Tanika N Kelly

List of Publications by Year in descending order

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97 8,144 27 84
papers citations h-index g-index

101 101 101 15864 all docs docs citations times ranked citing authors

#	Article	IF	CITATIONS
1	Global Disparities of Hypertension Prevalence and Control. Circulation, 2016, 134, 441-450.	1.6	2,345
2	Sequencing of 53,831 diverse genomes from the NHLBI TOPMed Program. Nature, 2021, 590, 290-299.	13.7	1,069
3	A systematic analysis of worldwide population-based data on the global burden of chronic kidney disease in 2010. Kidney International, 2015, 88, 950-957.	2.6	597
4	Systolic Blood Pressure Reduction and Risk of Cardiovascular Disease and Mortality. JAMA Cardiology, 2017, 2, 775.	3.0	515
5	Inherited causes of clonal haematopoiesis in 97,691 whole genomes. Nature, 2020, 586, 763-768.	13.7	376
6	Systematic Review: Glucose Control and Cardiovascular Disease in Type 2 Diabetes. Annals of Internal Medicine, 2009, 151, 394.	2.0	308
7	Trans-ancestry genome-wide association study identifies 12 genetic loci influencing blood pressure and implicates a role for DNA methylation. Nature Genetics, 2015, 47, 1282-1293.	9.4	294
8	Gut Microbiome Associates With Lifetime Cardiovascular Disease Risk Profile Among Bogalusa Heart Study Participants. Circulation Research, 2016, 119, 956-964.	2.0	264
9	Use of >100,000 NHLBI Trans-Omics for Precision Medicine (TOPMed) Consortium whole genome sequences improves imputation quality and detection of rare variant associations in admixed African and Hispanic/Latino populations. PLoS Genetics, 2019, 15, e1008500.	1.5	203
10	Gender difference in blood pressure responses to dietary sodium intervention in the GenSalt study. Journal of Hypertension, 2009, 27, 48-54.	0.3	180
11	Multi-ancestry genome-wide gene–smoking interaction study of 387,272 individuals identifies new loci associated with serum lipids. Nature Genetics, 2019, 51, 636-648.	9.4	112
12	Sex, gut microbiome, and cardiovascular disease risk. Biology of Sex Differences, 2019, 10, 29.	1.8	95
13	Novel genetic associations for blood pressure identified via gene-alcohol interaction in up to 570K individuals across multiple ancestries. PLoS ONE, 2018, 13, e0198166.	1.1	94
14	Hypertension Subtype and Risk of Cardiovascular Disease in Chinese Adults. Circulation, 2008, 118, 1558-1566.	1.6	88
15	Multiancestry Genome-Wide Association Study of Lipid Levels Incorporating Gene-Alcohol Interactions. American Journal of Epidemiology, 2019, 188, 1033-1054.	1.6	85
16	Cigarette Smoking and Risk of Stroke in the Chinese Adult Population. Stroke, 2008, 39, 1688-1693.	1.0	75
17	Interethnic analyses of blood pressure loci in populations of East Asian and European descent. Nature Communications, 2018, 9, 5052.	5.8	75
18	Genome-Wide Association Study Identifies 8 Novel Loci Associated With Blood Pressure Responses to Interventions in Han Chinese. Circulation: Cardiovascular Genetics, 2013, 6, 598-607.	5.1	64

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19	Multi-ancestry study of blood lipid levels identifies four loci interacting with physical activity. Nature Communications, 2019, 10, 376.	5.8	64
20	Genome-Wide Association Study Meta-Analysis Reveals Transethnic Replication of Mean Arterial and Pulse Pressure Loci. Hypertension, 2013, 62, 853-859.	1.3	63
21	Multi-ancestry sleep-by-SNP interaction analysis in 126,926 individuals reveals lipid loci stratified by sleep duration. Nature Communications, 2019, 10, 5121.	5.8	62
22	Endovascular Treatment with Stent-Retriever Devices for Acute Ischemic Stroke: A Meta-Analysis of Randomized Controlled Trials. PLoS ONE, 2016, 11, e0147287.	1.1	59
23	Genomic epidemiology of blood pressure salt sensitivity. Journal of Hypertension, 2012, 30, 861-873.	0.3	53
24	Metabolomic Markers of Kidney Function Decline in Patients With Diabetes: Evidence From the Chronic Renal Insufficiency Cohort (CRIC) Study. American Journal of Kidney Diseases, 2020, 76, 511-520.	2.1	45
25	Genome-Wide Gene–Sodium Interaction Analyses on Blood Pressure. Hypertension, 2016, 68, 348-355.	1.3	44
26	Association of Genetic Variants in the Apelin-APJ System and ACE2 With Blood Pressure Responses to Potassium Supplementation: The GenSalt Study. American Journal of Hypertension, 2010, 23, 606-613.	1.0	37
27	Mendelian randomization supports bidirectional causality between telomere length and clonal hematopoiesis of indeterminate potential. Science Advances, 2022, 8, eabl6579.	4.7	36
28	Leveraging linkage evidence to identify low-frequency and rare variants on 16p13 associated with blood pressure using TOPMed whole genome sequencing data. Human Genetics, 2019, 138, 199-210.	1.8	29
29	A System for Phenotype Harmonization in the National Heart, Lung, and Blood Institute Trans-Omics for Precision Medicine (TOPMed) Program. American Journal of Epidemiology, 2021, 190, 1977-1992.	1.6	29
30	Genetic variants in the renin–angiotensin–aldosterone system and blood pressure responses to potassium intake. Journal of Hypertension, 2011, 29, 1719-1730.	0.3	27
31	Genome-Wide Association Study Meta-Analysis of Long-Term Average Blood Pressure in East Asians. Circulation: Cardiovascular Genetics, 2017, 10, e001527.	5.1	26
32	Resequencing Epithelial Sodium Channel Genes Identifies Rare Variants Associated With Blood Pressure Salt-Sensitivity: The GenSalt Study. American Journal of Hypertension, 2018, 31, 205-211.	1.0	25
33	Analysis of Sex Hormone Genes Reveals Gender Differences in the Genetic Etiology of Blood Pressure Salt Sensitivity: The GenSalt Study. American Journal of Hypertension, 2013, 26, 191-200.	1.0	24
34	Variability and rapid increase in body mass index during childhood are associated with adult obesity. International Journal of Epidemiology, 2015, 44, 1943-1950.	0.9	24
35	Blood Pressure Genetic Risk Score Predicts Blood Pressure Responses to Dietary Sodium and Potassium. Hypertension, 2017, 70, 1106-1112.	1.3	24
36	Rare coding variants in 35 genes associate with circulating lipid levels—A multi-ancestry analysis of 170,000 exomes. American Journal of Human Genetics, 2022, 109, 81-96.	2.6	24

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37	Novel Genetic Variants in the α-Adducin and Guanine Nucleotide Binding Protein β-Polypeptide 3 Genes and Salt Sensitivity of Blood Pressure. American Journal of Hypertension, 2009, 22, 985-992.	1.0	23
38	The role of renin–angiotensin–aldosterone system genes in the progression of chronic kidney disease: findings from the Chronic Renal Insufficiency Cohort (CRIC) study. Nephrology Dialysis Transplantation, 2015, 30, 1711-1718.	0.4	22
39	An untargeted metabolomics study of blood pressure: findings from the Bogalusa Heart Study. Journal of Hypertension, 2020, 38, 1302-1311.	0.3	22
40	Human epithelial Na ⁺ channel missense variants identified in the GenSalt study alter channel activity. American Journal of Physiology - Renal Physiology, 2016, 311, F908-F914.	1.3	21
41	A Gene-Based Analysis of Variants in the Serum/Glucocorticoid Regulated Kinase (SGK) Genes with Blood Pressure Responses to Sodium Intake: The GenSalt Study. PLoS ONE, 2014, 9, e98432.	1.1	21
42	Predicting Long-Term Absence of Coronary Artery Calcium in Metabolic Syndrome and Diabetes. JACC: Cardiovascular Imaging, 2021, 14, 219-229.	2.3	19
43	A History of Asthma From Childhood andÂLeft Ventricular Mass in AsymptomaticÂYoung Adults. JACC: Heart Failure, 2017, 5, 497-504.	1.9	17
44	Novel Metabolites Are Associated With Augmentation Index and Pulse Wave Velocity: Findings From the Bogalusa Heart Study. American Journal of Hypertension, 2019, 32, 547-556.	1.0	17
45	Gene-educational attainment interactions in a multi-ancestry genome-wide meta-analysis identify novel blood pressure loci. Molecular Psychiatry, 2020, 26, 2111-2125.	4.1	17
46	Associations of Epithelial Sodium Channel Genes With Blood Pressure Changes and Hypertension Incidence: The GenSalt Study. American Journal of Hypertension, 2014, 27, 1370-1376.	1.0	16
47	Whole-Genome Sequencing Association Analyses of Stroke and Its Subtypes in Ancestrally Diverse Populations From Trans-Omics for Precision Medicine Project. Stroke, 2021, , STROKEAHA120031792.	1.0	16
48	Pseudouridine and N-formylmethionine associate with left ventricular mass index: Metabolome-wide association analysis of cardiac remodeling. Journal of Molecular and Cellular Cardiology, 2020, 140, 22-29.	0.9	15
49	Novel serum metabolites associate with cognition phenotypes among Bogalusa Heart Study participants. Aging, 2019, 11, 5124-5139.	1.4	15
50	Whole genome sequence analyses of eGFR in 23,732 people representing multiple ancestries in the NHLBI trans-omics for precision medicine (TOPMed) consortium. EBioMedicine, 2021, 63, 103157.	2.7	14
51	Sodium Sensitivity, Sodium Resistance, and Incidence of Hypertension: A Longitudinal Follow-Up Study of Dietary Sodium Intervention. Hypertension, 2021, 78, 155-164.	1.3	14
52	Metabolites Associated with Coffee Consumption and Incident Chronic Kidney Disease. Clinical Journal of the American Society of Nephrology: CJASN, 2021, 16, 1620-1629.	2.2	14
53	Blood pressure response to potassium supplementation is associated with genetic variation in endothelin 1 and interactions with E selectin in rural Chinese. Journal of Hypertension, 2010, 28, 748-755.	0.3	13
54	Genome-Wide Linkage and Positional Candidate Gene Study of Blood Pressure Response to Dietary Potassium Intervention. Circulation: Cardiovascular Genetics, 2010, 3, 539-547.	5.1	13

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55	Genome-Wide Linkage and Positional Association Analyses Identify Associations of Novel AFF3 and NTM Genes with Triglycerides: The GenSalt Study. Journal of Genetics and Genomics, 2015, 42, 107-117.	1.7	13
56	Novel associations between blood metabolites and kidney function among Bogalusa Heart Study and Multi-Ethnic Study of Atherosclerosis participants. Metabolomics, 2019, 15, 149.	1.4	13
57	Multi-ancestry genome-wide gene–sleep interactions identify novel loci for blood pressure. Molecular Psychiatry, 2021, 26, 6293-6304.	4.1	13
58	Atherosclerotic cardiovascular disease events among statin eligible individuals with and without long-term healthy arterial aging. Atherosclerosis, 2021, 326, 56-62.	0.4	13
59	Associations of Variants in the <i>CACNA1A </i> and <i>CACNA1C </i> Genes With Longitudinal Blood Pressure Changes and Hypertension Incidence: The GenSalt Study. American Journal of Hypertension, 2016, 29, 1301-1306.	1.0	12
60	Coronary Artery Calcium and the Age-Specific Competing Risk of Cardiovascular Versus Cancer Mortality: The Coronary Artery Calcium Consortium. American Journal of Medicine, 2020, 133, e575-e583.	0.6	12
61	Plasma Metabolomic Signatures of Healthy Dietary Patterns in the Chronic Renal Insufficiency Cohort (CRIC) Study. Journal of Nutrition, 2021, 151, 2894-2907.	1.3	12
62	Insights From a Large-Scale Whole-Genome Sequencing Study of Systolic Blood Pressure, Diastolic Blood Pressure, and Hypertension. Hypertension, 2022, 79, 1656-1667.	1.3	12
63	Resequencing Study Identifies Rare Renin–Angiotensin–Aldosterone System Variants Associated With Blood Pressure Salt-Sensitivity: The GenSalt Study. American Journal of Hypertension, 2017, 30, 495-501.	1.0	11
64	Serum metabolites associate with physical performance among middle-aged adults: Evidence from the Bogalusa Heart Study. Aging, 2020, 12, 11914-11941.	1.4	11
65	Fatty liver index and left ventricular mass: prospective associations from two independent cohorts. Journal of Hypertension, 2021, 39, 961-969.	0.3	10
66	Metabolomics study of blood pressure salt-sensitivity and hypertension. Nutrition, Metabolism and Cardiovascular Diseases, 2022, 32, 1681-1692.	1,1	10
67	Variation in Genes that Regulate Blood Pressure Are Associated with Glomerular Filtration Rate in Chinese. PLoS ONE, 2014, 9, e92468.	1.1	9
68	Genome-Wide Gene–Potassium Interaction Analyses on Blood Pressure. Circulation: Cardiovascular Genetics, 2017, 10, .	5.1	9
69	Left Ventricular Mass Index Is Associated With Cognitive Function in Middle-Age. Circulation: Cardiovascular Imaging, 2020, 13, e010335.	1.3	9
70	Maternal History of Hypertension and Blood Pressure Response to Potassium Intake: The GenSalt Study. American Journal of Epidemiology, 2012, 176, S55-S63.	1.6	8
71	Associations of Renin–Angiotensin–Aldosterone System Genes With Blood Pressure Changes and Hypertension Incidence. American Journal of Hypertension, 2015, 28, 1310-1315.	1.0	8
72	History of Asthma From Childhood and Arterial Stiffness in Asymptomatic Young Adults. Hypertension, 2018, 71, 928-936.	1.3	8

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73	Statistical tests for latent class in censored data due to detection limit. Statistical Methods in Medical Research, 2020, 29, 2179-2197.	0.7	8
74	Secular Trends in Cardiovascular Health in US Adults (from NHANES 2007 to 2018). American Journal of Cardiology, 2021, 159, 121-128.	0.7	8
75	Trans-ethnic genome-wide association study of blood metabolites in the Chronic Renal Insufficiency Cohort (CRIC) study. Kidney International, 2022, 101, 814-823.	2.6	8
76	Associations of the Serum/Glucocorticoid Regulated Kinase Genes With BP Changes and Hypertension Incidence: The Gensalt Study. American Journal of Hypertension, 2017, 30, 95-101.	1.0	7
77	Differential sex effects of systolic blood pressure and lowâ€density lipoprotein cholesterol on type 2 diabetes: Life course data from the Bogalusa Heart Study. Journal of Diabetes, 2018, 10, 449-457.	0.8	7
78	Serum metabolites associate with lipid phenotypes among Bogalusa Heart Study participants. Nutrition, Metabolism and Cardiovascular Diseases, 2020, 30, 777-787.	1.1	6
79	Pooled cohort equations heart failure risk score predicts cardiovascular disease and all-cause mortality in a nationally representative sample of US adults. BMC Cardiovascular Disorders, 2020, 20, 202.	0.7	6
80	Robust, flexible, and scalable tests for Hardy–Weinberg equilibrium across diverse ancestries. Genetics, 2021, 218, .	1.2	6
81	Genome-Wide Linkage and Regional Association Study of Blood Pressure Response to the Cold Pressor Test in Han Chinese. Circulation: Cardiovascular Genetics, 2014, 7, 521-528.	5.1	5
82	Associations of Endothelial System Genes With Blood Pressure Changes and Hypertension Incidence: The GenSalt Study. American Journal of Hypertension, 2015, 28, 780-788.	1.0	5
83	Examination of serum metabolome altered by cigarette smoking identifies novel metabolites mediating smokingâ€BMI association. Obesity, 2022, 30, 943-952.	1.5	5
84	Advances in Genomics Research of Blood Pressure Responses to Dietary Sodium and Potassium Intakes. Hypertension, 2021, 78, 4-15.	1.3	4
85	Association of Genome-Wide Polygenic Risk Score for Body Mass Index With Cardiometabolic Health From Childhood Through Midlife. Circulation Genomic and Precision Medicine, 2022, 15, .	1.6	4
86	Gene–Sodium Interaction and Blood Pressure: Findings from Genomics Research of Blood Pressure Salt Sensitivity. Progress in Molecular Biology and Translational Science, 2012, 108, 237-260.	0.9	3
87	Genetic variants of cGMP-dependent protein kinase genes and salt sensitivity of blood pressure: the GenSalt study. Journal of Human Hypertension, 2019, 33, 62-68.	1.0	3
88	Consumption of animal and plant foods and risk of left ventricular diastolic dysfunction: the Bogalusa Heart Study. ESC Heart Failure, 2020, 7, 2700-2710.	1.4	3
89	Association of Blood Pressure Genetic Risk Score with Cardiovascular Disease and CKD Progression: Findings from the CRIC Study. Kidney360, 2021, 2, 1251-1260.	0.9	3
90	Association of Kir genes with blood pressure responses to dietary sodium intervention: the GenSalt study. Hypertension Research, 2018, 41, 1045-1053.	1.5	2

Tanika N Kelly

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91	Discordantly normal ApoB relative to elevated LDL-C in persons with metabolic disorders: A marker of atherogenic heterogeneity. American Journal of Preventive Cardiology, 2021, 7, 100190.	1.3	2
92	Abstract 55: Sodium Sensitivity, Sodium Resistance, and Incidence of Hypertension. Circulation, 2020, 141, .	1.6	2
93	Association Between Baseline Buccal Telomere Length and Progression of Kidney Function: The Health and Retirement Study. Journals of Gerontology - Series A Biological Sciences and Medical Sciences, 2022, 77, 471-476.	1.7	2
94	Rare coding variants in RCN3 are associated with blood pressure. BMC Genomics, 2022, 23, 148.	1.2	2
95	Branched-chain amino acids, history of gestational diabetes, and breastfeeding: The Bogalusa Heart Study. Nutrition, Metabolism and Cardiovascular Diseases, 2020, 30, 2077-2084.	1.1	1
96	Race modifies the association between animal protein metabolite 1-methylhistidine and blood pressure in middle-aged adults: the Bogalusa Heart Study. Journal of Hypertension, 2020, 38, 2435-2442.	0.3	1
97	3384 Serum Metabolites from the Trimethylamine Pathway Associate with Left Ventricular Diastolic Function: The Bogalusa Heart Study. Journal of Clinical and Translational Science, 2019, 3, 53-54.	0.3	0