

Deepti Gurdasani

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2392742/publications.pdf>

Version: 2024-02-01

39
papers

6,859
citations

304743

22
h-index

289244

40
g-index

45
all docs

45
docs citations

45
times ranked

15746
citing authors

#	ARTICLE	IF	CITATIONS
1	Discovery and refinement of loci associated with lipid levels. <i>Nature Genetics</i> , 2013, 45, 1274-1283.	21.4	2,641
2	Common variants associated with plasma triglycerides and risk for coronary artery disease. <i>Nature Genetics</i> , 2013, 45, 1345-1352.	21.4	754
3	A General Approach for Haplotype Phasing across the Full Spectrum of Relatedness. <i>PLoS Genetics</i> , 2014, 10, e1004234.	3.5	553
4	The African Genome Variation Project shapes medical genetics in Africa. <i>Nature</i> , 2015, 517, 327-332.	27.8	473
5	Long reads: their purpose and place. <i>Human Molecular Genetics</i> , 2018, 27, R234-R241.	2.9	249
6	Genomics of disease risk in globally diverse populations. <i>Nature Reviews Genetics</i> , 2019, 20, 520-535.	16.3	217
7	Scientific consensus on the COVID-19 pandemic: we need to act now. <i>Lancet, The</i> , 2020, 396, e71-e72.	13.7	189
8	Association of HIV and ART with cardiometabolic traits in sub-Saharan Africa: a systematic review and meta-analysis. <i>International Journal of Epidemiology</i> , 2013, 42, 1754-1771.	1.9	158
9	The COVID-19 vaccines rush: participatory community engagement matters more than ever. <i>Lancet, The</i> , 2021, 397, 8-10.	13.7	156
10	Polymorphisms of large effect explain the majority of the host genetic contribution to variation of HIV-1 virus load. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 14658-14663.	7.1	154
11	Tracing the Route of Modern Humans out of Africa by Using 225 Human Genome Sequences from Ethiopians and Egyptians. <i>American Journal of Human Genetics</i> , 2015, 96, 986-991.	6.2	152
12	Uganda Genome Resource Enables Insights into Population History and Genomic Discovery in Africa. <i>Cell</i> , 2019, 179, 984-1002.e36.	28.9	152
13	Lipoprotein(a) and Risk of Coronary, Cerebrovascular, and Peripheral Artery Disease. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2012, 32, 3058-3065.	2.4	146
14	A systematic review of definitions of extreme phenotypes of HIV control and progression. <i>Aids</i> , 2014, 28, 149-162.	2.2	83
15	The Association Between Circulating Lipoprotein(a) and Type 2 Diabetes: Is It Causal?. <i>Diabetes</i> , 2014, 63, 332-342.	0.6	82
16	An Evaluation of HIV Elite Controller Definitions within a Large Seroconverter Cohort Collaboration. <i>PLoS ONE</i> , 2014, 9, e86719.	2.5	80
17	Linear mixed model for heritability estimation that explicitly addresses environmental variation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 7377-7382.	7.1	75
18	The transferability of lipid loci across African, Asian and European cohorts. <i>Nature Communications</i> , 2019, 10, 4330.	12.8	75

#	ARTICLE	IF	CITATIONS
19	Calling for pan-European commitment for rapid and sustained reduction in SARS-CoV-2 infections. <i>Lancet, The</i> , 2021, 397, 92-93.	13.7	71
20	Herd immunity by infection is not an option. <i>Science</i> , 2021, 371, 230-231.	12.6	47
21	School reopening without robust COVID-19 mitigation risks accelerating the pandemic. <i>Lancet, The</i> , 2021, 397, 1177-1178.	13.7	46
22	Vaccinating adolescents against SARS-CoV-2 in England: a risk-benefit analysis. <i>Journal of the Royal Society of Medicine</i> , 2021, 114, 513-524.	2.0	32
23	Complimentary Methods for Multivariate Genome-Wide Association Study Identify New Susceptibility Genes for Blood Cell Traits. <i>Frontiers in Genetics</i> , 2019, 10, 334.	2.3	31
24	Distinct genetic architectures and environmental factors associate with host response to the β 2-herpesvirus infections. <i>Nature Communications</i> , 2020, 11, 3849.	12.8	24
25	Mass infection is not an option: we must do more to protect our young. <i>Lancet, The</i> , 2021, 398, 297-298.	13.7	24
26	Health equality, race and pharmacogenomics. <i>British Journal of Clinical Pharmacology</i> , 2022, 88, 27-33.	2.4	24
27	The UK needs a sustainable strategy for COVID-19. <i>Lancet, The</i> , 2020, 396, 1800-1801.	13.7	23
28	Evaluating the Impact of Functional Genetic Variation on HIV-1 Control. <i>Journal of Infectious Diseases</i> , 2017, 216, 1063-1069.	4.0	20
29	The First Norovirus Longitudinal Seroepidemiological Study From Sub-Saharan Africa Reveals High Seroprevalence of Diverse Genotypes Associated With Host Susceptibility Factors. <i>Journal of Infectious Diseases</i> , 2018, 218, 716-725.	4.0	20
30	The ferroportin Q248H mutation protects from anemia, but not malaria or bacteremia. <i>Science Advances</i> , 2019, 5, eaaw0109.	10.3	20
31	Delaying the second dose of covid-19 vaccines. <i>BMJ, The</i> , 2021, 372, n710.	6.0	20
32	On the fallibility of simulation models in informing pandemic responses. <i>The Lancet Global Health</i> , 2020, 8, e776-e777.	6.3	15
33	HIV treatment is associated with a twofold higher probability of raised triglycerides: pooled analyses in 21,023 individuals in sub-Saharan Africa. <i>Global Health, Epidemiology and Genomics</i> , 2018, 3, .	0.8	11
34	Long COVID in children. <i>The Lancet Child and Adolescent Health</i> , 2022, 6, e2.	5.6	10
35	Association between early life antibiotic use and childhood overweight and obesity: a narrative review. <i>Global Health, Epidemiology and Genomics</i> , 2018, 3, e18.	0.8	8
36	Modelling the impact of lockdown-easing measures on cumulative COVID-19 cases and deaths in England. <i>BMJ Open</i> , 2021, 11, e042483.	1.9	5

#	ARTICLE	IF	CITATIONS
37	The World Health Network: a global citizens' initiative. <i>Lancet, The</i> , 2021, 398, 1567-1568.	13.7	3
38	Insights into the genetic architecture of haematological traits from deep phenotyping and whole-genome sequencing for two Mediterranean isolated populations. <i>Scientific Reports</i> , 2022, 12, 1131.	3.3	2
39	Visual Vignette. <i>Endocrine Practice</i> , 2008, 14, 255.	2.1	1