

Fabiola Del Greco M

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/10404448/publications.pdf>

Version: 2024-02-01

24
papers

5,904
citations

430874

18
h-index

610901

24
g-index

28
all docs

28
docs citations

28
times ranked

7032
citing authors

#	ARTICLE	IF	CITATIONS
1	A framework for the investigation of pleiotropy in two-sample summary data Mendelian randomization. <i>Statistics in Medicine</i> , 2017, 36, 1783-1802.	1.6	975
2	Genetic analysis of over 1 million people identifies 535 new loci associated with blood pressure traits. <i>Nature Genetics</i> , 2018, 50, 1412-1425.	21.4	924
3	Assessing the suitability of summary data for two-sample Mendelian randomization analyses using MR-Egger regression: the role of the I ² statistic. <i>International Journal of Epidemiology</i> , 2016, 45, dyw220.	1.9	787
4	Detecting pleiotropy in Mendelian randomisation studies with summary data and a continuous outcome. <i>Statistics in Medicine</i> , 2015, 34, 2926-2940.	1.6	671
5	Mendelian Randomization as an Approach to Assess Causality Using Observational Data. <i>Journal of the American Society of Nephrology: JASN</i> , 2016, 27, 3253-3265.	6.1	639
6	Genome-wide association study identifies six new loci influencing pulse pressure and mean arterial pressure. <i>Nature Genetics</i> , 2011, 43, 1005-1011.	21.4	403
7	Improving the visualization, interpretation and analysis of two-sample summary data Mendelian randomization via the Radial plot and Radial regression. <i>International Journal of Epidemiology</i> , 2018, 47, 1264-1278.	1.9	389
8	Improving the accuracy of two-sample summary-data Mendelian randomization: moving beyond the NOME assumption. <i>International Journal of Epidemiology</i> , 2019, 48, 728-742.	1.9	346
9	The use of two-sample methods for Mendelian randomization analyses on single large datasets. <i>International Journal of Epidemiology</i> , 2021, 50, 1651-1659.	1.9	150
10	Novel Blood Pressure Locus and Gene Discovery Using Genome-Wide Association Study and Expression Data Sets From Blood and the Kidney. <i>Hypertension</i> , 2017, 70, .	2.7	123
11	The Effect of Iron Status on Risk of Coronary Artery Disease. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2017, 37, 1788-1792.	2.4	72
12	Age at menarche and adult body mass index: a Mendelian randomization study. <i>International Journal of Obesity</i> , 2018, 42, 1574-1581.	3.4	68
13	Mendelian Randomization. <i>Methods in Molecular Biology</i> , 2017, 1666, 581-628.	0.9	65
14	Mendelian Randomization using Public Data from Genetic Consortia. <i>International Journal of Biostatistics</i> , 2016, 12, .	0.7	59
15	Genome-wide association analysis and fine mapping of NT-proBNP level provide novel insight into the role of the MTHFR-CLCN6-NPPA-NPPB gene cluster. <i>Human Molecular Genetics</i> , 2011, 20, 1660-1671.	2.9	47
16	Thyroid Function Affects the Risk of Stroke via Atrial Fibrillation: A Mendelian Randomization Study. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2020, 105, 2634-2641.	3.6	31
17	Variation in Normal Range Thyroid Function Affects Serum Cholesterol Levels, Blood Pressure, and Type 2 Diabetes Risk: A Mendelian Randomization Study. <i>Thyroid</i> , 2021, 31, 721-731.	4.5	31
18	Serum iron level and kidney function: a Mendelian randomization study. <i>Nephrology Dialysis Transplantation</i> , 2016, 32, gfw215.	0.7	23

#	ARTICLE	IF	CITATIONS
19	Thyroid Function and Mood Disorders: A Mendelian Randomization Study. <i>Thyroid</i> , 2021, 31, 1171-1181.	4.5	23
20	Risky behaviors and Parkinson disease. <i>Neurology</i> , 2019, 93, e1412-e1424.	1.1	18
21	Evaluating the current state of Mendelian randomization studies: a protocol for a systematic review on methodological and clinical aspects using neurodegenerative disorders as outcome. <i>Systematic Reviews</i> , 2018, 7, 145.	5.3	16
22	A multi-omics study of circulating phospholipid markers of blood pressure. <i>Scientific Reports</i> , 2022, 12, 574.	3.3	10
23	Whole Exome Sequencing Enhanced Imputation Identifies 85 Metabolite Associations in the Alpine CHRIS Cohort. <i>Metabolites</i> , 2022, 12, 604.	2.9	6
24	Bayesian analysis of censored response data in family-based genetic association studies. <i>Biometrical Journal</i> , 2016, 58, 1039-1053.	1.0	5