

Jacqueline A Robinson

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5055974/publications.pdf>

Version: 2024-02-01

16
papers

1,156
citations

759233

12
h-index

940533

16
g-index

22
all docs

22
docs citations

22
times ranked

1706
citing authors

#	ARTICLE	IF	CITATIONS
1	Genomic Flatlining in the Endangered Island Fox. <i>Current Biology</i> , 2016, 26, 1183-1189.	3.9	201
2	Genomic signatures of extensive inbreeding in Isle Royale wolves, a population on the threshold of extinction. <i>Science Advances</i> , 2019, 5, eaau0757.	10.3	173
3	Genome-wide Evidence Reveals that African and Eurasian Golden Jackals Are Distinct Species. <i>Current Biology</i> , 2015, 25, 2158-2165.	3.9	156
4	Whole-genome sequence analysis shows that two endemic species of North American wolf are admixtures of the coyote and gray wolf. <i>Science Advances</i> , 2016, 2, e1501714.	10.3	150
5	Purging of Strongly Deleterious Mutations Explains Long-Term Persistence and Absence of Inbreeding Depression in Island Foxes. <i>Current Biology</i> , 2018, 28, 3487-3494.e4.	3.9	140
6	Genetics and extinction and the example of Isle Royale wolves. <i>Animal Conservation</i> , 2019, 22, 302-309.	2.9	56
7	Reference genome and demographic history of the most endangered marine mammal, the vaquita. <i>Molecular Ecology Resources</i> , 2021, 21, 1008-1020.	4.8	54
8	The critically endangered vaquita is not doomed to extinction by inbreeding depression. <i>Science</i> , 2022, 376, 635-639.	12.6	49
9	Targeted capture and resequencing of 1040 genes reveal environmentally driven functional variation in grey wolves. <i>Molecular Ecology</i> , 2016, 25, 357-379.	3.9	47
10	Genome-wide diversity in the California condor tracks its prehistoric abundance and decline. <i>Current Biology</i> , 2021, 31, 2939-2946.e5.	3.9	35
11	Analysis of 100 high-coverage genomes from a pedigreed captive baboon colony. <i>Genome Research</i> , 2019, 29, 848-856.	5.5	27
12	Accurate assembly of the olive baboon (<i>Papio anubis</i>) genome using long-read and Hi-C data. <i>GigaScience</i> , 2020, 9, .	6.4	18
13	A Genome-Wide Perspective on the Persistence of Red Wolf Ancestry in Southeastern Canids. <i>Journal of Heredity</i> , 2020, 111, 277-286.	2.4	16
14	Attempted DNA extraction from a R ancho L a B rea C olumbian mammoth (<i>M ammuthus columbi</i>): prospects for ancient DNA from asphalt deposits. <i>Ecology and Evolution</i> , 2014, 4, 329-336.	1.9	9
15	High-Resolution Estimates of Crossover and Noncrossover Recombination from a Captive Baboon Colony. <i>Genome Biology and Evolution</i> , 2022, 14, .	2.5	9
16	Response to Bakker et al.. <i>Current Biology</i> , 2022, 32, R358-R359.	3.9	1