Keith L Keene

List of Publications by Year
in descending order

31 papers	2,693 citations	471371 17 h-index	395590 33 g-index
33 all docs	33 docs citations	33 times ranked	6725 citing authors

#	Article	IF	Citations
1	Estimating Clinical Research Project Duration from Idea to Publication. Journal of Investigative Medicine, 2022, 70, 108-109.	0.7	3
2	Multiâ€phenotype analyses of hemostatic traits with cardiovascular events reveal novel genetic associations. Journal of Thrombosis and Haemostasis, 2022, 20, 1331-1349.	1.9	12
3	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.	0.6	3
4	Multi-omic analysis of stroke recurrence in African Americans from the Vitamin Intervention for Stroke Prevention (VISP) clinical trial. PLoS ONE, 2021, 16, e0247257.	1.1	4
5	Genetic landscape of Gullah African Americans. American Journal of Physical Anthropology, 2021, 175, 905-919.	2.1	9
6	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.	1.1	5
7	Metabolic Traits and Stroke Risk in Individuals of African Ancestry: Mendelian Randomization Analysis. Stroke, 2021, 52, 2680-2684.	1.0	22
8	Genome-Wide Association Study Meta-Analysis of Stroke in 22 000 Individuals of African Descent Identifies Novel Associations With Stroke. Stroke, 2020, 51, 2454-2463.	1.0	26
9	Differential expression of PHACTR1 in atheromatous versus normal carotid artery tissue. Journal of Clinical Neuroscience, 2020, 74, 265-267.	0.8	3
10	Cervical Artery Dissection in Patients of African Ancestry. Cerebrovascular Diseases, 2018, 46, 218-222.	0.8	3
11	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.	1.1	12
12	Multiancestry genome-wide association study of 520,000 subjects identifies 32 loci associated with stroke and stroke subtypes. Nature Genetics, 2018, 50, 524-537.	9.4	1,124
13	Genetic Drivers of von Willebrand Factor Levels in an Ischemic Stroke Population and Association With Risk for Recurrent Stroke. Stroke, 2017, 48, 1444-1450.	1.0	21
14	Type 2 Diabetes Variants Disrupt Function of SLC16A11 through Two Distinct Mechanisms. Cell, 2017, 170, 199-212.e20.	13.5	121
15	Loci associated with ischaemic stroke and its subtypes (SiGN): a genome-wide association study. Lancet Neurology, The, 2016, 15, 174-184.	4.9	217
16	Genetic associations at 53 loci highlight cell types and biological pathways relevant for kidney function. Nature Communications, 2016, 7, 10023.	5.8	412
17	Shared genetic susceptibility of vascular-related biomarkers with ischemic and recurrent stroke. Neurology, 2016, 86, 351-359.	1.5	33
18	Meta-Analysis of Genome-Wide Association Studies Identifies Genetic Risk Factors for Stroke in African Americans. Stroke, 2015, 46, 2063-2068.	1.0	63

#	Article	IF	Citations
19	Genetic Associations with Plasma B12, B6, and Folate Levels in an Ischemic Stroke Population from the Vitamin Intervention for Stroke Prevention (VISP) Trial. Frontiers in Public Health, 2014, 2, 112.	1.3	23
20	Genome-Wide Association Study for Circulating Tissue Plasminogen Activator Levels and Functional Follow-Up Implicates Endothelial STXBP5 and STX2. Arteriosclerosis, Thrombosis, and Vascular Biology, 2014, 34, 1093-1101.	1.1	43
21	Genome-Wide Meta-Analysis of Homocysteine and Methionine Metabolism Identifies Five One Carbon Metabolism Loci and a Novel Association of ALDH1L1 with Ischemic Stroke. PLoS Genetics, 2014, 10, e1004214.	1.5	69
22	Meta-Analysis of Genome-Wide Association Studies in African Americans Provides Insights into the Genetic Architecture of Type 2 Diabetes. PLoS Genetics, 2014, 10, e1004517.	1.5	191
23	Fine Mapping and Functional Studies of Risk Variants for Type 1 Diabetes at Chromosome 16p13.13. Diabetes, 2014, 63, 4360-4368.	0.3	17
24	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
25	Comprehensive evaluation of the estrogen receptor Î± gene reveals further evidence for association with type 2 diabetes enriched for nephropathy in an African American population. Human Genetics, 2008, 123, 333-341.	1.8	28
26	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.	1.8	14
27	Exploration of the utility of ancestry informative markers for genetic association studies of African Americans with type 2 diabetes and end stage renal disease. Human Genetics, 2008, 124, 147-154.	1.8	29
28	Association of the Distal Region of the Ectonucleotide Pyrophosphatase/Phosphodiesterase 1 Gene With Type 2 Diabetes in an African-American Population Enriched for Nephropathy. Diabetes, 2008, 57, 1057-1062.	0.3	28
29	Association Analysis of the Ephrin-B2 Gene in African-Americans with End-Stage Renal Disease. American Journal of Nephrology, 2008, 28, 914-920.	1.4	7
30	Variants of the Transcription Factor 7-Like 2 (TCF7L2) Gene Are Associated With Type 2 Diabetes in an African-American Population Enriched for Nephropathy. Diabetes, 2007, 56, 2638-2642.	0.3	89
31	Investigation of the Estrogen Receptor-Â Gene With Type 2 Diabetes and/or Nephropathy in African-American and European-American Populations. Diabetes, 2007, 56, 675-684.	0.3	30