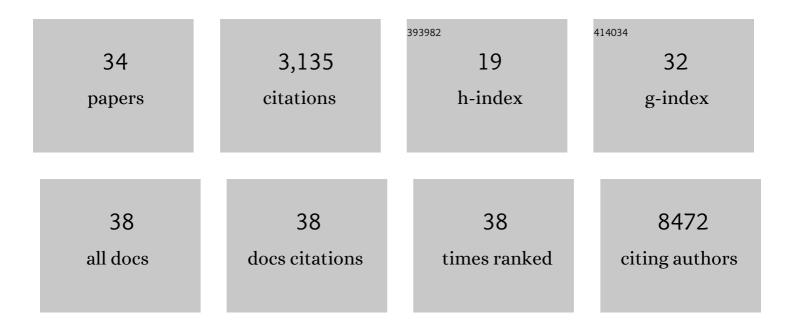
Joanne Knight

List of Publications by Year in descending order

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#	ARTICLE	IF	CITATIONS
1	Genetic analysis of over 1 million people identifies 535 new loci associated with blood pressure traits. Nature Genetics, 2018, 50, 1412-1425.	9.4	924
2	Genome-wide association analysis identifies novel blood pressure loci and offers biological insights into cardiovascular risk. Nature Genetics, 2017, 49, 403-415.	9.4	492
3	Fine-mapping inflammatory bowel disease loci to single-variant resolution. Nature, 2017, 547, 173-178.	13.7	473
4	Complex host genetics influence the microbiome in inflammatory bowel disease. Genome Medicine, 2014, 6, 107.	3.6	322
5	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.	3.9	150
6	Novel Blood Pressure Locus and Gene Discovery Using Genome-Wide Association Study and Expression Data Sets From Blood and the Kidney. Hypertension, 2017, 70, .	1.3	123
7	Cross-disorder analysis of schizophrenia and 19 immune-mediated diseases identifies shared genetic risk. Human Molecular Genetics, 2019, 28, 3498-3513.	1.4	65
8	Genome-Wide Association Studies Suggest Limited Immune Gene Enrichment in Schizophrenia Compared to 5 Autoimmune Diseases. Schizophrenia Bulletin, 2016, 42, 1176-1184.	2.3	62
9	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.	1.5	59
10	Dopamine D2 receptor gene variants and response to rasagiline in early Parkinson's disease: a pharmacogenetic study. Brain, 2016, 139, 2050-2062.	3.7	53
11	Can we accurately classify schizophrenia patients from healthy controls using magnetic resonance imaging and machine learning? A multi-method and multi-dataset study. Schizophrenia Research, 2019, 214, 3-10.	1.1	53
12	Genomeâ€wide association study of a nicotine metabolism biomarker in African American smokers: impact of chromosome 19 genetic influences. Addiction, 2018, 113, 509-523.	1.7	45
13	<i>APOE</i> ‵4 associates with hippocampal volume, learning, and memory across the spectrum of Alzheimer's disease and dementia with Lewy bodies. Alzheimer's and Dementia, 2018, 14, 1137-1147.	0.4	39
14	A genome-wide association study of suicide severity scores in bipolar disorder. Journal of Psychiatric Research, 2015, 65, 23-29.	1.5	36
15	IBD Genetic Risk Profile in Healthy First-Degree Relatives of Crohn's Disease Patients. Journal of Crohn's and Colitis, 2016, 10, 209-215.	0.6	32
16	A Bayesian Method to Incorporate Hundreds of Functional Characteristics with Association Evidence to Improve Variant Prioritization. PLoS ONE, 2014, 9, e98122.	1.1	29
17	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.	1.5	25
18	Examining the role of common and rare mitochondrial variants in schizophrenia. PLoS ONE, 2018, 13, e0191153.	1.1	23

JOANNE KNIGHT

#	Article	IF	CITATIONS
19	Allele-Skewed DNA Modification in the Brain: Relevance to a Schizophrenia GWAS. American Journal of Human Genetics, 2016, 98, 956-962.	2.6	20
20	Heritability estimates of cortical anatomy: The influence and reliability of different estimation strategies. Neurolmage, 2018, 178, 78-91.	2.1	18
21	Transferability of Ancestryâ€Specific and Crossâ€Ancestry CYP2A6 Activity Genetic Risk Scores in African and European Populations. Clinical Pharmacology and Therapeutics, 2021, 110, 975-985.	2.3	15
22	Pharmacogenetic Analysis of Functional Glutamate System Gene Variants and Clinical Response to Clozapine. Molecular Neuropsychiatry, 2016, 2, 185-197.	3.0	14
23	Late-onset epilepsy predicts stroke: Systematic review and meta-analysis. Epilepsy and Behavior, 2021, 115, 107634.	0.9	10
24	Genetic epistasis regulates amyloid deposition in resilient aging. Alzheimer's and Dementia, 2017, 13, 1107-1116.	0.4	8
25	Routinely collected patient data in neurology research: a systematic mapping review. BMC Neurology, 2020, 20, 431.	0.8	6
26	A Genome-Wide Association Study of Nausea Incidence in Varenicline-Treated Cigarette Smokers. Nicotine and Tobacco Research, 2021, 23, 1805-1809.	1.4	3
27	Problematising characteristicness. International Journal of Corpus Linguistics, 2021, 26, 305-335.	0.6	3
28	Variation in waiting times by diagnostic category: an observational study of 1,951 referrals to a neurology outpatient clinic. BMJ Neurology Open, 2021, 3, e000133.	0.7	2
29	COVID-19 exposes the urgent need for coding of outpatient neurology episodes. BMJ Neurology Open, 2020, 2, e000080.	0.7	1
30	A modelling framework for developing early warning systems of COPD emergency admissions. Spatial and Spatio-temporal Epidemiology, 2021, 36, 100392.	0.9	1
31	Accuracy and applications of sequencing and genotyping approaches for CYP2A6 and homologous genes. Pharmacogenetics and Genomics, 2022, Publish Ahead of Print, .	0.7	1
32	Comment: Unraveling DNA sequence to identify cerebral indicators of dementia risk. Neurology, 2018, 90, 109-109.	1.5	0
33	Are child health information services a viable source of accurate vaccination data for clinicians working in paediatric emergency departments in England?. BMJ Health and Care Informatics, 2021, 28, e100486.	1.4	0
34	Late-onset epilepsy predicts future stroke: a systematic review and meta-analysis. Journal of Neurology, Neurosurgery and Psychiatry, 2022, 93, A5.3-A5.	0.9	0