

Anders Bergström

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4357432/publications.pdf>

Version: 2024-02-01

28
papers

4,684
citations

279798

23
h-index

501196

28
g-index

30
all docs

30
docs citations

30
times ranked

6650
citing authors

#	ARTICLE	IF	CITATIONS
1	Grey wolf genomic history reveals a dual ancestry of dogs. <i>Nature</i> , 2022, 607, 313-320.	27.8	48
2	Origins of modern human ancestry. <i>Nature</i> , 2021, 590, 229-237.	27.8	166
3	Million-year-old DNA sheds light on the genomic history of mammoths. <i>Nature</i> , 2021, 591, 265-269.	27.8	179
4	Genome-scale sequencing and analysis of human, wolf, and bison DNA from 25,000-year-old sediment. <i>Current Biology</i> , 2021, 31, 3564-3574.e9.	3.9	34
5	Origins and genetic legacy of prehistoric dogs. <i>Science</i> , 2020, 370, 557-564.	12.6	152
6	Archaeological Central American maize genomes suggest ancient gene flow from South America. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 33124-33129.	7.1	36
7	Population Structure, Stratification, and Introgression of Human Structural Variation. <i>Cell</i> , 2020, 182, 189-199.e15.	28.9	79
8	Insights into human genetic variation and population history from 929 diverse genomes. <i>Science</i> , 2020, 367, .	12.6	534
9	Uganda Genome Resource Enables Insights into Population History and Genomic Discovery in Africa. <i>Cell</i> , 2019, 179, 984-1002.e36.	28.9	152
10	Y Chromosome Sequences Reveal a Short Beringian Standstill, Rapid Expansion, and early Population structure of Native American Founders. <i>Current Biology</i> , 2019, 29, 149-157.e3.	3.9	94
11	Genome evolution across 1,011 <i>Saccharomyces cerevisiae</i> isolates. <i>Nature</i> , 2018, 556, 339-344.	27.8	952
12	Copy number variation arising from gene conversion on the human Y chromosome. <i>Human Genetics</i> , 2018, 137, 73-83.	3.8	9
13	Human Genetics: Busy Subway Networks in Remote Oceania?. <i>Current Biology</i> , 2018, 28, R549-R551.	3.9	2
14	Y-chromosomal sequences of diverse Indian populations and the ancestry of the Andamanese. <i>Human Genetics</i> , 2017, 136, 499-510.	3.8	18
15	Contrasting evolutionary genome dynamics between domesticated and wild yeasts. <i>Nature Genetics</i> , 2017, 49, 913-924.	21.4	340
16	Paleolithic networking. <i>Science</i> , 2017, 358, 586-587.	12.6	1
17	Clonal Heterogeneity Influences the Fate of New Adaptive Mutations. <i>Cell Reports</i> , 2017, 21, 732-744.	6.4	70
18	A Neolithic expansion, but strong genetic structure, in the independent history of New Guinea. <i>Science</i> , 2017, 357, 1160-1163.	12.6	45

#	ARTICLE	IF	CITATIONS
19	Chad Genetic Diversity Reveals an African History Marked by Multiple Holocene Eurasian Migrations. <i>American Journal of Human Genetics</i> , 2016, 99, 1316-1324.	6.2	37
20	A genomic history of Aboriginal Australia. <i>Nature</i> , 2016, 538, 207-214.	27.8	439
21	Deep Roots for Aboriginal Australian Y Chromosomes. <i>Current Biology</i> , 2016, 26, 809-813.	3.9	54
22	Extensive Recombination of a Yeast Diploid Hybrid through Meiotic Reversion. <i>PLoS Genetics</i> , 2016, 12, e1005781.	3.5	60
23	Genomic evidence for the Pleistocene and recent population history of Native Americans. <i>Science</i> , 2015, 349, aab3884.	12.6	449
24	A High-Definition View of Functional Genetic Variation from Natural Yeast Genomes. <i>Molecular Biology and Evolution</i> , 2014, 31, 872-888.	8.9	328
25	High quality de novo sequencing and assembly of the <i>Saccharomyces arboricolus</i> genome. <i>BMC Genomics</i> , 2013, 14, 69.	2.8	87
26	High-Resolution Mapping of Complex Traits with a Four-Parent Advanced Intercross Yeast Population. <i>Genetics</i> , 2013, 195, 1141-1155.	2.9	164
27	Inferring Genome-Wide Recombination Landscapes from Advanced Intercross Lines: Application to Yeast Crosses. <i>PLoS ONE</i> , 2013, 8, e62266.	2.5	29
28	The Genetic Basis of Natural Variation in Oenological Traits in <i>Saccharomyces cerevisiae</i> . <i>PLoS ONE</i> , 2012, 7, e49640.	2.5	107