Nora Franceschini

List of Publications by Citations

145 papers	7,896 citations	41 h-index	87 g-index
163 ext. papers	10,995 ext. citations	10.8 avg, IF	4.56 L-index

#	Paper	IF	Citations
145	A comprehensive 1,000 Genomes-based genome-wide association meta-analysis of coronary artery disease. *Nature Genetics*, **2015**, 47, 1121-1130	36.3	1290
144	Loss-of-function mutations in APOC3, triglycerides, and coronary disease. *New England Journal of Medicine*, **2014**, 371, 22-31	59.2	721
143	Parent-of-origin-specific allelic associations among 106 genomic loci for age at menarche. *Nature*, **2014**, 514, 92-97	50.4	401
142	Sequencing of 53,831 diverse genomes from the NHLBI TOPMed Program. *Nature*, **2021**, 590, 290-299	50.4	268
141	Genomic analyses identify hundreds of variants associated with age at menarche and support a role for puberty timing in cancer risk. *Nature Genetics*, **2017**, 49, 834-841	36.3	257
140	The genetics of blood pressure regulation and its target organs from association studies in 342,415 individuals. *Nature Genetics*, **2016**, 48, 1171-1184	36.3	251
139	Large-scale genomic analyses link reproductive aging to hypothalamic signaling, breast cancer susceptibility and BRCA1-mediated DNA repair. *Nature Genetics*, **2015**, 47, 1294-1303	36.3	226
138	Genetic loci associated with chronic obstructive pulmonary disease overlap with loci for lung function and pulmonary fibrosis. *Nature Genetics*, **2017**, 49, 426-432	36.3	201
137	Meta-analysis of correlated traits via summary statistics from GWASs with an application in hypertension. *American Journal of Human Genetics*, **2015**, 96, 21-36	11	186
136	Fifteen new risk loci for coronary artery disease highlight arterial-wall-specific mechanisms. *Nature Genetics*, **2017**, 49, 1113-1119	36.3	184
135	Genetic Diversity and Association Studies in US Hispanic/Latino Populations: Applications in the Hispanic Community Health Study/Study of Latinos. *American Journal of Human Genetics*, **2016**, 98, 165-84	11	181
134	Whole-exome sequencing identifies rare and low-frequency coding variants associated with LDL cholesterol. *American Journal of Human Genetics*, **2014**, 94, 233-45	11	170
133	Meta-analysis identifies common and rare variants influencing blood pressure and overlapping with metabolic trait loci. *Nature Genetics*, **2016**, 48, 1162-70	36.3	152
132	Genome-wide association analysis of blood-pressure traits in African-ancestry individuals reveals common associated genes in African and non-African populations. *American Journal of Human Genetics*, **2013**, 93, 545-54	11	145
131	Genetic evidence for a normal-weight "metabolically obese" phenotype linking insulin resistance, hypertension, coronary artery disease, and type 2 diabetes. *Diabetes*, **2014**, 63, 4369-77	0.9	131
130	Gene-centric meta-analysis in 87,736 individuals of European ancestry identifies multiple blood-pressure-related loci. *American Journal of Human Genetics*, **2014**, 94, 349-60	11	131
129	Genome-wide association analysis identifies six new loci associated with forced vital capacity. *Nature Genetics*, **2014**, 46, 669-77	36.3	104

128	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	11.6	100
127	Genome-wide meta-analysis associates HLA-DQA1/DRB1 and LPA and lifestyle factors with human longevity. *Nature Communications*, **2017**, 8, 910	17.4	78
126	Uganda Genome Resource Enables Insights into Population History and Genomic Discovery in Africa. *Cell*, **2019**, 179, 984-1002.e36	56.2	76
125	NPHS2 gene, nephrotic syndrome and focal segmental glomerulosclerosis: a HuGE review. *Genetics in Medicine*, **2006**, 8, 63-75	8.1	71
124	Combined admixture mapping and association analysis identifies a novel blood pressure genetic locus on 5p13: contributions from the CARe consortium. *Human Molecular Genetics*, **2011**, 20, 2285-95	5.6	70
123	Trans-ethnic meta-regression of genome-wide association studies accounting for ancestry increases power for discovery and improves fine-mapping resolution. *Human Molecular Genetics*, **2017**, 26, 3639-3650	5.6	67
122	Cystatin C and Cardiovascular Disease: A Mendelian Randomization Study. *Journal of the American College of Cardiology*, **2016**, 68, 934-45	15.1	65
121	GWAS and colocalization analyses implicate carotid intima-media thickness and carotid plaque loci in cardiovascular outcomes. *Nature Communications*, **2018**, 9, 5141	17.4	64
120	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	36.3	59
119	A Large-Scale Multi-ancestry Genome-wide Study Accounting for Smoking Behavior Identifies Multiple Significant Loci for Blood Pressure. *American Journal of Human Genetics*, **2018**, 102, 375-400	11	59
118	Discovery and fine mapping of serum protein loci through transethnic meta-analysis. *American Journal of Human Genetics*, **2012**, 91, 744-53	11	58
117	Single-trait and multi-trait genome-wide association analyses identify novel loci for blood pressure in African-ancestry populations. *PLoS Genetics*, **2017**, 13, e1006728	6	58
116	Genome-wide Association Study of Platelet Count Identifies Ancestry-Specific Loci in Hispanic/Latino Americans. *American Journal of Human Genetics*, **2016**, 98, 229-42	11	54
115	Genome-Wide Association Study for Incident Myocardial Infarction and Coronary Heart Disease in Prospective Cohort Studies: The CHARGE Consortium. *PLoS ONE*, **2016**, 11, e0144997	3.7	53
114	Molecular mechanisms underlying variations in lung function: a systems genetics analysis. *Lancet Respiratory Medicine,the*, **2015**, 3, 782-95	35.1	52
113	Trans-ethnic kidney function association study reveals putative causal genes and effects on kidney-specific disease aetiologies. *Nature Communications*, **2019**, 10, 29	17.4	51
112	Association of genetic variants and incident coronary heart disease in multiethnic cohorts: the PAGE study. *Circulation: Cardiovascular Genetics*, **2011**, 4, 661-72		48
111	Orthostatic hypotension and incident chronic kidney disease: the atherosclerosis risk in communities study. *Hypertension*, **2010**, 56, 1054-9	8.5	46

110	Multiethnic meta-analysis identifies ancestry-specific and cross-ancestry loci for pulmonary function. *Nature Communications*, **2018**, 9, 2976	17.4	45
109	HDAC9 is implicated in atherosclerotic aortic calcification and affects vascular smooth muscle cell phenotype. *Nature Genetics*, **2019**, 51, 1580-1587	36.3	45
108	Leveraging population admixture to characterize the heritability of complex traits. *Nature Genetics*, **2014**, 46, 1356-62	36.3	45
107	Meta-analysis of loci associated with age at natural menopause in African-American women. *Human Molecular Genetics*, **2014**, 23, 3327-42	5.6	44
106	Trans-ethnic Fine Mapping Highlights Kidney-Function Genes Linked to Salt Sensitivity. *American Journal of Human Genetics*, **2016**, 99, 636-646	11	44
105	Large-scale genome-wide association studies and meta-analyses of longitudinal change in adult lung function. *PLoS ONE*, **2014**, 9, e100776	3.7	42
104	Multi-ancestry study of blood lipid levels identifies four loci interacting with physical activity. *Nature Communications*, **2019**, 10, 376	17.4	41
103	Interleukin-6 Signaling Effects on Ischemic Stroke and Other Cardiovascular Outcomes: A Mendelian Randomization Study. *Circulation Genomic and Precision Medicine*, **2020**, 13, e002872	5.2	41
102	Trans-ethnic Meta-analysis and Functional Annotation Illuminates the Genetic Architecture of Fasting Glucose and Insulin. *American Journal of Human Genetics*, **2016**, 99, 56-75	11	41
101	Associations of autozygosity with a broad range of human phenotypes. *Nature Communications*, **2019**, 10, 4957	17.4	40
100	A whole genome association study of mother-to-child transmission of HIV in Malawi. *Genome Medicine*, **2010**, 2, 17	14.4	40
99	Multiancestry Genome-Wide Association Study of Lipid Levels Incorporating Gene-Alcohol Interactions. *American Journal of Epidemiology*, **2019**, 188, 1033-1054	3.8	39
98	Phosphodiesterase 1 regulation is a key mechanism in vascular aging. *Clinical Science*, **2015**, 129, 1061-75	6.5	39
97	Strategies for Enriching Variant Coverage in Candidate Disease Loci on a Multiethnic Genotyping Array. *PLoS ONE*, **2016**, 11, e0167758	3.7	39
96	GWAS for male-pattern baldness identifies 71 susceptibility loci explaining 38% of the risk. *Nature Communications*, **2017**, 8, 1584	17.4	37
95	African Ancestry-Specific Alleles and Kidney Disease Risk in Hispanics/Latinos. *Journal of the American Society of Nephrology: JASN*, **2017**, 28, 915-922	12.7	37
94	The association of genetic variants of type 2 diabetes with kidney function. *Kidney International*, **2012**, 82, 220-5	9.9	36
93	Rare Exome Sequence Variants in CLCN6 Reduce Blood Pressure Levels and Hypertension Risk. *Circulation: Cardiovascular Genetics*, **2016**, 9, 64-70		35

92	Recent findings in the genetics of blood pressure and hypertension traits. *American Journal of Hypertension*, **2011**, 24, 392-400	2.3	35
91	Prevalence and Correlates of CKD in Hispanics/Latinos in the United States. *Clinical Journal of the American Society of Nephrology: CJASN*, **2015**, 10, 1757-66	6.9	34
90	Serum Calcification Propensity and Fetuin-A: Biomarkers of Cardiovascular Disease in Kidney Transplant Recipients. *American Journal of Nephrology*, **2018**, 48, 21-31	4.6	33
89	Prevention and treatment strategies for glucocorticoid-induced osteoporotic fractures. *Clinical Rheumatology*, **2007**, 26, 144-53	3.9	32
88	Novel genetic associations for blood pressure identified via gene-alcohol interaction in up to 570K individuals across multiple ancestries. *PLoS ONE*, **2018**, 13, e0198166	3.7	31
87	Multi-ancestry sleep-by-SNP interaction analysis in 126,926 individuals reveals lipid loci stratified by sleep duration. *Nature Communications*, **2019**, 10, 5121	17.4	31
86	Immunosuppression, hepatitis C infection, and acute renal failure in HIV-infected patients. *Journal of Acquired Immune Deficiency Syndromes (1999)*, **2006**, 42, 368-72	3.1	31
85	and Loci Identified through Large-Scale Exome Chip Analysis Regulate Kidney Development and Function. *Journal of the American Society of Nephrology: JASN*, **2017**, 28, 981-994	12.7	30
84	Cadmium body burden, hypertension, and changes in blood pressure over time: results from a prospective cohort study in American Indians. *Journal of the American Society of Hypertension*, **2018**, 12, 426-437.e9		30
83	Genetic insights into biological mechanisms governing human ovarian ageing. *Nature*, **2021**, 596, 393-397	50.4	28
82	Discovery of rare variants associated with blood pressure regulation through meta-analysis of 1.3 million individuals. *Nature Genetics*, **2020**, 52, 1314-1332	36.3	26
81	Genetics of hypertension: discoveries from the bench to human populations. *American Journal of Physiology - Renal Physiology*, **2014**, 306, F1-F11	4.3	24
80	Genetics, ancestry, and hypertension: implications for targeted antihypertensive therapies. *Current Hypertension Reports*, **2014**, 16, 461	4.7	24
79	Trans-ethnic fine-mapping of genetic loci for body mass index in the diverse ancestral populations of the Population Architecture using Genomics and Epidemiology (PAGE) Study reveals evidence for multiple signals at established loci. *Human Genetics*, **2017**, 136, 771-800	6.3	23
78	Sequence variation in telomerase reverse transcriptase (TERT) as a determinant of risk of cardiovascular disease: the Atherosclerosis Risk in Communities (ARIC) study. *BMC Medical Genetics*, **2015**, 16, 52	2.1	23
77	Rare coding variants and X-linked loci associated with age at menarche. *Nature Communications*, **2015**, 6, 7756	17.4	23
76	The effects of HIV type-1 viral suppression and non-viral factors on quantitative proteinuria in the highly active antiretroviral therapy era. *Antiviral Therapy*, **2009**, 14, 543-549	1.6	23
75	Replication of the effect of SLC2A9 genetic variation on serum uric acid levels in American Indians. *European Journal of Human Genetics*, **2014**, 22, 938-43	5.3	22

74	Cystatin C and preeclampsia: a case control study. *Renal Failure*, **2008**, 30, 89-95	2.9	22
73	Serum Uromodulin: A Biomarker of Long-Term Kidney Allograft Failure. *American Journal of Nephrology*, **2018**, 47, 275-282	4.6	21
72	Adiposity patterns and the risk for ESRD in postmenopausal women. *Clinical Journal of the American Society of Nephrology: CJASN*, **2015**, 10, 241-50	6.9	21
71	The association of the MYH9 gene and kidney outcomes in American Indians: the Strong Heart Family Study. *Human Genetics*, **2010**, 127, 295-301	6.3	19
70	Variant Discovery and Fine Mapping of Genetic Loci Associated with Blood Pressure Traits in Hispanics and African Americans. *PLoS ONE*, **2016**, 11, e0164132	3.7	19
69	Genome-wide association study identifies novel loci for type 2 diabetes-attributed end-stage kidney disease in African Americans. *Human Genomics*, **2019**, 13, 21	6.8	18
68	Admixture Mapping Identifies an Amerindian Ancestry Locus Associated with Albuminuria in Hispanics in the United States. *Journal of the American Society of Nephrology: JASN*, **2017**, 28, 2211-2220	12.7	17
67	Prospective associations of coronary heart disease loci in African Americans using the MetaboChip: the PAGE study. *PLoS ONE*, **2014**, 9, e113203	3.7	17
66	Admixture mapping in the Hispanic Community Health Study/Study of Latinos reveals regions of genetic associations with blood pressure traits. *PLoS ONE*, **2017**, 12, e0188400	3.7	17
65	A quantitative trait loci-specific gene-by-sex interaction on systolic blood pressure among American Indians: the Strong Heart Family Study. *Hypertension*, **2006**, 48, 266-70	8.5	16
64	Meta-analysis of exome array data identifies six novel genetic loci for lung function. *Wellcome Open Research*, **2018**, 3, 4	4.8	16
63	Generalization of associations of kidney-related genetic loci to American Indians. *Clinical Journal of the American Society of Nephrology: CJASN*, **2014**, 9, 150-8	6.9	15
62	Diabetes Mellitus, Glycemic Traits, and Cerebrovascular Disease: A Mendelian Randomization Study. *Neurology*, **2021**, 96, e1732-e1742	6.5	15
61	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	6.3	14
60	Genetics of Chronic Kidney Disease Stages Across Ancestries: The PAGE Study. *Frontiers in Genetics*, **2019**, 10, 494	4.5	14
59	Linkage Analysis of Urine Arsenic Species Patterns in the Strong Heart Family Study. *Toxicological Sciences*, **2015**, 148, 89-100	4.4	14
58	The genetic underpinnings of variation in ages at menarche and natural menopause among women from the multi-ethnic Population Architecture using Genomics and Epidemiology (PAGE) Study: A trans-ethnic meta-analysis. *PLoS ONE*, **2018**, 13, e0200486	3.7	14
57	The association of the vanin-1 N131S variant with blood pressure is mediated by endoplasmic reticulum-associated degradation and loss of function. *PLoS Genetics*, **2014**, 10, e1004641	6	14

56	Genome-wide linkage analysis of pulse pressure in American Indians: the Strong Heart Study. *American Journal of Hypertension*, **2008**, 21, 194-9	2.3	14
55	Genome-Wide Association Study of Blood Pressure Traits by Hispanic/Latino Background: the Hispanic Community Health Study/Study of Latinos. *Scientific Reports*, **2017**, 7, 10348	4.9	13
54	A genome-wide association study meta-analysis of clinical fracture in 10,012 African American women. *Bone Reports*, **2016**, 5, 233-242	2.6	12
53	Genetic variants associated with earlier age at menopause increase the risk of cardiovascular events in women. *Menopause*, **2018**, 25, 451-457	2.5	12
52	Association of APOL1 With Heart Failure With Preserved Ejection Fraction in Postmenopausal African American Women. *JAMA Cardiology*, **2018**, 3, 712-720	16.2	12
51	Genetic Testing in Clinical Settings. *American Journal of Kidney Diseases*, **2018**, 72, 569-581	7.4	11
50	Diabetes-specific genetic effects on obesity traits in American Indian populations: the Strong Heart Family Study. *BMC Medical Genetics*, **2008**, 9, 90	2.1	11
49	Rare variants in fox-1 homolog A (RBFOX1) are associated with lower blood pressure. *PLoS Genetics* , **2017**, 13, e1006678	6	11
48	Role of Rare and Low-Frequency Variants in Gene-Alcohol Interactions on Plasma Lipid Levels. *Circulation Genomic and Precision Medicine*, **2020**, 13, e002772	5.2	8
47	Smoking patterns and chronic kidney disease in US Hispanics: Hispanic Community Health Study/Study of Latinos. *Nephrology Dialysis Transplantation*, **2016**, 31, 1670-6	4.3	8
46	Low agreement between modified-Schwartz and CKD-EPI eGFR in young adults: a retrospective longitudinal cohort study. *BMC Nephrology*, **2018**, 19, 194	2.7	7
45	Cholesteryl ester transfer protein (CETP) as a drug target for cardiovascular disease. *Nature Communications*, **2021**, 12, 5640	17.4	7
44	Interleukin-6 signaling effects on ischemic stroke and other cardiovascular outcomes: a Mendelian Randomization study		7
43	Variants Associated with the Ankle Brachial Index Differ by Hispanic/Latino Ethnic Group: a genome-wide association study in the Hispanic Community Health Study/Study of Latinos. *Scientific Reports*, **2019**, 9, 11410	4.9	6
42	Analysis of Sequence Data Under Multivariate Trait-Dependent Sampling. *Journal of the American Statistical Association*, **2015**, 110, 560-572	2.8	6
41	Genetic diversity is a predictor of mortality in humans. *BMC Genetics*, **2014**, 15, 159	2.6	6
40	Social- and behavioral-specific genetic effects on blood pressure traits: the Strong Heart Family Study. *Circulation: Cardiovascular Genetics*, **2009**, 2, 396-401		6
39	Analyses of biomarker traits in diverse UK biobank participants identify associations missed by European-centric analysis strategies. *Journal of Human Genetics*, **2021**,	4.3	6

38	Current Smoking Raises Risk of Incident Hypertension: Hispanic Community Health Study-Study of Latinos. *American Journal of Hypertension*, **2021**, 34, 190-197	2.3	4
37	Plasma Protein Profile of Carotid Artery Atherosclerosis and Atherosclerotic Outcomes: Meta-Analyses and Mendelian Randomization Analyses. *Arteriosclerosis, Thrombosis, and Vascular Biology*, **2021**, 41, 1777-1788	9.4	4
36	Genetic Variants Related to Cardiometabolic Traits Are Associated to B Cell Function, Insulin Resistance, and Diabetes Among AmeriCan Indians: The Strong Heart Family Study. *Frontiers in Genetics*, **2018**, 9, 466	4.5	4
35	Incident Chronic Kidney Disease Risk among Hispanics/Latinos in the United States: The Hispanic Community Health Study/Study of Latinos (HCHS/SOL). *Journal of the American Society of Nephrology: JASN*, **2020**, 31, 1315-1324	12.7	3
34	Arsenic-gene interactions and beta-cell function in the Strong Heart Family Study. *Toxicology and Applied Pharmacology*, **2018**, 348, 123-129	4.6	3
33	Genome-wide association of trajectories of systolic blood pressure change. *BMC Proceedings*, **2016**, 10, 321-327	2.3	3
32	Mapping of a blood pressure QTL on chromosome 17 in American Indians of the strong heart family study. *BMC Cardiovascular Disorders*, **2014**, 14, 158	2.3	3
31	Discovery and fine-mapping of kidney function loci in first genome-wide association study in Africans. *Human Molecular Genetics*, **2021**, 30, 1559-1568	5.6	3
30	Epigenome-wide association study of kidney function identifies trans-ethnic and ethnic-specific loci. *Genome Medicine*, **2021**, 13, 74	14.4	3
29	Multi-ancestry genome-wide gene-sleep interactions identify novel loci for blood pressure. *Molecular Psychiatry*, **2021**,	15.1	3
28	Engulfment and cell motility 1 (ELMO1) and apolipoprotein A1 (APOA1) as candidate genes for sickle cell nephropathy. *British Journal of Haematology*, **2021**, 193, 628-632	4.5	3
27	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	8.8	3
26	Association of Sickle Cell Trait With Incidence of Coronary Heart Disease Among African American Individuals. *JAMA Network Open*, **2021**, 4, e2030435	10.4	3
25	Genetic analysis of hsCRP in American Indians: The Strong Heart Family Study. *PLoS ONE*, **2019**, 14, e0223574	3.7	2
24	Association of Variants with Blood Pressure in the Atherosclerosis Risk in Communities Study. *International Journal of Hypertension*, **2019**, 2019, 2137629	2.4	2
23	Rare variant associations with waist-to-hip ratio in European-American and African-American women from the NHLBI-Exome Sequencing Project. *European Journal of Human Genetics*, **2016**, 24, 1181-7	5.3	2
22	Effect of Sickle Cell Trait and Genotype on the Association of Soluble uPAR with Kidney Function Measures in Black Americans. *Clinical Journal of the American Society of Nephrology: CJASN*, **2021**, 16, 287-289	6.9	2
21	When the At-Risk Do Not Develop Heart Failure: Understanding Positive Deviance Among Postmenopausal African American and Hispanic Women. *Journal of Cardiac Failure*, **2021**, 27, 217-223	3.3	2

20	Genome-Wide Association of Kidney Traits in Hispanics/Latinos Using Dense Imputed Whole-Genome Sequencing Data: The Hispanic Community Health Study/Study of Latinos. *Circulation Genomic and Precision Medicine*, **2020**, 13, e002891	5.2	1
19	Genome-wide association study and meta-analysis identify loci associated with ventricular and supraventricular ectopy. *Scientific Reports*, **2018**, 8, 5675	4.9	1
18	The association of cell cycle checkpoint 2 variants and kidney function: findings of the Family Blood Pressure Program and the Atherosclerosis Risk In Communities study. *American Journal of Hypertension*, **2009**, 22, 552-8	2.3	1
17	Integrative analysis of 3604 GWAS reveals multiple novel cell type-specific regulatory associations.. *Genome Biology*, **2022**, 23, 13	18.3	1
16	Genome-Wide Admixture Mapping of Estimated Glomerular Filtration Rate and Chronic Kidney Disease Identify European and African Ancestry-of-Origin Loci in Hispanic and Latino Individuals in the United States. *Journal of the American Society of Nephrology: JASN*, **2021**,	12.7	1
15	Soluble Urokinase Plasminogen Activator Receptor: Genetic Variation and Cardiovascular Disease Risk in Blacks. *Circulation Genomic and Precision Medicine*, **2021**, CIRCGEN121003421	5.2	1
14	Genetic variation and urine cadmium levels: ABCC1 effects in the Strong Heart Family Study. *Environmental Pollution*, **2021**, 276, 116717	9.3	1
13	DNAm-based signatures of accelerated aging and mortality in blood are associated with low renal function. *Clinical Epigenetics*, **2021**, 13, 121	7.7	1
12	Fine mapping and identification of serum urate loci in American Indians: The Strong Heart Family Study. *Scientific Reports*, **2019**, 9, 17899	4.9	1
11	Trans-ethnic Mendelian-randomization study reveals causal relationships between cardiometabolic factors and chronic kidney disease. *International Journal of Epidemiology*, **2021**,	7.8	1
10	Comparison of 2 models for gene-environment interactions: an example of simulated gene-medication interactions on systolic blood pressure in family-based data. *BMC Proceedings*, **2016**, 10, 371-377	2.3	0
9	Comparison of strategies for identification of regulatory quantitative trait loci of transcript expression traits. *BMC Proceedings*, **2007**, 1 Suppl 1, S85	2.3	0
8	Demographic and sociocultural risk factors for adulthood weight gain in Hispanic/Latinos: results from the Hispanic Community Health Study / Study of Latinos (HCHS/SOL). *BMC Public Health*, **2021**, 21, 2064	4.1	0
7	Sedentary Behavior and Change in Kidney Function: The Hispanic Community Health Study/Study of Latinos (HCHS/SOL). *Kidney360*, **2021**, 2, 245-253	1.8	0
6	Multiethnic Genome-Wide Association Study of Subclinical Atherosclerosis in Individuals With Type 2 Diabetes. *Circulation Genomic and Precision Medicine*, **2021**, 14, e003258	5.2	0
5	Reply to ‘Misestimation of heritability and prediction accuracy of male-pattern baldness’. *Nature Communications*, **2018**, 9, 2538	17.4	
4	, Sickle Cell Trait, and CKD in the Jackson Heart Study.. *Kidney Medicine*, **2021**, 3, 962-973.e1	2.8	
3	Rare coding variants in RCN3 are associated with blood pressure.. *BMC Genomics*, **2022**, 23, 148	4.5	

2	Racial disparities in COVID-19 test positivity among people living with HIV in the United States.. *International Journal of STD and AIDS*, **2022**, 9564624221074468	1.4
1	Hepatitis C infection and chronic kidney disease among Hispanics/Latinos. *Medicine (United States)*, **2021**, 100, e28089	1.8