## Romain Derelle

List of Publications by Year in descending order

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567281 888059 2,447 17 15 17 citations h-index g-index papers 19 19 19 2855 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Analysis of diverse eukaryotes suggests the existence of an ancestral mitochondrial apparatus derived from the bacterial type II secretion system. Nature Communications, 2021, 12, 2947.	12.8	19
2	A Eukaryote-Wide Perspective on the Diversity and Evolution of the ARF GTPase Protein Family. Genome Biology and Evolution, $2021,13,.$	2.5	18
3	The draft nuclear genome sequence and predicted mitochondrial proteome of Andalucia godoyi, a protist with the most gene-rich and bacteria-like mitochondrial genome. BMC Biology, 2020, 18, 22.	3.8	43
4	Spatial and temporal variation in river corridor exchange across a 5th-order mountain stream network. Hydrology and Earth System Sciences, 2019, 23, 5199-5225.	4.9	23
5	Co-located contemporaneous mapping of morphological, hydrological, chemical, and biological conditions in a 5th-order mountain stream network, Oregon, USA. Earth System Science Data, 2019, 11, 1567-1581.	9.9	14
6	Nuclear genetic codes with a different meaning of the UAG and the UAA codon. BMC Biology, 2017, 15, 8.	3.8	25
7	A Phylogenomic Framework to Study the Diversity and Evolution of Stramenopiles (=Heterokonts). Molecular Biology and Evolution, 2016, 33, 2890-2898.	8.9	125
8	A paneukaryotic genomic analysis of the small GTPase RABL2 underscores the significance of recurrent gene loss in eukaryote evolution. Biology Direct, 2016, 11, 5.	4.6	22
9	Bacterial proteins pinpoint a single eukaryotic root. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, E693-9.	7.1	159
10	The ctenophore genome and the evolutionary origins of neural systems. Nature, 2014, 510, 109-114.	27.8	606
11	The Capsaspora genome reveals a complex unicellular prehistory of animals. Nature Communications, 2013, 4, 2325.	12.8	244
12	Rooting the Eukaryotic Tree with Mitochondrial and Bacterial Proteins. Molecular Biology and Evolution, 2012, 29, 1277-1289.	8.9	121
13	Phylogenetic Relationships within the Opisthokonta Based on Phylogenomic Analyses of Conserved Single-Copy Protein Domains. Molecular Biology and Evolution, 2012, 29, 531-544.	8.9	166
14	Convergent origins and rapid evolution of spliced leader <i>trans</i> -splicing in Metazoa: Insights from the Ctenophora and Hydrozoa. Rna, 2010, 16, 696-707.	3.5	35
15	WNT/ $\hat{l}^2$ -Catenin Signalling and Epithelial Patterning in the Homoscleromorph Sponge Oscarella. PLoS ONE, 2009, 4, e5823.	2.5	68
16	Phylogenomics Revives Traditional Views on Deep Animal Relationships. Current Biology, 2009, 19, 706-712.	3.9	611
17	A maternally localised Wnt ligand required for axial patterning in the cnidarian <i>Clytia hemisphaerica</i> . Development (Cambridge), 2008, 135, 2105-2113.	2.5	144