

Julien Soubrier

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2084116/publications.pdf>

Version: 2024-02-01

21
papers

3,211
citations

430442

18
h-index

752256

20
g-index

23
all docs

23
docs citations

23
times ranked

6390
citing authors

#	ARTICLE	IF	CITATIONS
1	Time-dependent rates of molecular evolution. <i>Molecular Ecology</i> , 2011, 20, 3087-3101.	2.0	473
2	Neanderthal behaviour, diet, and disease inferred from ancient DNA in dental calculus. <i>Nature</i> , 2017, 544, 357-361.	13.7	398
3	Ancient mitochondrial DNA provides high-resolution time scale of the peopling of the Americas. <i>Science Advances</i> , 2016, 2, e1501385.	4.7	306
4	The Influence of Rate Heterogeneity among Sites on the Time Dependence of Molecular Rates. <i>Molecular Biology and Evolution</i> , 2012, 29, 3345-3358.	3.5	275
5	Environmental metabarcodes for insects: <i>in silico</i> PCR reveals potential for taxonomic bias. <i>Molecular Ecology Resources</i> , 2014, 14, 1160-1170.	2.2	261
6	Multiple Geographic Origins of Commensalism and Complex Dispersal History of Black Rats. <i>PLoS ONE</i> , 2011, 6, e26357.	1.1	250
7	Ancient DNA reveals elephant birds and kiwi are sister taxa and clarifies ratite bird evolution. <i>Science</i> , 2014, 344, 898-900.	6.0	247
8	Aboriginal mitogenomes reveal 50,000 years of regionalism in Australia. <i>Nature</i> , 2017, 544, 180-184.	13.7	195
9	Neolithic mitochondrial haplogroup H genomes and the genetic origins of Europeans. <i>Nature Communications</i> , 2013, 4, 1764.	5.8	180
10	Rates of Phenotypic and Genomic Evolution during the Cambrian Explosion. <i>Current Biology</i> , 2013, 23, 1889-1895.	1.8	140
11	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.	3.3	101
12	Early cave art and ancient DNA record the origin of European bison. <i>Nature Communications</i> , 2016, 7, 13158.	5.8	81
13	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.	3.4	78
14	Experimental conditions improving <i>in silico</i> resolution target enrichment for ancient DNA. <i>Molecular Ecology Resources</i> , 2017, 17, 508-522.	2.2	67
15	The origins of the enigmatic Falkland Islands wolf. <i>Nature Communications</i> , 2013, 4, 1552.	5.8	40
16	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.	1.1	34
17	Modular tagging of amplicons using a single PCR for high-throughput sequencing. <i>Molecular Ecology Resources</i> , 2014, 14, 117-121.	2.2	29
18	Pleistocene Chinese cave hyenas and the recent Eurasian history of the spotted hyena, <i>Crocuta crocuta</i> . <i>Molecular Ecology</i> , 2014, 23, 522-533.	2.0	29

#	ARTICLE	IF	CITATIONS
19	Whole-Genome Sequencing of SARS-CoV-2 from Quarantine Hotel Outbreak. <i>Emerging Infectious Diseases</i> , 2021, 27, 2219-2221.	2.0	9
20	Did the historical range of the European bison (<i>Bison bonasus</i> L.) extend further south?â€”a new finding from the YenikapÄ± Metro and Marmaray excavation, Turkey. <i>Mammal Research</i> , 2017, 62, 103-109.	0.6	4
21	Review of 12 months of copy number variant calling on a clinical next generation sequencing pipeline. <i>Pathology</i> , 2020, 52, S108.	0.3	0