

Diptavo Dutta

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5570860/publications.pdf>

Version: 2024-02-01

11
papers

274
citations

1478505

6
h-index

1372567

10
g-index

17
all docs

17
docs citations

17
times ranked

419
citing authors

#	ARTICLE	IF	CITATIONS
1	Genome-wide association studies of 27 accelerometry-derived physical activity measurements identified novel loci and genetic mechanisms. <i>Genetic Epidemiology</i> , 2022, 46, 122-138.	1.3	7
2	Trans-ethnic genome-wide association study of blood metabolites in the Chronic Renal Insufficiency Cohort (CRIC) study. <i>Kidney International</i> , 2022, 101, 814-823.	5.2	8
3	Rare Genetic Variants Associated With Myocardial Fibrosis: Multi-Ethnic Study of Atherosclerosis. <i>Frontiers in Cardiovascular Medicine</i> , 2022, 9, 804788.	2.4	6
4	Plasma proteome analyses in individuals of European and African ancestry identify cis-pQTLs and models for proteome-wide association studies. <i>Nature Genetics</i> , 2022, 54, 593-602.	21.4	98
5	A powerful subset-based method identifies gene set associations and improves interpretation in UK Biobank. <i>American Journal of Human Genetics</i> , 2021, 108, 669-681.	6.2	8
6	Genome-Wide Association Study of Pelvic Organ Prolapse Using the Michigan Genomics Initiative. <i>Female Pelvic Medicine and Reconstructive Surgery</i> , 2021, 27, 502-506.	1.1	0
7	Proteins Associated with Risk of Kidney Function Decline in the General Population. <i>Journal of the American Society of Nephrology: JASN</i> , 2021, 32, 2291-2302.	6.1	23
8	Heritability of the Fibromyalgia Phenotype Varies by Age. <i>Arthritis and Rheumatology</i> , 2020, 72, 815-823.	5.6	15
9	Genome-wide association and multi-omic analyses reveal ACTN2 as a gene linked to heart failure. <i>Nature Communications</i> , 2020, 11, 1122.	12.8	57
10	Meta-MultiSKAT: Multiple phenotype meta-analysis for region-based association test. <i>Genetic Epidemiology</i> , 2019, 43, 800-814.	1.3	9
11	MultiSKAT: General framework to test for rare-variant association with multiple phenotypes. <i>Genetic Epidemiology</i> , 2019, 43, 4-23.	1.3	40