Jun Liu

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/137911/publications.pdf

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		566801	676716
28	9,247	15	22
papers	citations	h-index	g-index
38	38	38	23713
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	Fat metabolism is associated with telomere length in six population-based studies. Human Molecular Genetics, 2022, 31, 1159-1170.	1.4	7
2	A multi-omics study of circulating phospholipid markers of blood pressure. Scientific Reports, 2022, 12, 574.	1.6	10
3	Mendelian randomization analyses for PCOS: evidence, opportunities, and challenges. Trends in Genetics, 2022, 38, 468-482.	2.9	21
4	Multi-ancestry genome-wide association study of gestational diabetes mellitus highlights genetic links with type 2 diabetes. Human Molecular Genetics, 2022, 31, 3377-3391.	1.4	47
5	Breast cancer risk in patients with polycystic ovary syndrome: a Mendelian randomization analysis. Breast Cancer Research and Treatment, 2021, 185, 799-806.	1.1	19
6	The trans-ancestral genomic architecture of glycemic traits. Nature Genetics, 2021, 53, 840-860.	9.4	341
7	Identification of Biomarkers for the Prevention of Chronic Disease. SpringerBriefs in Public Health, 2021, , 9-32.	0.2	1
8	The power of genetic diversity in genome-wide association studies of lipids. Nature, 2021, 600, 675-679.	13.7	353
9	Profiling the metabolome of patients with dementia in the UK Biobank. Alzheimer's and Dementia, 2021, 17, .	0.4	O
10	Ethnicity influences risk of dementia in the UK Biobank. Alzheimer's and Dementia, 2021, 17 , .	0.4	1
11	Lipidomic profiling identifies signatures of metabolic risk. EBioMedicine, 2020, 51, 102520.	2.7	56
12	Smoking-by-genotype interaction in type 2 diabetes risk and fasting glucose. PLoS ONE, 2020, 15, e0230815.	1.1	10
13	Integration of epidemiologic, pharmacologic, genetic and gut microbiome data in a drug–metabolite atlas. Nature Medicine, 2020, 26, 110-117.	15.2	54
14	Smoking-by-genotype interaction in type 2 diabetes risk and fasting glucose., 2020, 15, e0230815.		0
15	Smoking-by-genotype interaction in type 2 diabetes risk and fasting glucose., 2020, 15, e0230815.		O
16	Smoking-by-genotype interaction in type 2 diabetes risk and fasting glucose., 2020, 15, e0230815.		0
17	Smoking-by-genotype interaction in type 2 diabetes risk and fasting glucose., 2020, 15, e0230815.		0
18	Target genes, variants, tissues and transcriptional pathways influencing human serum urate levels. Nature Genetics, 2019, 51, 1459-1474.	9.4	251

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19	A catalog of genetic loci associated with kidney function from analyses of a million individuals. Nature Genetics, 2019, 51, 957-972.	9.4	549
20	An integrative cross-omics analysis of DNA methylation sites of glucose and insulin homeostasis. Nature Communications, 2019, 10, 2581.	5.8	62
21	Large-scale plasma metabolome analysis reveals alterations in HDL metabolism in migraine. Neurology, 2019, 92, e1899-e1911.	1.5	42
22	Refining the accuracy of validated target identification through coding variant fine-mapping in type 2 diabetes. Nature Genetics, 2018, 50, 559-571.	9.4	356
23	Large-scale whole-exome sequencing association studies identify rare functional variants influencing serum urate levels. Nature Communications, 2018, 9, 4228.	5.8	43
24	Carotid intima-media thickness and plagues are associated with indicators of peripheral artery diseases in patients with diabetes. Diabetes Research and Clinical Practice, 2018, 144, 245-251.	1.1	3
25	A Mendelian Randomization Study of Metabolite Profiles, Fasting Glucose, and Type 2 Diabetes. Diabetes, 2017, 66, 2915-2926.	0.3	40
26	Metabolomics based markers predict type 2 diabetes in a 14-year follow-up study. Metabolomics, 2017, 13, 104.	1.4	82
27	Association between fat mass and obesity associated (FTO) gene rs9939609 A/T polymorphism and polycystic ovary syndrome: a systematic review and meta-analysis. BMC Medical Genetics, 2017, 18, 89.	2.1	63
28	The Genotype-Tissue Expression (GTEx) project. Nature Genetics, 2013, 45, 580-585.	9.4	6,815