

Keith L Keene

List of Publications by Year in descending order

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31
papers

2,693
citations

471509
17
h-index

395702
33
g-index

33
all docs

33
docs citations

33
times ranked

6725
citing authors

#	ARTICLE	IF	CITATIONS
1	Multiancestry genome-wide association study of 520,000 subjects identifies 32 loci associated with stroke and stroke subtypes. <i>Nature Genetics</i> , 2018, 50, 524-537.	21.4	1,124
2	Genetic associations at 53 loci highlight cell types and biological pathways relevant for kidney function. <i>Nature Communications</i> , 2016, 7, 10023.	12.8	412
3	Loci associated with ischaemic stroke and its subtypes (SiGN): a genome-wide association study. <i>Lancet Neurology</i> , The, 2016, 15, 174-184.	10.2	217
4	Meta-Analysis of Genome-Wide Association Studies in African Americans Provides Insights into the Genetic Architecture of Type 2 Diabetes. <i>PLoS Genetics</i> , 2014, 10, e1004517.	3.5	191
5	Type 2 Diabetes Variants Disrupt Function of SLC16A11 through Two Distinct Mechanisms. <i>Cell</i> , 2017, 170, 199-212.e20.	28.9	121
6	Variants of the Transcription Factor 7-Like 2 (TCF7L2) Gene Are Associated With Type 2 Diabetes in an African-American Population Enriched for Nephropathy. <i>Diabetes</i> , 2007, 56, 2638-2642.	0.6	89
7	Genome-Wide Meta-Analysis of Homocysteine and Methionine Metabolism Identifies Five One Carbon Metabolism Loci and a Novel Association of ALDH1L1 with Ischemic Stroke. <i>PLoS Genetics</i> , 2014, 10, e1004214.	3.5	69
8	Meta-Analysis of Genome-Wide Association Studies Identifies Genetic Risk Factors for Stroke in African Americans. <i>Stroke</i> , 2015, 46, 2063-2068.	2.0	63
9	Genome-Wide Association Study for Circulating Tissue Plasminogen Activator Levels and Functional Follow-Up Implicates Endothelial <i>STXBP5</i> and <i>STX2</i> . <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2014, 34, 1093-1101.	2.4	43
10	Shared genetic susceptibility of vascular-related biomarkers with ischemic and recurrent stroke. <i>Neurology</i> , 2016, 86, 351-359.	1.1	33
11	Investigation of the Estrogen Receptor- α Gene With Type 2 Diabetes and/or Nephropathy in African-American and European-American Populations. <i>Diabetes</i> , 2007, 56, 675-684.	0.6	30
12	Exploration of the utility of ancestry informative markers for genetic association studies of African Americans with type 2 diabetes and end stage renal disease. <i>Human Genetics</i> , 2008, 124, 147-154.	3.8	29
13	Comprehensive evaluation of the estrogen receptor α gene reveals further evidence for association with type 2 diabetes enriched for nephropathy in an African American population. <i>Human Genetics</i> , 2008, 123, 333-341.	3.8	28
14	Association of the Distal Region of the Ectonucleotide Pyrophosphatase/Phosphodiesterase 1 Gene With Type 2 Diabetes in an African-American Population Enriched for Nephropathy. <i>Diabetes</i> , 2008, 57, 1057-1062.	0.6	28
15	Genome-Wide Association Study Meta-Analysis of Stroke in 22 000 Individuals of African Descent Identifies Novel Associations With Stroke. <i>Stroke</i> , 2020, 51, 2454-2463.	2.0	26
16	Genetic Associations with Plasma B12, B6, and Folate Levels in an Ischemic Stroke Population from the Vitamin Intervention for Stroke Prevention (VISP) Trial. <i>Frontiers in Public Health</i> , 2014, 2, 112.	2.7	23
17	Metabolic Traits and Stroke Risk in Individuals of African Ancestry: Mendelian Randomization Analysis. <i>Stroke</i> , 2021, 52, 2680-2684.	2.0	22
18	Genetic Drivers of von Willebrand Factor Levels in an Ischemic Stroke Population and Association With Risk for Recurrent Stroke. <i>Stroke</i> , 2017, 48, 1444-1450.	2.0	21

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19	Fine Mapping and Functional Studies of Risk Variants for Type 1 Diabetes at Chromosome 16p13.13. Diabetes, 2014, 63, 4360-4368.	0.6	17
20	Evaluation of a SNP map of 6q24â€“27 confirms diabetic nephropathy loci and identifies novel associations in type 2 diabetes patients with nephropathy from an African-American population. Human Genetics, 2008, 124, 63-71.	3.8	14
21	Chromosome 7p linkage and association study for diabetes related traits and type 2 diabetes in an African-American population enriched for nephropathy. BMC Medical Genetics, 2010, 11, 22.	2.1	13
22	Epigenome-Wide Analyses Identify Two Novel Associations With Recurrent Stroke in the Vitamin Intervention for Stroke Prevention Clinical Trial. Frontiers in Genetics, 2018, 9, 358.	2.3	12
23	Multi-phenotype analyses of hemostatic traits with cardiovascular events reveal novel genetic associations. Journal of Thrombosis and Haemostasis, 2022, 20, 1331-1349.	3.8	12
24	Genetic landscape of Gullah African Americans. American Journal of Physical Anthropology, 2021, 175, 905-919.	2.1	9
25	Association Analysis of the Ephrin-B2 Gene in African-Americans with End-Stage Renal Disease. American Journal of Nephrology, 2008, 28, 914-920.	3.1	7
26	DNA methylation analyses identify an intronic ZDHHC6 locus associated with time to recurrent stroke in the Vitamin Intervention for Stroke Prevention (VISP) clinical trial. PLoS ONE, 2021, 16, e0254562.	2.5	5
27	Multi-omic analysis of stroke recurrence in African Americans from the Vitamin Intervention for Stroke Prevention (VISP) clinical trial. PLoS ONE, 2021, 16, e0247257.	2.5	4
28	Cervical Artery Dissection in Patients of African Ancestry. Cerebrovascular Diseases, 2018, 46, 218-222.	1.7	3
29	Differential expression of PHACTR1 in atheromatous versus normal carotid artery tissue. Journal of Clinical Neuroscience, 2020, 74, 265-267.	1.5	3
30	The Impact of COVID-19 on Racial-Ethnic Health Disparities in the US: Now Is the Time To Address the Problem. Journal of the National Medical Association, 2021, 113, 195-198.	0.8	3
31	Estimating Clinical Research Project Duration from Idea to Publication. Journal of Investigative Medicine, 2022, 70, 108-109.	1.6	3