## Fernando P Hartwig

List of Publications by Year in descending order

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78 papers

7,173 citations

147726 31 h-index 71 g-index

95 all docs 95
docs citations

95 times ranked 10297 citing authors

#	Article	IF	CITATIONS
1	Robust inference in summary data Mendelian randomization via the zero modal pleiotropy assumption. International Journal of Epidemiology, 2017, 46, 1985-1998.	0.9	1,407
2	Guidelines for performing Mendelian randomization investigations. Wellcome Open Research, 2019, 4, 186.	0.9	661
3	Guidelines for performing Mendelian randomization investigations. Wellcome Open Research, 2019, 4, 186.	0.9	511
4	Two-sample Mendelian randomization: avoiding the downsides of a powerful, widely applicable but potentially fallible technique. International Journal of Epidemiology, 2016, 45, 1717-1726.	0.9	458
5	The trans-ancestral genomic architecture of glycemic traits. Nature Genetics, 2021, 53, 840-860.	9.4	341
6	PCSK9 genetic variants and risk of type 2 diabetes: a mendelian randomisation study. Lancet Diabetes and Endocrinology,the, 2017, 5, 97-105.	5.5	298
7	SARS-CoV-2 antibody prevalence in Brazil: results from two successive nationwide serological household surveys. The Lancet Global Health, 2020, 8, e1390-e1398.	2.9	292
8	Life-Course Genome-wide Association Study Meta-analysis of Total Body BMD and Assessment of Age-Specific Effects. American Journal of Human Genetics, 2018, 102, 88-102.	2.6	252
9	Genome-wide analyses using UK Biobank data provide insights into the genetic architecture of osteoarthritis. Nature Genetics, 2018, 50, 549-558.	9.4	223
10	Avoiding dynastic, assortative mating, and population stratification biases in Mendelian randomization through within-family analyses. Nature Communications, 2020, 11, 3519.	5.8	213
11	Inflammatory Biomarkers and Risk of Schizophrenia. JAMA Psychiatry, 2017, 74, 1226.	6.0	204
12	Education and coronary heart disease: mendelian randomisation study. BMJ: British Medical Journal, 2017, 358, j3542.	2.4	191
13	Population-based surveys of antibodies against SARS-CoV-2 in Southern Brazil. Nature Medicine, 2020, 26, 1196-1199.	15.2	132
14	A Large-Scale Multi-ancestry Genome-wide Study Accounting for Smoking Behavior Identifies Multiple Significant Loci for Blood Pressure. American Journal of Human Genetics, 2018, 102, 375-400.	2.6	123
15	Multi-ancestry genome-wide gene–smoking interaction study of 387,272 individuals identifies new loci associated with serum lipids. Nature Genetics, 2019, 51, 636-648.	9.4	112
16	Associations of Mitochondrial and Nuclear Mitochondrial Variants and Genes with Seven Metabolic Traits. American Journal of Human Genetics, 2019, 104, 112-138.	2.6	106
17	Novel genetic associations for blood pressure identified via gene-alcohol interaction in up to 570K individuals across multiple ancestries. PLoS ONE, 2018, 13, e0198166.	1.1	94
18	Multiethnic meta-analysis identifies ancestry-specific and cross-ancestry loci for pulmonary function. Nature Communications, 2018, 9, 2976.	5.8	85

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19	Multiancestry Genome-Wide Association Study of Lipid Levels Incorporating Gene-Alcohol Interactions. American Journal of Epidemiology, 2019, 188, 1033-1054.	1.6	85
20	Bias in Mendelian randomization due to assortative mating. Genetic Epidemiology, 2018, 42, 608-620.	0.6	81
21	Bias in two-sample Mendelian randomization when using heritable covariable-adjusted summary associations. International Journal of Epidemiology, 2021, 50, 1639-1650.	0.9	65
22	Multi-ancestry study of blood lipid levels identifies four loci interacting with physical activity. Nature Communications, 2019, 10, 376.	5 <b>.</b> 8	64
23	Efficacy of Regular Exercise During Pregnancy on the Prevention of Postpartum Depression. JAMA Network Open, 2019, 2, e186861.	2.8	52
24	Breastfeeding effects on DNA methylation in the offspring: A systematic literature review. PLoS ONE, 2017, 12, e0173070.	1.1	52
25	Body mass index and psychiatric disorders: a Mendelian randomization study. Scientific Reports, 2016, 6, 32730.	1.6	47
26	Assessing causality in the association between attention-deficit/hyperactivity disorder and obesity: a Mendelian randomization study. International Journal of Obesity, 2019, 43, 2500-2508.	1.6	45
27	Missed childhood immunizations during the COVID-19 pandemic in Brazil: Analyses of routine statistics and of a national household survey. Vaccine, 2021, 39, 3404-3409.	1.7	43
28	African ancestry, lung function and the effect of genetics. European Respiratory Journal, 2015, 45, 1582-1589.	3.1	39
29	Why internal weights should be avoided (not only) in MR-Egger regression. International Journal of Epidemiology, 2016, 45, 1676-1678.	0.9	37
30	A Genome-Wide Association Study in Hispanics/Latinos Identifies Novel Signals for Lung Function. The Hispanic Community Health Study/Study of Latinos. American Journal of Respiratory and Critical Care Medicine, 2018, 198, 208-219.	2.5	37
31	Prevalence of antibodies against SARS-CoV-2 according to socioeconomic and ethnic status in a nationwide Brazilian survey. Revista Panamericana De Salud Publica/Pan American Journal of Public Health, 2020, 44, 1-7.	0.6	37
32	Effects of early-life poverty on health and human capital in children and adolescents: analyses of national surveys and birth cohort studies in LMICs. Lancet, The, 2022, 399, 1741-1752.	6.3	37
33	Health and development from preconception to 20 years of age and human capital. Lancet, The, 2022, 399, 1730-1740.	6.3	37
34	From stem cells to the law courts: DNA methylation, the forensic epigenome and the possibility of a biosocial archive. International Journal of Epidemiology, 2015, 44, 1083-1093.	0.9	33
35	Association of lactase persistence genotype with milk consumption, obesity and blood pressure: a Mendelian randomization study in the 1982 Pelotas (Brazil) Birth Cohort, with a systematic review and meta-analysis. International Journal of Epidemiology, 2016, 45, 1573-1587.	0.9	31
36	A multi-ancestry genome-wide study incorporating gene–smoking interactions identifies multiple new loci for pulse pressure and mean arterial pressure. Human Molecular Genetics, 2019, 28, 2615-2633.	1.4	31

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37	Genome-Wide Association Study of Blood Pressure Traits by Hispanic/Latino Background: the Hispanic Community Health Study/Study of Latinos. Scientific Reports, 2017, 7, 10348.	1.6	24
38	Phenome-wide association analysis of LDL-cholesterol lowering genetic variants in PCSK9. BMC Cardiovascular Disorders, 2019, 19, 240.	0.7	22
39	Breastfeeding moderates FTO related adiposity: a birth cohort study with 30 years of follow-up. Scientific Reports, 2018, 8, 2530.	1.6	18
40	Telomeres and Tissue Engineering: The Potential Roles of TERT in VEGF-mediated Angiogenesis. Stem Cell Reviews and Reports, 2012, 8, 1275-1281.	5.6	17
41	Mendelian randomization evaluation of causal effects of fibrinogen on incident coronary heart disease. PLoS ONE, 2019, 14, e0216222.	1.1	17
42	Gene-educational attainment interactions in a multi-ancestry genome-wide meta-analysis identify novel blood pressure loci. Molecular Psychiatry, 2020, 26, 2111-2125.	4.1	17
43	EPICOVID19 protocol: repeated serological surveys on SARS-CoV-2 antibodies in Brazil. Ciencia E Saude Coletiva, 2020, 25, 3573-3578.	0.1	15
44	COVID-19 and outpatient care: a nationwide household survey. Cadernos De Saude Publica, 2022, 38, e00194121.	0.4	15
45	Suggestive association between variants in IL1RAPL and asthma symptoms in Latin American children. European Journal of Human Genetics, 2017, 25, 439-445.	1.4	14
46	The median and the mode as robust metaâ€analysis estimators in the presence of smallâ€study effects and outliers. Research Synthesis Methods, 2020, 11, 397-412.	4.2	14
47	Assortative mating and within-spouse pair comparisons. PLoS Genetics, 2021, 17, e1009883.	1.5	13
48	The Confidence Interval Method for Selecting Valid Instrumental Variables. Journal of the Royal Statistical Society Series B: Statistical Methodology, 2021, 83, 752-776.	1.1	12
49	Telomere dysfunction and tumor suppression responses in dyskeratosis congenita: Balancing cancer and tissue renewal impairment. Ageing Research Reviews, 2013, 12, 642-652.	5.0	11
50	Doenças crônicas não transmissÃveis e covid-19: resultados do estudo Epicovid-19 Brasil. Revista De Saude Publica, 2021, 55, 38.	0.7	11
51	Mendelian Randomization Concerns—Reply. JAMA Psychiatry, 2018, 75, 407.	6.0	10
52	Genome-wide burden and association analyses implicate copy number variations in asthma risk among children and young adults from Latin America. Scientific Reports, 2018, 8, 14475.	1.6	10
53	Geospatial estimation of reproductive, maternal, newborn and child health indicators: a systematic review of methodological aspects of studies based on household surveys. International Journal of Health Geographics, 2020, 19, 41.	1.2	10
54	Association between Breastfeeding and DNA Methylation over the Life Course: Findings from the Avon Longitudinal Study of Parents and Children (ALSPAC). Nutrients, 2020, 12, 3309.	1.7	10

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55	High prevalence of symptoms among Brazilian subjects with antibodies against SARS-CoV-2. Scientific Reports, 2021, 11, 13279.	1.6	10
56	Genomic ancestry and the social pathways leading to major depression in adulthood: the mediating effect of socioeconomic position and discrimination. BMC Psychiatry, 2016, 16, 308.	1.1	9
57	Mental disorders, comorbidities, and suicidality at 30†years of age in a Brazilian birth cohort. Comprehensive Psychiatry, 2020, 102, 152194.	1.5	9
58	Patterns of Growth in Childhood in Relation to Adult Schooling Attainment and Intelligence Quotient in 6 Birth Cohorts in Low- and Middle-Income Countries: Evidence from the Consortium of Health-Oriented Research in Transitioning Societies (COHORTS). Journal of Nutrition, 2021, 151, 2342-2352.	1.3	9
59	Time-dependent decay of detectable antibodies against SARS-CoV-2: A comparison of ELISA with two batches of a lateral-flow test. Brazilian Journal of Infectious Diseases, 2021, 25, 101601.	0.3	9
60	Breastfeeding and intelligence in adulthood: due to genetic confounding?. The Lancet Global Health, 2018, 6, e1276-e1277.	2.9	7
61	Population-level seropositivity trend for SARS-Cov-2 in Rio Grande do Sul, Brazil. Revista De Saude Publica, 2021, 55, 78.	0.7	7
62	Effect modification of <i>FADS2 </i> polymorphisms on the association between breastfeeding and intelligence: protocol for a collaborative meta-analysis. BMJ Open, 2016, 6, e010067.	0.8	6
63	Lactase Persistence and Body Mass Index: The Contribution of Mendelian Randomization. Clinical Chemistry, 2018, 64, 4-6.	1.5	6
64	Slow Spread of SARS-CoV-2 in Southern Brazil Over a 6-Month Period: Report on 8 Sequential Statewide Serological Surveys Including 35 611 Participants. American Journal of Public Health, 2021, 111, 1542-1550.	1.5	6
65	Influence of maternal pre-pregnancy nutritional status on offspring anthropometric measurements and body composition in three Brazilian Birth Cohorts. Public Health Nutrition, 2021, 24, 882-894.	1.1	6
66	Effect modification of <i>FADS2</i> polymorphisms on the association between breastfeeding and intelligence: results from a collaborative meta-analysis. International Journal of Epidemiology, 2019, 48, 45-57.	0.9	5
67	COVID-19 and social distancing among children and adolescents in Brazil. Revista De Saude Publica, 2021, 55, 42.	0.7	5
68	Uso de m $\tilde{A}_i$ scara durante a pandemia de COVID-19 no Brasil: resultados do estudo EPICOVID19-BR. Cadernos De Saude Publica, 2022, 38, .	0.4	5
69	The challenge of conducting epidemiological research in times of pandemic and denialism: 1-year anniversary of the EPICOVID-19 project in Brazil. International Journal of Epidemiology, 2021, 50, 1049-1052.	0.9	4
70	Evidence for an Epistatic Effect between TP53 R72P and MDM2 T309G SNPs in HIV Infection: A Cross-Sectional Study in Women from South Brazil. PLoS ONE, 2014, 9, e89489.	1.1	4
71	Effectiveness of a large-scale home visiting programme (PIM) on early child development in Brazil: quasi-experimental study nested in a birth cohort. BMJ Global Health, 2022, 7, e007116.	2.0	4
72	Up-Regulating Telomerase and Tumor Suppressors: Focusing on Anti-Aging Interventions at the Population Level. , 2014, 5, 17-26.		3

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73	Ancestral diversity improves discovery and fine-mapping of genetic loci for anthropometric traits—The Hispanic/Latino Anthropometry Consortium. Human Genetics and Genomics Advances, 2022, 3, 100099.	1.0	3
74	Multi-ancestry genome-wide association study accounting for gene-psychosocial factor interactions identifies novel loci for blood pressure traits. Human Genetics and Genomics Advances, 2021, 2, 100013.	1.0	2
75	Letter by Hartwig et al Regarding Article, "Evaluation of the Pleiotropic Effects of Statins: A Reanalysis of the Randomized Trial Evidence Using Egger Regression― Arteriosclerosis, Thrombosis, and Vascular Biology, 2018, 38, e85-e86.	1.1	1
76	Association between cesarean section and human capital in adulthood: 1982 and 1993 Pelotas birth cohorts, Rio Grande do Sul State, Brazil. Cadernos De Saude Publica, 2021, 37, e00235520.	0.4	1
77	The median and the mode as robust meta-analysis estimators in the presence of small-study effects and outliers. , 2020, $11$ , $397$ .		1
78	Prevalência de sintomas caracterÃsticos de covid-19 no Rio Grande do Sul: resultados de um estudo de base populacional com 18 mil participantes. Revista De Saude Publica, 2021, 55, 82.	0.7	1