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List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8579636/publications.pdf>

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25
papers

2,005
citations

516561

16
h-index

610775

24
g-index

25
all docs

25
docs citations

25
times ranked

5311
citing authors

#	ARTICLE	IF	CITATIONS
1	Genomic insights in ascending aortic size and distensibility. <i>EBioMedicine</i> , 2022, 75, 103783.	2.7	22
2	Twenty-Five Novel Loci for Carotid Intima-Media Thickness: A Genome-Wide Association Study in >45â€™%000 Individuals and Meta-Analysis of >100â€™%000 Individuals. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2022, 42, 484-501.	1.1	17
3	Genome-Wide Association Study and Identification of a Protective Missense Variant on Lipoprotein(a) Concentration. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2021, 41, 1792-1800.	1.1	29
4	Search for a Functional Genetic Variant Mimicking the Effect of SGLT2 Inhibitor Treatment. <i>Genes</i> , 2021, 12, 1174.	1.0	3
5	Polygenic risk score and coronary artery disease: A meta-analysis of 979,286 participant data. <i>Atherosclerosis</i> , 2021, 333, 48-55.	0.4	18
6	An Erythropoietin-Independent Mechanism of Erythrocytic Precursor Proliferation Underlies Hypoxia Tolerance in Sea Nomads. <i>Frontiers in Physiology</i> , 2021, 12, 760851.	1.3	0
7	Associations of Observational and Genetically Determined Caffeine Intake With Coronary Artery Disease and Diabetes Mellitus. <i>Journal of the American Heart Association</i> , 2020, 9, e016808.	1.6	21
8	Multi-ancestry GWAS of the electrocardiographic PR interval identifies 202 loci underlying cardiac conduction. <i>Nature Communications</i> , 2020, 11, 2542.	5.8	59
9	Genetically Determined High Levels of Iron Parameters Are Protective for Coronary Artery Disease. <i>Circulation Genomic and Precision Medicine</i> , 2020, 13, e002544.	1.6	2
10	Genetically Determined ABO Blood Group and its Associations With Health and Disease. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2020, 40, 830-838.	1.1	90
11	Genome-wide association studies and Mendelian randomization analyses for leisure sedentary behaviours. <i>Nature Communications</i> , 2020, 11, 1770.	5.8	66
12	Lipidomics, Atrial Conduction, and Body Mass Index. <i>Circulation Genomic and Precision Medicine</i> , 2019, 12, e002384.	1.6	9
13	Contributions of Interactions Between Lifestyle and Genetics on Coronary Artery Disease Risk. <i>Current Cardiology Reports</i> , 2019, 21, 89.	1.3	27
14	Effects of Calcium, Magnesium, and Potassium Concentrations on Ventricular Repolarization in Unselected Individuals. <i>Journal of the American College of Cardiology</i> , 2019, 73, 3118-3131.	1.2	27
15	Phenome-wide association analysis of LDL-cholesterol lowering genetic variants in PCSK9. <i>BMC Cardiovascular Disorders</i> , 2019, 19, 240.	0.7	22
16	Genetically Determined Physical Activity and Its Association with Circulating Blood Cells. <i>Genes</i> , 2019, 10, 908.	1.0	4
17	Effect of Systolic Blood Pressure on Left Ventricular Structure and Function. <i>Hypertension</i> , 2019, 74, 826-832.	1.3	23
18	Multiancestry Genome-Wide Association Study of Lipid Levels Incorporating Gene-Alcohol Interactions. <i>American Journal of Epidemiology</i> , 2019, 188, 1033-1054.	1.6	85

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19	Smoking does not accelerate leucocyte telomere attrition: a meta-analysis of 18 longitudinal cohorts. Royal Society Open Science, 2019, 6, 190420.	1.1	33
20	Multi-ancestry genome-wide gene-smoking interaction study of 387,272 individuals identifies new loci associated with serum lipids. Nature Genetics, 2019, 51, 636-648.	9.4	112
21	Relationship of Arterial Stiffness Index and Pulse Pressure With Cardiovascular Disease and Mortality. Journal of the American Heart Association, 2018, 7, .	1.6	142
22	Associations of Combined Genetic and Lifestyle Risks With Incident Cardiovascular Disease and Diabetes in the UK Biobank Study. JAMA Cardiology, 2018, 3, 693.	3.0	310
23	Novel genetic associations for blood pressure identified via gene-alcohol interaction in up to 570K individuals across multiple ancestries. PLoS ONE, 2018, 13, e0198166.	1.1	94
24	Genome-wide association analysis identifies novel blood pressure loci and offers biological insights into cardiovascular risk. Nature Genetics, 2017, 49, 403-415.	9.4	492
25	PCSK9 genetic variants and risk of type 2 diabetes: a mendelian randomisation study. Lancet Diabetes and Endocrinology, the, 2017, 5, 97-105.	5.5	298