

Desmond D Campbell

List of Publications by Year in descending order

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Version: 2024-02-01

29
papers

2,368
citations

516710

16
h-index

501196

28
g-index

30
all docs

30
docs citations

30
times ranked

5653
citing authors

#	ARTICLE	IF	CITATIONS
1	Reconstructing Native American population history. <i>Nature</i> , 2012, 488, 370-374.	27.8	699
2	Mental health and health behaviours before and during the initial phase of the COVID-19 lockdown: longitudinal analyses of the UK Household Longitudinal Study. <i>Journal of Epidemiology and Community Health</i> , 2021, 75, jech-2020-215060.	3.7	323
3	Partitioning the Heritability of Tourette Syndrome and Obsessive Compulsive Disorder Reveals Differences in Genetic Architecture. <i>PLoS Genetics</i> , 2013, 9, e1003864.	3.5	241
4	Identifying Gene-Environment Interactions in Schizophrenia: Contemporary Challenges for Integrated, Large-scale Investigations. <i>Schizophrenia Bulletin</i> , 2014, 40, 729-736.	4.3	229
5	DSM-IV combined type ADHD shows familial association with sibling trait scores: A sampling strategy for QTL linkage. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2008, 147B, 1450-1460.	1.7	129
6	Cross-Disorder Genome-Wide Analyses Suggest a Complex Genetic Relationship Between Tourette's Syndrome and OCD. <i>American Journal of Psychiatry</i> , 2015, 172, 82-93.	7.2	117
7	CNV Analysis in Tourette Syndrome Implicates Large Genomic Rearrangements in COL8A1 and NRXN1. <i>PLoS ONE</i> , 2013, 8, e59061.	2.5	70
8	<i>CYP2C19</i> genotype predicts steady state escitalopram concentration in GENDEP. <i>Journal of Psychopharmacology</i> , 2012, 26, 398-407.	4.0	69
9	The emerging molecular architecture of schizophrenia, polygenic risk scores and the clinical implications for GxE research. <i>Social Psychiatry and Psychiatric Epidemiology</i> , 2014, 49, 169-182.	3.1	68
10	Interplay between Schizophrenia Polygenic Risk Score and Childhood Adversity in First-Presentation Psychotic Disorder: A Pilot Study. <i>PLoS ONE</i> , 2016, 11, e0163319.	2.5	52
11	Two subtypes of intervertebral disc degeneration distinguished by large-scale population-based study. <i>Spine Journal</i> , 2016, 16, 1079-1089.	1.3	51
12	Amerind Ancestry, Socioeconomic Status and the Genetics of Type 2 Diabetes in a Colombian Population. <i>PLoS ONE</i> , 2012, 7, e33570.	2.5	47
13	Suicidal ideation during treatment of depression with escitalopram and nortriptyline in Genome-Based Therapeutic Drugs for Depression (GENDEP): a clinical trial. <i>BMC Medicine</i> , 2009, 7, 60.	5.5	43
14	Linkage to Chromosome 1p36 for Attention-Deficit/Hyperactivity Disorder Traits in School and Home Settings. <i>Biological Psychiatry</i> , 2008, 64, 571-576.	1.3	41
15	Cognitive impairment among children at-risk for schizophrenia. <i>Journal of Psychiatric Research</i> , 2014, 50, 92-99.	3.1	34
16	The causal effects of health conditions and risk factors on social and socioeconomic outcomes: Mendelian randomization in UK Biobank. <i>International Journal of Epidemiology</i> , 2020, 49, 1661-1681.	1.9	33
17	Clinical burden, risk factor impact and outcomes following myocardial infarction and stroke: A 25-year individual patient level linkage study. <i>Lancet Regional Health - Europe</i> , The, 2021, 7, 100141.	5.6	18
18	Effects of depression on employment and social outcomes: a Mendelian randomisation study. <i>Journal of Epidemiology and Community Health</i> , 2022, 76, 563-571.	3.7	17

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19	Sacral agenesis: a pilot whole exome sequencing and copy number study. BMC Medical Genetics, 2016, 17, 98.	2.1	15
20	Local True Discovery Rate Weighted Polygenic Scores Using GWAS Summary Data. Behavior Genetics, 2016, 46, 573-582.	2.1	15
21	Gene Variants at Loci Related to Blood Pressure Account for Variation in Response to Antihypertensive Drugs Between Black and White Individuals. Hypertension, 2019, 74, 614-622.	2.7	14
22	Software for generating liability distributions for pedigrees conditional on their observed disease states and covariates. Genetic Epidemiology, 2010, 34, 159-170.	1.3	11
23	Genetic study of congenital bile-duct dilatation identifies de novo and inherited variants in functionally related genes. BMC Medical Genomics, 2016, 9, 75.	1.5	11
24	Salt stress in the renal tubules is linked to TAL-specific expression of uromodulin and an upregulation of heat shock genes. Physiological Genomics, 2018, 50, 964-972.	2.3	7
25	Multifactorial disease risk calculator: Risk prediction for multifactorial disease pedigrees. Genetic Epidemiology, 2018, 42, 130-133.	1.3	6
26	Effects of increased body mass index on employment status: a Mendelian randomisation study. International Journal of Obesity, 2021, 45, 1790-1801.	3.4	4
27	Cost effective assay choice for rare disease study designs. Orphanet Journal of Rare Diseases, 2015, 10, 10.	2.7	3
28	A Single Nucleotide Variant in HNF-1 β is Associated with Maturity-Onset Diabetes of the Young in a Large Chinese Family. Iranian Journal of Public Health, 2016, 45, 170-8.	0.5	1
29	Error in sample size formula. Journal of Human Reproductive Sciences, 2014, 7, 155.	0.9	0