins in prokaryotic elongation factor Tu prolyl-hydroxylation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014 , 111, 13331-6	11.5	52
24	Evolutionary history of the Tricladida and the Platyhelminthes: an up-to-date phylogenetic and systematic account. <i>International Journal of Developmental Biology</i> , 2012 , 56, 5-17	1.9	46
23	Hagfish and lamprey Hox genes reveal conservation of temporal colinearity in vertebrates. <i>Nature Ecology and Evolution</i> , 2018 , 2, 859-866	12.3	39
22	Widespread patterns of gene loss in the evolution of the animal kingdom. <i>Nature Ecology and Evolution</i> , 2020 , 4, 519-523	12.3	37

21	The Origin of Land Plants Is Rooted in Two Bursts of Genomic Novelty. <i>Current Biology</i> , 2020 , 30, 530-536. 32	36
20	A genome-wide view of transcription factor gene diversity in chordate evolution: less gene loss in amphioxus?. <i>Briefings in Functional Genomics</i> , 2012 , 11, 177-86	4.9 34
19	Hox and ParaHox genes in Nemertodermatida, a basal bilaterian clade. <i>International Journal of Developmental Biology</i> , 2006 , 50, 675-9	1.9 34
18	Reinforcing the egg-timer: recruitment of novel lophotrochozoa homeobox genes to early and late development in the pacific oyster. <i>Genome Biology and Evolution</i> , 2015 , 7, 677-88	3.9 31
17	Genomic adaptations to aquatic and aerial life in mayflies and the origin of insect wings. <i>Nature Communications</i> , 2020 , 11, 2631	17.4 27
16	Metabarcoding analysis on European coastal samples reveals new molecular metazoan diversity. <i>Scientific Reports</i> , 2018 , 8, 9106	4.9 24
15	Acoelomorpha: earliest branching bilaterians or deuterostomes?. <i>Organisms Diversity and Evolution</i> , 2016 , 16, 391-399	1.7 15
14	Novel and divergent genes in the evolution of placental mammals. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2017 , 284,	4.4 15
13	The phylogenetic utility and functional constraint of microRNA flanking sequences. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2015 , 282, 20142983	4.4 15
12	Expression of the pupal determinant broad during metamorphic and neotenic development of the strepsipteran <i>Xenos vesparum</i> Rossi. <i>PLoS ONE</i> , 2014 , 9, e93614	3.7 14
11	A cytosolic copper storage protein provides a second level of copper tolerance in <i>Streptomyces lividans</i> . <i>Metallomics</i> , 2018 , 10, 180-193	4.5 13
10	Molecular phylogeny of the phylum Gastrotricha: new data brings together molecules and morphology. <i>Molecular Phylogenetics and Evolution</i> , 2012 , 63, 208-12	4.1 13
9	Plant Evolution: Assembling Land Plants. <i>Current Biology</i> , 2020 , 30, R81-R83	6.3 12
8	What Makes an Animal? The Molecular Quest for the Origin of the Animal Kingdom. <i>Integrative and Comparative Biology</i> , 2018 , 58, 654-665	2.8 10
7	The evolutionary emergence of land plants. <i>Current Biology</i> , 2021 , 31, R1281-R1298	6.3 7
6	One fold, two functions: cytochrome P460 and cytochrome W from the methanotroph (Bath). <i>Chemical Science</i> , 2019 , 10, 3031-3041	9.4 6
5	Animals and Their Unicellular Ancestors 2010 ,	4
4	Discovery and classification of homeobox genes in animal genomes. <i>Methods in Molecular Biology</i> , 2014 , 1196, 3-18	1.4 4

- 3 Water-related innovations in land plants evolved by different patterns of gene co-option and novelty.. *New Phytologist*, **2022**, 9.8 2
- 2 Evolutionary Origins of Drought Tolerance in Spermatophytes. *Frontiers in Plant Science*, **2021**, 12, 6559242 0
- 1 Unravelling body plan and axial evolution in the Bilateria with molecular phylogenetic markers217-238