

Julien Soubrier

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.

21
papers

2,278
citations

17
h-index

23
g-index

23
ext. papers

2,886
ext. citations

13.7
avg, IF

4.39
L-index

#	Paper	IF	Citations
21	Whole-Genome Sequencing of SARS-CoV-2 from Quarantine Hotel Outbreak. <i>Emerging Infectious Diseases</i> , 2021 , 27, 2219-2221	10.2	4
20	Genome of the Tasmanian tiger provides insights into the evolution and demography of an extinct marsupial carnivore. <i>Nature Ecology and Evolution</i> , 2018 , 2, 182-192	12.3	48
19	Aboriginal mitogenomes reveal 50,000 years of regionalism in Australia. <i>Nature</i> , 2017 , 544, 180-184	50.4	122
18	Neanderthal behaviour, diet, and disease inferred from ancient DNA in dental calculus. <i>Nature</i> , 2017 , 544, 357-361	50.4	263
17	Experimental conditions improving in-solution target enrichment for ancient DNA. <i>Molecular Ecology Resources</i> , 2017 , 17, 508-522	8.4	48
16	Did the historical range of the European bison (<i>Bison bonasus</i> L.) extend further south? A new finding from the Yenikapı Metro and Marmaray excavation, Turkey. <i>Mammal Research</i> , 2017 , 62, 103-109	1.8	3
15	Ancient mitochondrial DNA provides high-resolution time scale of the peopling of the Americas. <i>Science Advances</i> , 2016 , 2, e1501385	14.3	211
14	Early cave art and ancient DNA record the origin of European bison. <i>Nature Communications</i> , 2016 , 7, 13158	17.4	63
13	Pleistocene Chinese cave hyenas and the recent Eurasian history of the spotted hyena, <i>Crocuta crocuta</i> . <i>Molecular Ecology</i> , 2014 , 23, 522-33	5.7	21
12	Environmental metabarcodes for insects: in silico PCR reveals potential for taxonomic bias. <i>Molecular Ecology Resources</i> , 2014 , 14, 1160-70	8.4	195
11	Ancient DNA reveals elephant birds and kiwi are sister taxa and clarifies ratite bird evolution. <i>Science</i> , 2014 , 344, 898-900	33.3	189
10	Modular tagging of amplicons using a single PCR for high-throughput sequencing. <i>Molecular Ecology Resources</i> , 2014 , 14, 117-21	8.4	25
9	Mitochondrial genome sequencing in Mesolithic North East Europe Unearths a new sub-clade within the broadly distributed human haplogroup C1. <i>PLoS ONE</i> , 2014 , 9, e87612	3.7	15
8	DNA capture and next-generation sequencing can recover whole mitochondrial genomes from highly degraded samples for human identification. <i>Investigative Genetics</i> , 2013 , 4, 26		78
7	Rates of phenotypic and genomic evolution during the Cambrian explosion. <i>Current Biology</i> , 2013 , 23, 1889-95	6.3	105
6	The origins of the enigmatic Falkland Islands wolf. <i>Nature Communications</i> , 2013 , 4, 1552	17.4	33
5	Neolithic mitochondrial haplogroup H genomes and the genetic origins of Europeans. <i>Nature Communications</i> , 2013 , 4, 1764	17.4	141

4	The influence of rate heterogeneity among sites on the time dependence of molecular rates. <i>Molecular Biology and Evolution</i> , 2012 , 29, 3345-58	8.3	140
3	Multiple geographic origins of commensalism and complex dispersal history of Black Rats. <i>PLoS ONE</i> , 2011 , 6, e26357	3.7	189
2	Time-dependent rates of molecular evolution. <i>Molecular Ecology</i> , 2011 , 20, 3087-101	5.7	383
1	Population size history from short genomic scaffolds: how short is too short?		2