

Pere Gelabert

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8626280/publications.pdf>

Version: 2024-02-01

14
papers

410
citations

1163117

8
h-index

1058476

14
g-index

20
all docs

20
docs citations

20
times ranked

989
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--|------|-----------|
| 1 | The dental proteome of Homo antecessor. Nature, 2020, 580, 235-238. | 27.8 | 100 |
| 2 | Differential DNA methylation of vocal and facial anatomy genes in modern humans. Nature Communications, 2020, 11, 1189. | 12.8 | 69 |
| 3 | Mitochondrial DNA from the eradicated European <i>Plasmodium vivax</i> and <i>P. falciparum</i> from 70-year-old slides from the Ebro Delta in Spain. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 11495-11500. | 7.1 | 41 |
| 4 | Plasmodium vivax Malaria Viewed through the Lens of an Eradicated European Strain. Molecular Biology and Evolution, 2020, 37, 773-785. | 8.9 | 38 |
| 5 | Genome-scale sequencing and analysis of human, wolf, and bison DNA from 25,000-year-old sediment. Current Biology, 2021, 31, 3564-3574.e9. | 3.9 | 34 |
| 6 | Malaria was a weak selective force in ancient Europeans. Scientific Reports, 2017, 7, 1377. | 3.3 | 32 |
| 7 | Evolutionary History, Genomic Adaptation to Toxic Diet, and Extinction of the Carolina Parakeet. Current Biology, 2020, 30, 108-114.e5. | 3.9 | 24 |
| 8 | The epidemiology of Plasmodium vivax among adults in the Democratic Republic of the Congo. Nature Communications, 2021, 12, 4169. | 12.8 | 18 |
| 9 | TKGWV2: an ancient DNA relatedness pipeline for ultra-low coverage whole genome shotgun data. Scientific Reports, 2021, 11, 21262. | 3.3 | 12 |
| 10 | Genetic affinities of an eradicated European Plasmodium falciparum strain. Microbial Genomics, 2019, 5, . | 2.0 | 9 |
| 11 | Mapping co-ancestry connections between the genome of a Medieval individual and modern Europeans. Scientific Reports, 2020, 10, 6843. | 3.3 | 8 |
| 12 | Genome-wide data from the Bubi of Bioko Island clarifies the Atlantic fringe of the Bantu dispersal. BMC Genomics, 2019, 20, 179. | 2.8 | 6 |
| 13 | Genomes from Vertebea cave suggest diversity within the Trypillians in Ukraine. Scientific Reports, 2022, 12, 7242. | 3.3 | 2 |
| 14 | Evolutionary History, Genomic Adaptation to Toxic Diet and Extinction of the Carolina Parakeet. SSRN Electronic Journal, 0, , . | 0.4 | 0 |