

George Hindy

List of Publications by Year in descending order

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Version: 2024-02-01

33
papers

4,538
citations

331259

21
h-index

476904

29
g-index

36
all docs

36
docs citations

36
times ranked

8371
citing authors

#	ARTICLE	IF	CITATIONS
1	The Odds Ratio is “portable” across baseline risk but not the Relative Risk: Time to do away with the log link in binomial regression. <i>Journal of Clinical Epidemiology</i> , 2022, 142, 288-293.	2.4	19
2	The prevalence of adaptive immunity to COVID-19 and reinfection after recovery “ a comprehensive systematic review and meta-analysis. <i>Pathogens and Global Health</i> , 2022, 116, 269-281.	1.0	27
3	Rare coding variants in 35 genes associate with circulating lipid levels” A multi-ancestry analysis of 170,000 exomes. <i>American Journal of Human Genetics</i> , 2022, 109, 81-96.	2.6	24
4	Association of Habitual Alcohol Intake With Risk of Cardiovascular Disease. <i>JAMA Network Open</i> , 2022, 5, e223849.	2.8	136
5	Transethnic Transferability of a Genome-Wide Polygenic Score for Coronary Artery Disease. <i>Circulation Genomic and Precision Medicine</i> , 2021, 14, e003092.	1.6	25
6	Methodological considerations for identifying multiple plasma proteins associated with all-cause mortality in a population-based prospective cohort. <i>Scientific Reports</i> , 2021, 11, 6734.	1.6	2
7	Efficacy of chloroquine and hydroxychloroquine in treating COVID-19 infection: A meta-review of systematic reviews and an updated meta-analysis. <i>Travel Medicine and Infectious Disease</i> , 2021, 43, 102135.	1.5	16
8	Genome-Wide Polygenic Score, Clinical Risk Factors, and Long-Term Trajectories of Coronary Artery Disease. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2020, 40, 2738-2746.	1.1	71
9	Dynamic incorporation of multiple in silico functional annotations empowers rare variant association analysis of large whole-genome sequencing studies at scale. <i>Nature Genetics</i> , 2020, 52, 969-983.	9.4	146
10	Limitations of Contemporary Guidelines for Managing Patients at High Genetic Risk of Coronary Artery Disease. <i>Journal of the American College of Cardiology</i> , 2020, 75, 2769-2780.	1.2	88
11	A missense variant in Mitochondrial Amidoxime Reducing Component 1 gene and protection against liver disease. <i>PLoS Genetics</i> , 2020, 16, e1008629.	1.5	101
12	Blood pressure and bladder cancer risk in men by use of survival analysis and in interaction with NAT2 genotype, and by Mendelian randomization analysis. <i>PLoS ONE</i> , 2020, 15, e0241711.	1.1	4
13	Title is missing!. , 2020, 15, e0241711.		0
14	Title is missing!. , 2020, 15, e0241711.		0
15	Title is missing!. , 2020, 15, e0241711.		0
16	Title is missing!. , 2020, 15, e0241711.		0
17	Genetic Link Between Arterial Stiffness and Atrial Fibrillation. <i>Circulation Genomic and Precision Medicine</i> , 2019, 12, e002453.	1.6	11
18	Cardiometabolic Polygenic Risk Scores and Osteoarthritis Outcomes: A Mendelian Randomization Study Using Data From the MalmÅ Diet and Cancer Study and the <sc>UK</sc> Biobank. <i>Arthritis and Rheumatology</i> , 2019, 71, 925-934.	2.9	33

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19	Polygenic Risk Score for Coronary Heart Disease Modifies the Elevated Risk by Cigarette Smoking for Disease Incidence. <i>Circulation Genomic and Precision Medicine</i> , 2018, 11, e001856.	1.6	27
20	Role of Blood Lipids in the Development of Ischemic Stroke and its Subtypes. <i>Stroke</i> , 2018, 49, 820-827.	1.0	132
21	Dietary and genetic risk scores and incidence of type 2 diabetes. <i>Genes and Nutrition</i> , 2018, 13, 13.	1.2	32
22	Soluble Urokinase-type Plasminogen Activator Receptor (suPAR) and Impaired Kidney Function in the Population-based Malmö Diet and Cancer Study. <i>Kidney International Reports</i> , 2017, 2, 239-247.	0.4	33
23	Type 2 diabetes, glucose, insulin, BMI, and ischemic stroke subtypes. <i>Neurology</i> , 2017, 89, 454-460.	1.5	84
24	High Level of Fasting Plasma Proenkephalin-A Predicts Deterioration of Kidney Function and Incidence of CKD. <i>Journal of the American Society of Nephrology: JASN</i> , 2017, 28, 291-303.	3.0	29
25	A prospective study of dietary and supplemental zinc intake and risk of type 2 diabetes depending on genetic variation in SLC30A8. <i>Genes and Nutrition</i> , 2017, 12, 30.	1.2	26
26	Novel genetic loci associated with long-term deterioration in blood lipid concentrations and coronary artery disease in European adults. <i>International Journal of Epidemiology</i> , 2016, 46, dyw245.	0.9	17
27	Several type 2 diabetes-associated variants in genes annotated to WNT signaling interact with dietary fiber in relation to incidence of type 2 diabetes. <i>Genes and Nutrition</i> , 2016, 11, 6.	1.2	25
28	Using genetics to test the causal relationship of total adiposity and periodontitis: Mendelian randomization analyses in the Gene-Lifestyle Interactions and Dental Endpoints (GLIDE) Consortium. <i>International Journal of Epidemiology</i> , 2015, 44, 638-650.	0.9	54
29	Smoking Modifies the Associated Increased Risk of Future Cardiovascular Disease by Genetic Variation on Chromosome 9p21. <i>PLoS ONE</i> , 2014, 9, e85893.	1.1	24
30	Association of Low-Density Lipoprotein Cholesterol-Related Genetic Variants With Aortic Valve Calcium and Incident Aortic Stenosis. <i>JAMA - Journal of the American Medical Association</i> , 2014, 312, 1764.	3.8	184
31	Association of Low-Frequency and Rare Coding-Sequence Variants with Blood Lipids and Coronary Heart Disease in 56,000 Whites and Blacks. <i>American Journal of Human Genetics</i> , 2014, 94, 223-232.	2.6	287
32	Loss-of-Function Mutations in <i>APOC3</i> , Triglycerides, and Coronary Disease. <i>New England Journal of Medicine</i> , 2014, 371, 22-31.	13.9	936
33	Plasma HDL cholesterol and risk of myocardial infarction: a mendelian randomisation study. <i>Lancet</i> , 2012, 380, 572-580.	6.3	1,937