

Manjinder S Sandhu

List of Publications by Year in descending order

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Version: 2024-02-01

45
papers

13,349
citations

201674

27
h-index

254184

43
g-index

46
all docs

46
docs citations

46
times ranked

22938
citing authors

#	ARTICLE	IF	CITATIONS
1	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
2	The trans-ancestral genomic architecture of glycemic traits. <i>Nature Genetics</i> , 2021, 53, 840-860.	21.4	341
3	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
4	Sociodemographic inequities associated with participation in leisure-time physical activity in sub-Saharan Africa: an individual participant data meta-analysis. <i>BMC Public Health</i> , 2020, 20, 927.	2.9	16
5	The transferability of lipid loci across African, Asian and European cohorts. <i>Nature Communications</i> , 2019, 10, 4330.	12.8	75
6	Sociodemographic patterns of health insurance coverage in Namibia. <i>International Journal for Equity in Health</i> , 2019, 18, 16.	3.5	24
7	Genomics of disease risk in globally diverse populations. <i>Nature Reviews Genetics</i> , 2019, 20, 520-535.	16.3	217
8	Characterisation and correlates of stunting among Malaysian children and adolescents aged 6â€“19 years. <i>Global Health, Epidemiology and Genomics</i> , 2019, 4, e2.	0.8	11
9	Objective measurement of physical activity: improving the evidence base to address non-communicable diseases in Africa. <i>BMJ Global Health</i> , 2018, 3, e001044.	4.7	9
10	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
11	A cross-sectional analysis of ITN and IRS coverage in Namibia in 2013. <i>Malaria Journal</i> , 2018, 17, 264.	2.3	5
12	HIV treatment is associated with a twofold higher probability of raised triglycerides: pooled analyses in 21Â023 individuals in sub-Saharan Africa. <i>Global Health, Epidemiology and Genomics</i> , 2018, 3, .	0.8	11
13	The Use of Different International References to Assess Child Anthropometric Status in a Malaysian Population. <i>Journal of Pediatrics</i> , 2017, 190, 63-68.e1.	1.8	13
14	Anthropometric and cardiometabolic risk factors in parents and child obesity in Segamat, Malaysia. <i>International Journal of Epidemiology</i> , 2017, 46, 1523-1532.	1.9	6
15	Ideal cardiovascular health influences cardiovascular disease risk associated with high lipoprotein(a) levels and genotype: The EPIC-Norfolk prospective population study. <i>Atherosclerosis</i> , 2017, 256, 47-52.	0.8	65
16	HDSS Profile: The South East Asia Community Observatory Health and Demographic Surveillance System (SEACO HDSS). <i>International Journal of Epidemiology</i> , 2017, 46, 1370-1371g.	1.9	37
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	Population and assay thresholds for the predictive value of lipoprotein (a) for coronary artery disease: the EPIC-Norfolk Prospective Population Study. <i>Journal of Lipid Research</i> , 2016, 57, 697-705.	4.2	24

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19	Burden of Diabetes and First Evidence for the Utility of HbA1c for Diagnosis and Detection of Diabetes in Urban Black South Africans: The Durban Diabetes Study. <i>PLoS ONE</i> , 2016, 11, e0161966.	2.5	38
20	Prevalence of Dyslipidaemia and Associated Risk Factors in a Rural Population in South-Western Uganda: A Community Based Survey. <i>PLoS ONE</i> , 2015, 10, e0126166.	2.5	45
21	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
22	The African Genome Variation Project shapes medical genetics in Africa. <i>Nature</i> , 2015, 517, 327-332.	27.8	473
23	A General Approach for Haplotype Phasing across the Full Spectrum of Relatedness. <i>PLoS Genetics</i> , 2014, 10, e1004234.	3.5	553
24	Urbanicity and Lifestyle Risk Factors for Cardiometabolic Diseases in Rural Uganda: A Cross-Sectional Study. <i>PLoS Medicine</i> , 2014, 11, e1001683.	8.4	53
25	Genetic characterization of Greek population isolates reveals strong genetic drift at missense and trait-associated variants. <i>Nature Communications</i> , 2014, 5, 5345.	12.8	60
26	Response to Comment on Ye et al. The Association Between Circulating Lipoprotein(a) and Type 2 Diabetes: Is It Causal? <i>Diabetes</i> 2014;63:332-342. <i>Diabetes</i> , 2014, 63, e15-e15.	0.6	3
27	Lipoprotein(a) Levels, Genotype, and Incident Aortic Valve Stenosis. <i>Circulation: Cardiovascular Genetics</i> , 2014, 7, 304-310.	5.1	219
28	Open-source electronic data capture system offered increased accuracy and cost-effectiveness compared with paper methods in Africa. <i>Journal of Clinical Epidemiology</i> , 2014, 67, 1358-1363.	5.0	49
29	The Association Between Circulating Lipoprotein(a) and Type 2 Diabetes: Is It Causal?. <i>Diabetes</i> , 2014, 63, 332-342.	0.6	82
30	Burden and Predictors of HIV /Hepatitis B Co-infection in Rural Uganda. <i>AIDS Research and Human Retroviruses</i> , 2014, 30, A276-A277.	1.1	0
31	Enabling the genomic revolution in Africa. <i>Science</i> , 2014, 344, 1346-1348.	12.6	361
32	Discovery and refinement of loci associated with lipid levels. <i>Nature Genetics</i> , 2013, 45, 1274-1283.	21.4	2,641
33	The general population cohort in rural south-western Uganda: a platform for communicable and non-communicable disease studies. <i>International Journal of Epidemiology</i> , 2013, 42, 129-141.	1.9	131
34	Effect modification by population dietary folate on the association between MTHFR genotype, homocysteine, and stroke risk: a meta-analysis of genetic studies and randomised trials. <i>Lancet</i> , The, 2011, 378, 584-594.	18.7	273
35	Biological, clinical and population relevance of 95 loci for blood lipids. <i>Nature</i> , 2010, 466, 707-713.	27.8	3,249
36	Association analyses of 249,796 individuals reveal 18 new loci associated with body mass index. <i>Nature Genetics</i> , 2010, 42, 937-948.	21.4	2,634

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37	Genomic risk prediction. <i>Lancet, The</i> , 2010, 376, 1366-1367.	13.7	7
38	LDL-cholesterol concentrations: a genome-wide association study. <i>Lancet, The</i> , 2008, 371, 483-491.	13.7	329
39	Abstract 3132: Paraoxonase-1 Activity Is not Independently Related with the Risk of Future Coronary Artery Disease. <i>Circulation</i> , 2008, 118, .	1.6	0
40	Insulin-Like Growth Factor-I and Risk of Type 2 Diabetes and Coronary Heart Disease: Molecular Epidemiology. , 2005, 9, 44-54.		34
41	INS VNTR Class Genotype and Indexes of Body Size and Obesity: Population-Based Studies of 7,999 Middle-Aged Men and Women. <i>Diabetes</i> , 2005, 54, 2812-2815.	0.6	14
42	Association between Insulin-Like Growth Factor-I: Insulin-Like Growth Factor-Binding Protein-1 Ratio and Metabolic and Anthropometric Factors in Men and Women. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2004, 13, 166-170.	2.5	49
43	Low Circulating IGF-II Concentrations Predict Weight Gain and Obesity in Humans. <i>Diabetes</i> , 2003, 52, 1403-1408.	0.6	86
44	Insulin, Insulin-Like Growth Factor-I (IGF-I), IGF Binding Proteins, Their Biologic Interactions, and Colorectal Cancer. <i>Journal of the National Cancer Institute</i> , 2002, 94, 972-980.	6.3	406
45	Circulating concentrations of insulin-like growth factor-I and development of glucose intolerance: a prospective observational study. <i>Lancet, The</i> , 2002, 359, 1740-1745.	13.7	438