

Erin B Ware

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

50
papers

2,467
citations

15
h-index

49
g-index

63
ext. papers

3,397
ext. citations

8.9
avg, IF

3.27
L-index

#	Paper	IF	Citations
50	Gene-educational attainment interactions in a multi-ancestry genome-wide meta-analysis identify novel blood pressure loci. <i>Molecular Psychiatry</i> , 2021 , 26, 2111-2125	15.1	3
49	Type 2 Diabetes and Cognitive Status in the Health and Retirement Study: A Mendelian Randomization Approach. <i>Frontiers in Genetics</i> , 2021 , 12, 634767	4.5	4
48	Cumulative Genetic Risk and Are Independently Associated With Dementia Status in a Multiethnic, Population-Based Cohort. <i>Neurology: Genetics</i> , 2021 , 7, e576	3.8	2
47	Discovery and fine-mapping of height loci via high-density imputation of GWASs in individuals of African ancestry. <i>American Journal of Human Genetics</i> , 2021 , 108, 564-582	11	7
46	A multi-ethnic epigenome-wide association study of leukocyte DNA methylation and blood lipids. <i>Nature Communications</i> , 2021 , 12, 3987	17.4	3
45	Linking gene regions jointly with environment and depression 2021 , 69-76		
44	Saliva cell type DNA methylation reference panel for epidemiological studies in children. <i>Epigenetics</i> , 2021 , 1-17	5.7	3
43	Type 2 diabetes and dementia in the Health and Retirement Study: A Mendelian randomization approach. <i>Alzheimer's and Dementia</i> , 2020 , 16, e041220	1.2	
42	Social regulation of inflammation related gene expression in the multi-ethnic study of atherosclerosis. <i>Psychoneuroendocrinology</i> , 2020 , 117, 104654	5	7
41	A data-driven prospective study of dementia among older adults in the United States. <i>PLoS ONE</i> , 2020 , 15, e0239994	3.7	4
40	Considering the APOE locus in Alzheimer's disease polygenic scores in the Health and Retirement Study: a longitudinal panel study. <i>BMC Medical Genomics</i> , 2020 , 13, 164	3.7	3
39	A data-driven prospective study of dementia among older adults in the United States 2020 , 15, e0239994		
38	A data-driven prospective study of dementia among older adults in the United States 2020 , 15, e0239994		
37	A data-driven prospective study of dementia among older adults in the United States 2020 , 15, e0239994		
36	A data-driven prospective study of dementia among older adults in the United States 2020 , 15, e0239994		
35	Expression of socially sensitive genes: The multi-ethnic study of atherosclerosis. <i>PLoS ONE</i> , 2019 , 14, e0214061	3.7	5
34	A multi-ancestry genome-wide study incorporating gene-smoking interactions identifies multiple new loci for pulse pressure and mean arterial pressure. <i>Human Molecular Genetics</i> , 2019 , 28, 2615-2633	5.6	14

33	Multi-ancestry genome-wide gene-smoking interaction study of 387,272 individuals identifies new loci associated with serum lipids. <i>Nature Genetics</i> , 2019 , 51, 636-648	36.3	59
32	A meta-analysis of genome-wide association studies identifies multiple longevity genes. <i>Nature Communications</i> , 2019 , 10, 3669	17.4	102
31	Genome-wide Association Study of 24-Hour Urinary Excretion of Calcium, Magnesium, and Uric Acid. <i>Mayo Clinic Proceedings Innovations, Quality & Outcomes</i> , 2019 , 3, 448-460	3.1	2
30	Combined linkage and association analysis identifies rare and low frequency variants for blood pressure at 1q31. <i>European Journal of Human Genetics</i> , 2019 , 27, 269-277	5.3	3
29	Testing cross-phenotype effects of rare variants in longitudinal studies of complex traits. <i>Genetic Epidemiology</i> , 2018 , 42, 320-332	2.6	3
28	A Large-Scale Multi-ancestry Genome-wide Study Accounting for Smoking Behavior Identifies Multiple Significant Loci for Blood Pressure. <i>American Journal of Human Genetics</i> , 2018 , 102, 375-400	11	59
27	Hormone therapy and urine protein excretion: a multiracial cohort study, systematic review, and meta-analysis. <i>Menopause</i> , 2018 , 25, 625-634	2.5	9
26	DNA methylation age is associated with an altered hemostatic profile in a multiethnic meta-analysis. <i>Blood</i> , 2018 , 132, 1842-1850	2.2	11
25	Novel genetic associations for blood pressure identified via gene-alcohol interaction in up to 570K individuals across multiple ancestries. <i>PLoS ONE</i> , 2018 , 13, e0198166	3.7	31
24	Protein-altering variants associated with body mass index implicate pathways that control energy intake and expenditure in obesity. <i>Nature Genetics</i> , 2018 , 50, 26-41	36.3	186
23	An Analysis of Two Genome-wide Association Meta-analyses Identifies a New Locus for Broad Depression Phenotype. <i>Biological Psychiatry</i> , 2017 , 82, 322-329	7.9	68
22	Polymorphisms in Renal Ammonia Metabolism Genes Correlate With 24-Hour Urine pH. <i>Kidney International Reports</i> , 2017 , 2, 1111-1121	4.1	6
21	Interaction between Social/Psychosocial Factors and Genetic Variants on Body Mass Index: A Gene-Environment Interaction Analysis in a Longitudinal Setting. <i>International Journal of Environmental Research and Public Health</i> , 2017 , 14,	4.6	8
20	Rare variants in fox-1 homolog A (RFX1) are associated with lower blood pressure. <i>PLoS Genetics</i> , 2017 , 13, e1006678	6	11
19	Discovery and fine-mapping of adiposity loci using high density imputation of genome-wide association studies in individuals of African ancestry: African Ancestry Anthropometry Genetics Consortium. <i>PLoS Genetics</i> , 2017 , 13, e1006719	6	60
18	Single-trait and multi-trait genome-wide association analyses identify novel loci for blood pressure in African-ancestry populations. <i>PLoS Genetics</i> , 2017 , 13, e1006728	6	58
17	Applying Novel Methods for Assessing Individual- and Neighborhood-Level Social and Psychosocial Environment Interactions with Genetic Factors in the Prediction of Depressive Symptoms in the Multi-Ethnic Study of Atherosclerosis. <i>Behavior Genetics</i> , 2016 , 46, 89-99	3.2	6
16	Epigenetic Signatures of Cigarette Smoking. <i>Circulation: Cardiovascular Genetics</i> , 2016 , 9, 436-447		442

15	Genetic variants linked to education predict longevity. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016 , 113, 13366-13371	11.5	90
14	Key influence of sex on urine volume and osmolality. <i>Biology of Sex Differences</i> , 2016 , 7, 12	9.3	13
13	Heritability of dietary traits that contribute to nephrolithiasis in a cohort of adult sibships. <i>Journal of Nephrology</i> , 2016 , 29, 45-51	4.8	7
12	A Statistical Approach for Testing Cross-Phenotype Effects of Rare Variants. <i>American Journal of Human Genetics</i> , 2016 , 98, 525-540	11	40
11	Genome-wide association study identifies 74 loci associated with educational attainment. <i>Nature</i> , 2016 , 533, 539-42	50.4	850
10	Directional dominance on stature and cognition in diverse human populations. <i>Nature</i> , 2015 , 523, 459-463	50.4	119
9	A statistical approach for rare-variant association testing in affected sibships. <i>American Journal of Human Genetics</i> , 2015 , 96, 543-54	11	14
8	Current Applications of Genetic Risk Scores to Cardiovascular Outcomes and Subclinical Phenotypes. <i>Current Epidemiology Reports</i> , 2015 , 2, 180-190	2.9	58
7	Effect of Demographics on Excretion of Key Urinary Factors Related to Kidney Stone Risk. <i>Urology</i> , 2015 , 86, 690-6	1.6	11
6	Comparative genome-wide association studies of a depressive symptom phenotype in a repeated measures setting by race/ethnicity in the Multi-Ethnic Study of Atherosclerosis. <i>BMC Genetics</i> , 2015 , 16, 118	2.6	11
5	Association between Stress Response Genes and Features of Diurnal Cortisol Curves in the Multi-Ethnic Study of Atherosclerosis: A New Multi-Phenotype Approach for Gene-Based Association Tests. <i>PLoS ONE</i> , 2015 , 10, e0126637	3.7	5
4	SLC2A9 Genotype Is Associated with SLC2A9 Gene Expression and Urinary Uric Acid Concentration. <i>PLoS ONE</i> , 2015 , 10, e0128593	3.7	9
3	Phenotypic and Genetic Markers of Psychopathology in a Population-Based Sample of Older Adults		1
2	Heterogeneity in polygenic scores for common human traits		49
1	Considering the APOE locus in polygenic scores for Alzheimer's disease		1