Laurence J Howe

List of Publications by Year
in descending order

25 papers	2,033 citations	567281 15 h-index	552781 26 g-index
35 all docs	35 docs citations	35 times ranked	6477 citing authors

#	Article	IF	Citations
1	Interpreting Mendelian-randomization estimates of the effects of categorical exposures such as disease status and educational attainment. International Journal of Epidemiology, 2022, 51, 948-957.	1.9	17
2	Modeling assortative mating and genetic similarities between partners, siblings, and in-laws. Nature Communications, 2022, 13, 1108.	12.8	23
3	Taller height and risk of coronary heart disease and cancer: A within-sibship Mendelian randomization study. ELife, 2022, 11, .	6.0	6
4	Within-sibship genome-wide association analyses decrease bias in estimates of direct genetic effects. Nature Genetics, 2022, 54, 581-592.	21.4	142
5	The impact of fatty acids biosynthesis on the risk of cardiovascular diseases in Europeans and East Asians: a Mendelian randomization study. Human Molecular Genetics, 2022, 31, 4034-4054.	2.9	5
6	Evaluating indirect genetic effects of siblings using singletons. PLoS Genetics, 2022, 18, e1010247.	3.5	7
7	A GWAS in Latin Americans identifies novel face shape loci, implicating VPS13B and a Denisovan introgressed region in facial variation. Science Advances, 2021, 7, .	10.3	32
8	Genome-wide analysis identifies novel susceptibility loci for myocardial infarction. European Heart Journal, 2021, 42, 919-933.	2.2	113
9	Assortative mating and within-spouse pair comparisons. PLoS Genetics, 2021, 17, e1009883.	3.5	13
10	Vitamin D and the risk of treatment-resistant and atypical depression: A Mendelian randomization study. Translational Psychiatry, 2021, 11, 561.	4.8	9
11	Is population structure in the genetic biobank era irrelevant, a challenge, or an opportunity?. Human Genetics, 2020, 139, 23-41.	3.8	72
12	Avoiding dynastic, assortative mating, and population stratification biases in Mendelian randomization through within-family analyses. Nature Communications, 2020, 11, 3519.	12.8	213
13	Evaluating shared genetic influences on nonsyndromic cleft lip/palate and oropharyngeal neoplasms. Genetic Epidemiology, 2020, 44, 924-933.	1.3	6
14	Cleft lip/palate and educational attainment: cause, consequence or correlation? A Mendelian randomization study. International Journal of Epidemiology, 2020, 49, 1282-1293.	1.9	21
15	Polygenic risk scores for coronary artery disease and subsequent event risk amongst established cases. Human Molecular Genetics, 2020, 29, 1388-1395.	2.9	23
16	Within family Mendelian randomization studies. Human Molecular Genetics, 2019, 28, R170-R179.	2.9	105
17	Prenatal alcohol exposure and facial morphology in a UK cohort. Drug and Alcohol Dependence, 2019, 197, 42-47.	3.2	15
18	Genetic evidence for assortative mating on alcohol consumption in the UK Biobank. Nature Communications, 2019, 10, 5039.	12.8	48

#	Article	IF	Citations
19	Evidence for DNA methylation mediating genetic liability to non-syndromic cleft lip/palate. Epigenomics, 2019, 11, 133-145.	2.1	25
20	Novel genetic loci affecting facial shape variation in humans. ELife, 2019, 8, .	6.0	58
21	Facial Genetics: A Brief Overview. Frontiers in Genetics, 2018, 9, 462.	2.3	79
22	Investigating the shared genetics of non-syndromic cleft lip/palate and facial morphology. PLoS Genetics, 2018, 14, e1007501.	3.5	44
23	Y Chromosome, Mitochondrial DNA and Childhood Behavioural Traits. Scientific Reports, 2017, 7, 11655.	3.3	4
24	Body mass index, body dissatisfaction and adolescent smoking initiation. Drug and Alcohol Dependence, 2017, 178, 143-149.	3.2	26
25	LD Hub: a centralized database and web interface to perform LD score regression that maximizes the potential of summary level GWAS data for SNP heritability and genetic correlation analysis. Bioinformatics, 2017, 33, 272-279.	4.1	822