Joey Ward

List of Publications by Year
in descending order

Source: https://exaly.com/author-pdf/7636999/publications.pdf

Version: 2024-02-01

20 papers	2,737 citations	623188 14 h-index	752256 20 g-index
33 all docs	33 docs citations	33 times ranked	5263 citing authors

#	Article	IF	Citations
1	Prediabetes blunts DPP4 genetic control of postprandial glycaemia and insulin secretion. Diabetologia, 2022, 65, 861-871.	2.9	3
2	Dietary patterns, genetic risk, and incidence of obesity: Application of reduced rank regression in 11,735 adults from the UK Biobank study. Preventive Medicine, 2022, 158, 107035.	1.6	7
3	Associations between three diet quality indices, genetic risk and body composition: A prospective cohort study. Clinical Nutrition, 2022, 41, 1942-1949.	2.3	2
4	Sex-stratified genome-wide association study of multisite chronic pain in UK Biobank. PLoS Genetics, 2021, 17, e1009428.	1.5	37
5	Genetic Variation in the ASTN2 Locus in Cardiovascular, Metabolic and Psychiatric Traits: Evidence for Pleiotropy Rather Than Shared Biology. Genes, 2021, 12, 1194.	1.0	4
6	The genomic basis of mood instability: identification of 46 loci in 363,705 UK Biobank participants, genetic correlation with psychiatric disorders, and association with gene expression and function. Molecular Psychiatry, 2020, 25, 3091-3099.	4.1	48
7	Association between APOE e4 and white matter hyperintensity volume, but not total brain volume or white matter integrity. Brain Imaging and Behavior, 2020, 14, 1468-1476.	1.1	62
8	Alzheimerâ€™s Disease Susceptibility Gene Apolipoprotein E (APOE) and Blood Biomarkers in UK Biobank (Nâ€Š=â€Š395,769). Journal of Alzheimer's Disease, 2020, 76, 1541-1551.	1.2	13
9	Association of SBP and BMI with cognitive and structural brain phenotypes in UK Biobank. Journal of Hypertension, 2020, 38, 2482-2489.	0.3	20
10	Genome-wide association study of multisite chronic pain in UK Biobank. PLoS Genetics, 2019, 15, e1008164.	1.5	144
11	Assessing for interaction between APOE Îµ4, sex, and lifestyle on cognitive abilities. Neurology, 2019, 92, e2691-e2698.	1.5	28
12	Identification of novel genome-wide associations for suicidality in UK Biobank, genetic correlation with psychiatric disorders and polygenic association with completed suicide. EBioMedicine, 2019, 41, 517-525.	2.7	87
13	Identification of novel common variants associated with chronic pain using conditional false discovery rate analysis with major depressive disorder and assessment of pleiotropic effects of LRFN5. Translational Psychiatry, 2019, 9, 310.	2.4	16
14	Novel genome-wide associations for anhedonia, genetic correlation with psychiatric disorders, and polygenic association with brain structure. Translational Psychiatry, 2019, 9, 327.	2.4	56
15	Genome-wide meta-analysis of depression identifies 102 independent variants and highlights the importance of the prefrontal brain regions. Nature Neuroscience, 2019, 22, 343-352.	7.1	1,589
16	Genome-wide association study of depression phenotypes in UK Biobank identifies variants in excitatory synaptic pathways. Nature Communications, 2018, 9, 1470.	5.8	415
17	Seasonality of depressive symptoms in women but not in men: A cross-sectional study in the UK Biobank cohort. Journal of Affective Disorders, 2018, 229, 296-305.	2.0	31
18	Polygenic risk scores for major depressive disorder and neuroticism as predictors of antidepressant response: Meta-analysis of three treatment cohorts. PLoS ONE, 2018, 13, e0203896.	1.1	37

#	Article	IF	Citations
19	Genome-Wide Association Study of Circadian Rhythmicity in 71,500 UK Biobank Participants and Polygenic Association with Mood Instability. EBioMedicine, 2018, 35, 279-287.	2.7	53
20	Genome-wide analysis in UK Biobank identifies four loci associated with mood instability and genetic correlation with major depressive disorder, anxiety disorder and schizophrenia. Translational Psychiatry, 2017, 7, 1264.	2.4	69