

# M Abdullah Said

## List of Publications by Year in descending order

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

Version: 2024-02-01

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	Genome-wide association analysis identifies novel blood pressure loci and offers biological insights into cardiovascular risk. <i>Nature Genetics</i> , 2017, 49, 403-415.	9.4	492
2	Associations of Combined Genetic and Lifestyle Risks With Incident Cardiovascular Disease and Diabetes in the UK Biobank Study. <i>JAMA Cardiology</i> , 2018, 3, 693.	3.0	310
3	PCSK9 genetic variants and risk of type 2 diabetes: a mendelian randomisation study. <i>Lancet Diabetes and Endocrinology</i> , 2017, 5, 97-105.	5.5	298
4	Relationship of Arterial Stiffness Index and Pulse Pressure With Cardiovascular Disease and Mortality. <i>Journal of the American Heart Association</i> , 2018, 7, .	1.6	142
5	Multi-ancestry genome-wide gene-smoking interaction study of 387,272 individuals identifies new loci associated with serum lipids. <i>Nature Genetics</i> , 2019, 51, 636-648.	9.4	112
6	Novel genetic associations for blood pressure identified via gene-alcohol interaction in up to 570K individuals across multiple ancestries. <i>PLoS ONE</i> , 2018, 13, e0198166.	1.1	94
7	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
8	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
9	Genome-wide association studies and Mendelian randomization analyses for leisure sedentary behaviours. <i>Nature Communications</i> , 2020, 11, 1770.	5.8	66
10	Multi-ancestry GWAS of the electrocardiographic PR interval identifies 202 loci underlying cardiac conduction. <i>Nature Communications</i> , 2020, 11, 2542.	5.8	59
11	Smoking does not accelerate leucocyte telomere attrition: a meta-analysis of 18 longitudinal cohorts. <i>Royal Society Open Science</i> , 2019, 6, 190420.	1.1	33
12	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
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	Effect of Systolic Blood Pressure on Left Ventricular Structure and Function. <i>Hypertension</i> , 2019, 74, 826-832.	1.3	23
16	Phenome-wide association analysis of LDL-cholesterol lowering genetic variants in PCSK9. <i>BMC Cardiovascular Disorders</i> , 2019, 19, 240.	0.7	22
17	Genomic insights in ascending aortic size and distensibility. <i>EBioMedicine</i> , 2022, 75, 103783.	2.7	22
18	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

#	ARTICLE	IF	CITATIONS
19	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
20	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
21	Lipidomics, Atrial Conduction, and Body Mass Index. <i>Circulation Genomic and Precision Medicine</i> , 2019, 12, e002384.	1.6	9
22	Genetically Determined Physical Activity and Its Association with Circulating Blood Cells. <i>Genes</i> , 2019, 10, 908.	1.0	4
23	Search for a Functional Genetic Variant Mimicking the Effect of SGLT2 Inhibitor Treatment. <i>Genes</i> , 2021, 12, 1174.	1.0	3
24	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
25	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