

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 multi-omics study of circulating phospholipid markers of blood pressure. <i>Scientific Reports</i> , 2022, 12, 574. | 3.3 | 10 |
| 2 | Whole Exome Sequencing Enhanced Imputation Identifies 85 Metabolite Associations in the Alpine CHRIS Cohort. <i>Metabolites</i> , 2022, 12, 604. | 2.9 | 6 |
| 3 | 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 |
| 4 | 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 |
| 5 | Thyroid Function and Mood Disorders: A Mendelian Randomization Study. <i>Thyroid</i> , 2021, 31, 1171-1181. | 4.5 | 23 |
| 6 | 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 |
| 7 | Risky behaviors and Parkinson disease. <i>Neurology</i> , 2019, 93, e1412-e1424. | 1.1 | 18 |
| 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 | 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 |
| 10 | 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 |
| 11 | 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 |
| 12 | 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 |
| 13 | 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 |
| 14 | Mendelian Randomization. <i>Methods in Molecular Biology</i> , 2017, 1666, 581-628. | 0.9 | 65 |
| 15 | 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 |
| 16 | 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 |
| 17 | Mendelian Randomization using Public Data from Genetic Consortia. <i>International Journal of Biostatistics</i> , 2016, 12, . | 0.7 | 59 |
| 18 | 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 |

| # | ARTICLE | IF | CITATIONS |
|----|---|------|-----------|
| 19 | 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 |
| 20 | Serum iron level and kidney function: a Mendelian randomization study. <i>Nephrology Dialysis Transplantation</i> , 2016, 32, gfw215. | 0.7 | 23 |
| 21 | Bayesian analysis of censored response data in family-based genetic association studies. <i>Biometrical Journal</i> , 2016, 58, 1039-1053. | 1.0 | 5 |
| 22 | 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 |
| 23 | 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 |
| 24 | 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 |