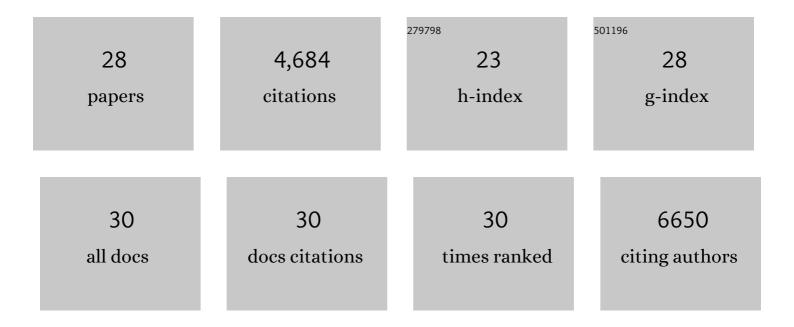
Anders Bergström

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4357432/publications.pdf Version: 2024-02-01



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

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