Deepti Gurdasani

List of Publications by Year
in descending order

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	Health equality, race and pharmacogenomics. British Journal of Clinical Pharmacology, 2022, 88, 27-33.	2.4	24
2	Insights into the genetic architecture of haematological traits from deep phenotyping and whole-genome sequencing for two Mediterranean isolated populations. Scientific Reports, 2022, 12, 1131.	3.3	2
3	Long COVID in children. The Lancet Child and Adolescent Health, 2022, 6, e2.	5.6	10
4	Calling for pan-European commitment for rapid and sustained reduction in SARS-CoV-2 infections. Lancet, The, 2021, 397, 92-93.	13.7	71
5	The COVID-19 vaccines rush: participatory community engagement matters more than ever. Lancet, The, 2021, 397, 8-10.	13.7	156
6	Herd immunity by infection is not an option. Science, 2021, 371, 230-231.	12.6	47
7	Delaying the second dose of covid-19 vaccines. BMJ, The, 2021, 372, n710.	6.0	20
8	School reopening without robust COVID-19 mitigation risks accelerating the pandemic. Lancet, The, 2021, 397, 1177-1178.	13.7	46
9	Mass infection is not an option: we must do more to protect our young. Lancet, The, 2021, 398, 297-298.	13.7	24
10	Modelling the impact of lockdown-easing measures on cumulative COVID-19 cases and deaths in England. BMJ Open, 2021, 11, e042483.	1.9	5
11	The World Health Network: a global citizens' initiative. Lancet, The, 2021, 398, 1567-1568.	13.7	3
12	Vaccinating adolescents against SARS-CoV-2 in England: a riskâ€“benefit analysis. Journal of the Royal Society of Medicine, 2021, 114, 513-524.	2.0	32
13	Scientific consensus on the COVID-19 pandemic: we need to act now. Lancet, The, 2020, 396, e71-e72.	13.7	189
14	The UK needs a sustainable strategy for COVID-19. Lancet, The, 2020, 396, 1800-1801.	13.7	23
15	Distinct genetic architectures and environmental factors associate with host response to the Î³2-herpesvirus infections. Nature Communications, 2020, 11, 3849.	12.8	24
16	On the fallibility of simulation models in informing pandemic responses. The Lancet Global Health, 2020, 8, e776-e777.	6.3	15
17	Uganda Genome Resource Enables Insights into Population History and Genomic Discovery in Africa. Cell, 2019, 179, 984-1002.e36.	28.9	152
18	The ferroportin Q248H mutation protects from anemia, but not malaria or bacteremia. Science Advances, 2019, 5, eaaw0109.	10.3	20

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19	The transferability of lipid loci across African, Asian and European cohorts. Nature Communications, 2019, 10, 4330.	12.8	75
20	Genomics of disease risk in globally diverse populations. Nature Reviews Genetics, 2019, 20, 520-535.	16.3	217
21	Complimentary Methods for Multivariate Genome-Wide Association Study Identify New Susceptibility Genes for Blood Cell Traits. Frontiers in Genetics, 2019, 10, 334.	2.3	31
22	Association between early life antibiotic use and childhood overweight and obesity: a narrative review. Global Health, Epidemiology and Genomics, 2018, 3, e18.	0.8	8
23	Long reads: their purpose and place. Human Molecular Genetics, 2018, 27, R234-R241.	2.9	249
24	The First Norovirus Longitudinal Seroepidemiological Study From Sub-Saharan Africa Reveals High Seroprevalence of Diverse Genotypes Associated With Host Susceptibility Factors. Journal of Infectious Diseases, 2018, 218, 716-725.	4.0	20
25	HIV treatment is associated with a twofold higher probability of raised triglycerides: pooled analyses in 21Â023 individuals in sub-Saharan Africa. Global Health, Epidemiology and Genomics, 2018, 3, .	0.8	11
26	Evaluating the Impact of Functional Genetic Variation on HIV-1 Control. Journal of Infectious Diseases, 2017, 216, 1063-1069.	4.0	20
27	Linear mixed model for heritability estimation that explicitly addresses environmental variation. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 7377-7382.	7.1	75
28	Tracing the Route of Modern Humans out of Africa by Using 225 Human Genome Sequences from Ethiopians and Egyptians. American Journal of Human Genetics, 2015, 96, 986-991.	6.2	152
29	Polymorphisms of large effect explain the majority of the host genetic contribution to variation of HIV-1 virus load. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 14658-14663.	7.1	154
30	The African Genome Variation Project shapes medical genetics in Africa. Nature, 2015, 517, 327-332.	27.8	473
31	A General Approach for Haplotype Phasing across the Full Spectrum of Relatedness. PLoS Genetics, 2014, 10, e1004234.	3.5	553
32	A systematic review of definitions of extreme phenotypes of HIV control and progression. Aids, 2014, 28, 149-162.	2.2	83
33	The Association Between Circulating Lipoprotein(a) and Type 2 Diabetes: Is It Causal?. Diabetes, 2014, 63, 332-342.	0.6	82
34	An Evaluation of HIV Elite Controller Definitions within a Large Seroconverter Cohort Collaboration. PLoS ONE, 2014, 9, e86719.	2.5	80
35	Discovery and refinement of loci associated with lipid levels. Nature Genetics, 2013, 45, 1274-1283.	21.4	2,641
36	Common variants associated with plasma triglycerides and risk for coronary artery disease. Nature Genetics, 2013, 45, 1345-1352.	21.4	754

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37	Association of HIV and ART with cardiometabolic traits in sub-Saharan Africa: a systematic review and meta-analysis. International Journal of Epidemiology, 2013, 42, 1754-1771.	1.9	158
38	Lipoprotein(a) and Risk of Coronary, Cerebrovascular, and Peripheral Artery Disease. Arteriosclerosis, Thrombosis, and Vascular Biology, 2012, 32, 3058-3065.	2.4	146
39	Visual Vignette. Endocrine Practice, 2008, 14, 255.	2.1	1