

Joanne Knight

List of Publications by Year in descending order

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Version: 2024-02-01

34
papers

3,135
citations

394421

19
h-index

414414

32
g-index

38
all docs

38
docs citations

38
times ranked

8472
citing authors

#	ARTICLE	IF	CITATIONS
1	Accuracy and applications of sequencing and genotyping approaches for CYP2A6 and homologous genes. <i>Pharmacogenetics and Genomics</i> , 2022, Publish Ahead of Print, .	1.5	1
2	Late-onset epilepsy predicts future stroke: a systematic review and meta-analysis. <i>Journal of Neurology, Neurosurgery and Psychiatry</i> , 2022, 93, A5.3-A5.	1.9	0
3	Transferability of Ancestry-Specific and Cross-Ancestry CYP2A6 Activity Genetic Risk Scores in African and European Populations. <i>Clinical Pharmacology and Therapeutics</i> , 2021, 110, 975-985.	4.7	15
4	Late-onset epilepsy predicts stroke: Systematic review and meta-analysis. <i>Epilepsy and Behavior</i> , 2021, 115, 107634.	1.7	10
5	A modelling framework for developing early warning systems of COPD emergency admissions. <i>Spatial and Spatio-temporal Epidemiology</i> , 2021, 36, 100392.	1.7	1
6	A Genome-Wide Association Study of Nausea Incidence in Varenicline-Treated Cigarette Smokers. <i>Nicotine and Tobacco Research</i> , 2021, 23, 1805-1809.	2.6	3
7	Variation in waiting times by diagnostic category: an observational study of 1,951 referrals to a neurology outpatient clinic. <i>BMJ Neurology Open</i> , 2021, 3, e000133.	1.6	2
8	Problematising characteristicness. <i>International Journal of Corpus Linguistics</i> , 2021, 26, 305-335.	1.4	3
9	Are child health information services a viable source of accurate vaccination data for clinicians working in paediatric emergency departments in England?. <i>BMJ Health and Care Informatics</i> , 2021, 28, e100486.	3.0	0
10	Routinely collected patient data in neurology research: a systematic mapping review. <i>BMC Neurology</i> , 2020, 20, 431.	1.8	6
11	COVID-19 exposes the urgent need for coding of outpatient neurology episodes. <i>BMJ Neurology Open</i> , 2020, 2, e000080.	1.6	1
12	Cross-disorder analysis of schizophrenia and 19 immune-mediated diseases identifies shared genetic risk. <i>Human Molecular Genetics</i> , 2019, 28, 3498-3513.	2.9	65
13	Can we accurately classify schizophrenia patients from healthy controls using magnetic resonance imaging and machine learning? A multi-method and multi-dataset study. <i>Schizophrenia Research</i> , 2019, 214, 3-10.	2.0	53
14	Comment: Unraveling DNA sequence to identify cerebral indicators of dementia risk. <i>Neurology</i> , 2018, 90, 109-109.	1.1	0
15	Genome-wide association study of a nicotine metabolism biomarker in African American smokers: impact of chromosome 19 genetic influences. <i>Addiction</i> , 2018, 113, 509-523.	3.3	45
16	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
17	<i>APOE</i> ϵ 4 associates with hippocampal volume, learning, and memory across the spectrum of Alzheimer's disease and dementia with Lewy bodies. <i>Alzheimer's and Dementia</i> , 2018, 14, 1137-1147.	0.8	39
18	Heritability estimates of cortical anatomy: The influence and reliability of different estimation strategies. <i>NeuroImage</i> , 2018, 178, 78-91.	4.2	18

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19	Examining the role of common and rare mitochondrial variants in schizophrenia. PLoS ONE, 2018, 13, e0191153.	2.5	23
20	Genome-wide association analysis identifies novel blood pressure loci and offers biological insights into cardiovascular risk. Nature Genetics, 2017, 49, 403-415.	21.4	492
21	Genetic epistasis regulates amyloid deposition in resilient aging. Alzheimer's and Dementia, 2017, 13, 1107-1116.	0.8	8
22	Novel Blood Pressure Locus and Gene Discovery Using Genome-Wide Association Study and Expression Data Sets From Blood and the Kidney. Hypertension, 2017, 70, .	2.7	123
23	Fine-mapping inflammatory bowel disease loci to single-variant resolution. Nature, 2017, 547, 173-178.	27.8	473
24	Dopamine D2 receptor gene variants and response to rasagiline in early Parkinson's disease: a pharmacogenetic study. Brain, 2016, 139, 2050-2062.	7.6	53
25	Allele-Skewed DNA Modification in the Brain: Relevance to a Schizophrenia GWAS. American Journal of Human Genetics, 2016, 98, 956-962.	6.2	20
26	Pharmacogenetic Analysis of Functional Glutamate System Gene Variants and Clinical Response to Clozapine. Molecular Neuropsychiatry, 2016, 2, 185-197.	2.9	14
27	Genome-Wide Association Studies Suggest Limited Immune Gene Enrichment in Schizophrenia Compared to 5 Autoimmune Diseases. Schizophrenia Bulletin, 2016, 42, 1176-1184.	4.3	62
28	IBD Genetic Risk Profile in Healthy First-Degree Relatives of Crohn's Disease Patients. Journal of Crohn's and Colitis, 2016, 10, 209-215.	1.3	32
29	Investigating the Causal Relationship of C-Reactive Protein with 32 Complex Somatic and Psychiatric Outcomes: A Large-Scale Cross-Consortium Mendelian Randomization Study. PLoS Medicine, 2016, 13, e1001976.	8.4	150
30	Pooled Sequencing of 531 Genes in Inflammatory Bowel Disease Identifies an Associated Rare Variant in BTNL2 and Implicates Other Immune Related Genes. PLoS Genetics, 2015, 11, e1004955.	3.5	59
31	A genome-wide association study of suicide severity scores in bipolar disorder. Journal of Psychiatric Research, 2015, 65, 23-29.	3.1	36
32	A Bayesian Method to Incorporate Hundreds of Functional Characteristics with Association Evidence to Improve Variant Prioritization. PLoS ONE, 2014, 9, e98122.	2.5	29
33	Complex host genetics influence the microbiome in inflammatory bowel disease. Genome Medicine, 2014, 6, 107.	8.2	322
34	The role of leptin, melanocortin, and neurotrophin system genes on body weight in anorexia nervosa and bulimia nervosa. Journal of Psychiatric Research, 2014, 55, 77-86.	3.1	25