

Sanni E Ruotsalainen

List of Publications by Year in descending order

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Version: 2024-02-01

17
papers

1,057
citations

858243

12
h-index

939365

18
g-index

24
all docs

24
docs citations

24
times ranked

1551
citing authors

#	ARTICLE	IF	CITATIONS
1	The power of genetic diversity in genome-wide association studies of lipids. <i>Nature</i> , 2021, 600, 675-679.	13.7	353
2	Genome-wide analysis of 102,084 migraine cases identifies 123 risk loci and subtype-specific risk alleles. <i>Nature Genetics</i> , 2022, 54, 152-160.	9.4	135
3	Sleep apnoea is a risk factor for severe COVID-19. <i>BMJ Open Respiratory Research</i> , 2021, 8, e000845.	1.2	92
4	Geographic Variation and Bias in the Polygenic Scores of Complex Diseases and Traits in Finland. <i>American Journal of Human Genetics</i> , 2019, 104, 1169-1181.	2.6	90
5	Genetic analysis of obstructive sleep apnoea discovers a strong association with cardiometabolic health. <i>European Respiratory Journal</i> , 2021, 57, 2003091.	3.1	85
6	Deep coverage whole genome sequences and plasma lipoprotein(a) in individuals of European and African ancestries. <i>Nature Communications</i> , 2018, 9, 2606.	5.8	79
7	How Communicating Polygenic and Clinical Risk for Atherosclerotic Cardiovascular Disease Impacts Health Behavior: an Observational Follow-up Study. <i>Circulation Genomic and Precision Medicine</i> , 2022, 15, CIRCGEN121003459.	1.6	53
8	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
9	High heritability of ascending aortic diameter and trans-ancestry prediction of thoracic aortic disease. <i>Nature Genetics</i> , 2022, 54, 772-782.	9.4	29
10	Genomic prediction of alcohol-related morbidity and mortality. <i>Translational Psychiatry</i> , 2020, 10, 23.	2.4	19
11	An expanded analysis framework for multivariate GWAS connects inflammatory biomarkers to functional variants and disease. <i>European Journal of Human Genetics</i> , 2021, 29, 309-324.	1.4	19
12	Chromosome Xq23 is associated with lower atherogenic lipid concentrations and favorable cardiometabolic indices. <i>Nature Communications</i> , 2021, 12, 2182.	5.8	17
13	Moyamoya angiopathy: long-term follow-up study in a Finnish population. <i>Journal of Neurology</i> , 2019, 266, 574-581.	1.8	15
14	Human essential hypertension: no significant association of polygenic risk scores with antihypertensive drug responses. <i>Scientific Reports</i> , 2020, 10, 11940.	1.6	11
15	Polygenic burden has broader impact on health, cognition, and socioeconomic outcomes than most rare and high-risk copy number variants. <i>Molecular Psychiatry</i> , 2021, 26, 4884-4895.	4.1	8
16	Genetic Risk Score for Serum 25-Hydroxyvitamin D Concentration Helps to Guide Personalized Vitamin D Supplementation in Healthy Finnish Adults. <i>Journal of Nutrition</i> , 2021, 151, 281-292.	1.3	8
17	Chromosomal Region 11p14.1 is Associated with Pharmacokinetics and Pharmacodynamics of Bisoprolol. <i>Pharmacogenomics and Personalized Medicine</i> , 2022, Volume 15, 249-260.	0.4	1