

Nora Franceschini

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.

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	Integrative analysis of 3604 GWAS reveals multiple novel cell type-specific regulatory associations.. <i>Genome Biology</i> , 2022 , 23, 13	18.3	1
144	Rare coding variants in RCN3 are associated with blood pressure.. <i>BMC Genomics</i> , 2022 , 23, 148	4.5	
143	Racial disparities in COVID-19 test positivity among people living with HIV in the United States.. <i>International Journal of STD and AIDS</i> , 2022 , 9564624221074468	1.4	
142	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. <i>Journal of the American Society of Nephrology: JASN</i> , 2021 ,	12.7	1
141	Demographic and sociocultural risk factors for adulthood weight gain in Hispanic/Latinos: results from the Hispanic Community Health Study / Study of Latinos (HCHS/SOL). <i>BMC Public Health</i> , 2021 , 21, 2064	4.1	0
140	Effect of Sickle Cell Trait and Genotype on the Association of Soluble uPAR with Kidney Function Measures in Black Americans. <i>Clinical Journal of the American Society of Nephrology: CJASN</i> , 2021 , 16, 287-289	6.9	2
139	Cholesteryl ester transfer protein (CETP) as a drug target for cardiovascular disease. <i>Nature Communications</i> , 2021 , 12, 5640	17.4	7
138	Soluble Urokinase Plasminogen Activator Receptor: Genetic Variation and Cardiovascular Disease Risk in Blacks. <i>Circulation Genomic and Precision Medicine</i> , 2021 , CIRCGEN121003421	5.2	1
137	Sedentary Behavior and Change in Kidney Function: The Hispanic Community Health Study/Study of Latinos (HCHS/SOL). <i>Kidney360</i> , 2021 , 2, 245-253	1.8	0
136	Current Smoking Raises Risk of Incident Hypertension: Hispanic Community Health Study-Study of Latinos. <i>American Journal of Hypertension</i> , 2021 , 34, 190-197	2.3	4
135	Discovery and fine-mapping of kidney function loci in first genome-wide association study in Africans. <i>Human Molecular Genetics</i> , 2021 , 30, 1559-1568	5.6	3
134	Epigenome-wide association study of kidney function identifies trans-ethnic and ethnic-specific loci. <i>Genome Medicine</i> , 2021 , 13, 74	14.4	3
133	Multi-ancestry genome-wide gene-sleep interactions identify novel loci for blood pressure. <i>Molecular Psychiatry</i> , 2021 ,	15.1	3
132	Genetic variation and urine cadmium levels: ABCC1 effects in the Strong Heart Family Study. <i>Environmental Pollution</i> , 2021 , 276, 116717	9.3	1
131	Plasma Protein Profile of Carotid Artery Atherosclerosis and Atherosclerotic Outcomes: Meta-Analyses and Mendelian Randomization Analyses. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2021 , 41, 1777-1788	9.4	4
130	DNAm-based signatures of accelerated aging and mortality in blood are associated with low renal function. <i>Clinical Epigenetics</i> , 2021 , 13, 121	7.7	1
129	Engulfment and cell motility 1 (ELMO1) and apolipoprotein A1 (APOA1) as candidate genes for sickle cell nephropathy. <i>British Journal of Haematology</i> , 2021 , 193, 628-632	4.5	3

128	Whole genome sequence analyses of eGFR in 23,732 people representing multiple ancestries in the NHLBI trans-omics for precision medicine (TOPMed) consortium. <i>EBioMedicine</i> , 2021 , 63, 103157	8.8	3
127	Diabetes Mellitus, Glycemic Traits, and Cerebrovascular Disease: A Mendelian Randomization Study. <i>Neurology</i> , 2021 , 96, e1732-e1742	6.5	15
126	When the At-Risk Do Not Develop Heart Failure: Understanding Positive Deviance Among Postmenopausal African American and Hispanic Women. <i>Journal of Cardiac Failure</i> , 2021 , 27, 217-223	3.3	2
125	Sequencing of 53,831 diverse genomes from the NHLBI TOPMed Program. <i>Nature</i> , 2021 , 590, 290-299	50.4	268
124	, Sickle Cell Trait, and CKD in the Jackson Heart Study.. <i>Kidney Medicine</i> , 2021 , 3, 962-973.e1	2.8	
123	Multiethnic Genome-Wide Association Study of Subclinical Atherosclerosis in Individuals With Type 2 Diabetes. <i>Circulation Genomic and Precision Medicine</i> , 2021 , 14, e003258	5.2	0
122	Genetic insights into biological mechanisms governing human ovarian ageing. <i>Nature</i> , 2021 , 596, 393-397	50.4	28
121	Analyses of biomarker traits in diverse UK biobank participants identify associations missed by European-centric analysis strategies. <i>Journal of Human Genetics</i> , 2021 ,	4.3	6
120	Association of Sickle Cell Trait With Incidence of Coronary Heart Disease Among African American Individuals. <i>JAMA Network Open</i> , 2021 , 4, e2030435	10.4	3
119	Trans-ethnic Mendelian-randomization study reveals causal relationships between cardiometabolic factors and chronic kidney disease. <i>International Journal of Epidemiology</i> , 2021 ,	7.8	1
118	Hepatitis C infection and chronic kidney disease among Hispanics/Latinos. <i>Medicine (United States)</i> , 2021 , 100, e28089	1.8	
117	Interleukin-6 Signaling Effects on Ischemic Stroke and Other Cardiovascular Outcomes: A Mendelian Randomization Study. <i>Circulation Genomic and Precision Medicine</i> , 2020 , 13, e002872	5.2	41
116	Incident Chronic Kidney Disease Risk among Hispanics/Latinos in the United States: The Hispanic Community Health Study/Study of Latinos (HCHS/SOL). <i>Journal of the American Society of Nephrology: JASN</i> , 2020 , 31, 1315-1324	12.7	3
115	Role of Rare and Low-Frequency Variants in Gene-Alcohol Interactions on Plasma Lipid Levels. <i>Circulation Genomic and Precision Medicine</i> , 2020 , 13, e002772	5.2	8
114	Genome-Wide Association of Kidney Traits in Hispanics/Latinos Using Dense Imputed Whole-Genome Sequencing Data: The Hispanic Community Health Study/Study of Latinos. <i>Circulation Genomic and Precision Medicine</i> , 2020 , 13, e002891	5.2	1
113	Discovery of rare variants associated with blood pressure regulation through meta-analysis of 1.3 million individuals. <i>Nature Genetics</i> , 2020 , 52, 1314-1332	36.3	26
112	Genetic analysis of hsCRP in American Indians: The Strong Heart Family Study. <i>PLoS ONE</i> , 2019 , 14, e0223574	5.74	2
111	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. <i>Scientific Reports</i> , 2019 , 9, 11410	4.9	6

110	Multiancestry Genome-Wide Association Study of Lipid Levels Incorporating Gene-Alcohol Interactions. <i>American Journal of Epidemiology</i> , 2019 , 188, 1033-1054	3.8	39
109	Multi-ancestry study of blood lipid levels identifies four loci interacting with physical activity. <i>Nature Communications</i> , 2019 , 10, 376	17.4	41
108	Leveraging linkage evidence to identify low-frequency and rare variants on 16p13 associated with blood pressure using TOPMed whole genome sequencing data. <i>Human Genetics</i> , 2019 , 138, 199-210	6.3	14
107	Genetics of Chronic Kidney Disease Stages Across Ancestries: The PAGE Study. <i>Frontiers in Genetics</i> , 2019 , 10, 494	4.5	14
106	Genome-wide association study identifies novel loci for type 2 diabetes-attributed end-stage kidney disease in African Americans. <i>Human Genomics</i> , 2019 , 13, 21	6.8	18
105	Association of Variants with Blood Pressure in the Atherosclerosis Risk in Communities Study. <i>International Journal of Hypertension</i> , 2019 , 2019, 2137629	2.4	2
104	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
103	Uganda Genome Resource Enables Insights into Population History and Genomic Discovery in Africa. <i>Cell</i> , 2019 , 179, 984-1002.e36	56.2	76
102	Associations of autozygosity with a broad range of human phenotypes. <i>Nature Communications</i> , 2019 , 10, 4957	17.4	40
101	Multi-ancestry sleep-by-SNP interaction analysis in 126,926 individuals reveals lipid loci stratified by sleep duration. <i>Nature Communications</i> , 2019 , 10, 5121	17.4	31
100	HDAC9 is implicated in atherosclerotic aortic calcification and affects vascular smooth muscle cell phenotype. <i>Nature Genetics</i> , 2019 , 51, 1580-1587	36.3	45
99	Fine mapping and identification of serum urate loci in American Indians: The Strong Heart Family Study. <i>Scientific Reports</i> , 2019 , 9, 17899	4.9	1
98	Trans-ethnic kidney function association study reveals putative causal genes and effects on kidney-specific disease aetiologies. <i>Nature Communications</i> , 2019 , 10, 29	17.4	51
97	Arsenic-gene interactions and beta-cell function in the Strong Heart Family Study. <i>Toxicology and Applied Pharmacology</i> , 2018 , 348, 123-129	4.6	3
96	Genome-wide association study and meta-analysis identify loci associated with ventricular and supraventricular ectopy. <i>Scientific Reports</i> , 2018 , 8, 5675	4.9	1
95	Genetic Testing in Clinical Settings. <i>American Journal of Kidney Diseases</i> , 2018 , 72, 569-581	7.4	11
94	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
93	Serum Uromodulin: A Biomarker of Long-Term Kidney Allograft Failure. <i>American Journal of Nephrology</i> , 2018 , 47, 275-282	4.6	21

92	Cadmium body burden, hypertension, and changes in blood pressure over time: results from a prospective cohort study in American Indians. <i>Journal of the American Society of Hypertension</i> , 2018 , 12, 426-437.e9		30
91	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. <i>PLoS ONE</i> , 2018 , 13, e0200486	3.7	14
90	Serum Calcification Propensity and Fetuin-A: Biomarkers of Cardiovascular Disease in Kidney Transplant Recipients. <i>American Journal of Nephrology</i> , 2018 , 48, 21-31	4.6	33
89	Multiethnic meta-analysis identifies ancestry-specific and cross-ancestry loci for pulmonary function. <i>Nature Communications</i> , 2018 , 9, 2976	17.4	45
88	Low agreement between modified-Schwartz and CKD-EPI eGFR in young adults: a retrospective longitudinal cohort study. <i>BMC Nephrology</i> , 2018 , 19, 194	2.7	7
87	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
86	Meta-analysis of exome array data identifies six novel genetic loci for lung function. <i>Wellcome Open Research</i> , 2018 , 3, 4	4.8	16
85	Genetic variants associated with earlier age at menopause increase the risk of cardiovascular events in women. <i>Menopause</i> , 2018 , 25, 451-457	2.5	12
84	GWAS and colocalization analyses implicate carotid intima-media thickness and carotid plaque loci in cardiovascular outcomes. <i>Nature Communications</i> , 2018 , 9, 5141	17.4	64
83	Genetic Variants Related to Cardiometabolic Traits Are Associated to B Cell Function, Insulin Resistance, and Diabetes Among American Indians: The Strong Heart Family Study. <i>Frontiers in Genetics</i> , 2018 , 9, 466	4.5	4
82	Association of APOL1 With Heart Failure With Preserved Ejection Fraction in Postmenopausal African American Women. <i>JAMA Cardiology</i> , 2018 , 3, 712-720	16.2	12
81	Reply to Misestimation of heritability and prediction accuracy of male-pattern baldness. <i>Nature Communications</i> , 2018 , 9, 2538	17.4	
80	Admixture Mapping Identifies an Amerindian Ancestry Locus Associated with Albuminuria in Hispanics in the United States. <i>Journal of the American Society of Nephrology: JASN</i> , 2017 , 28, 2211-2220	12.7	17
79	Genetic loci associated with chronic obstructive pulmonary disease overlap with loci for lung function and pulmonary fibrosis. <i>Nature Genetics</i> , 2017 , 49, 426-432	36.3	201
78	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. <i>Human Genetics</i> , 2017 , 136, 771-800	6.3	23
77	Genomic analyses identify hundreds of variants associated with age at menarche and support a role for puberty timing in cancer risk. <i>Nature Genetics</i> , 2017 , 49, 834-841	36.3	257
76	Fifteen new risk loci for coronary artery disease highlight arterial-wall-specific mechanisms. <i>Nature Genetics</i> , 2017 , 49, 1113-1119	36.3	184
75	and Loci Identified through Large-Scale Exome Chip Analysis Regulate Kidney Development and Function. <i>Journal of the American Society of Nephrology: JASN</i> , 2017 , 28, 981-994	12.7	30

74	Genome-wide meta-analysis associates HLA-DQA1/DRB1 and LPA and lifestyle factors with human longevity. <i>Nature Communications</i> , 2017 , 8, 910	17.4	78
73	Genome-Wide Association Study of Blood Pressure Traits by Hispanic/Latino Background: the Hispanic Community Health Study/Study of Latinos. <i>Scientific Reports</i> , 2017 , 7, 10348	4.9	13
72	GWAS for male-pattern baldness identifies 71 susceptibility loci explaining 38% of the risk. <i>Nature Communications</i> , 2017 , 8, 1584	17.4	37
71	African Ancestry-Specific Alleles and Kidney Disease Risk in Hispanics/Latinos. <i>Journal of the American Society of Nephrology: JASN</i> , 2017 , 28, 915-922	12.7	37
70	Trans-ethnic meta-regression of genome-wide association studies accounting for ancestry increases power for discovery and improves fine-mapping resolution. <i>Human Molecular Genetics</i> , 2017 , 26, 3639-3650	5.6	67
69	Rare variants in fox-1 homolog A (RFX1) are associated with lower blood pressure. <i>PLoS Genetics</i> , 2017 , 13, e1006678	6	11
68	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
67	Admixture mapping in the Hispanic Community Health Study/Study of Latinos reveals regions of genetic associations with blood pressure traits. <i>PLoS ONE</i> , 2017 , 12, e0188400	3.7	17
66	Meta-analysis identifies common and rare variants influencing blood pressure and overlapping with metabolic trait loci. <i>Nature Genetics</i> , 2016 , 48, 1162-70	36.3	152
65	The genetics of blood pressure regulation and its target organs from association studies in 342,415 individuals. <i>Nature Genetics</i> , 2016 , 48, 1171-1184	36.3	251
64	Genome-wide association of trajectories of systolic blood pressure change. <i>BMC Proceedings</i> , 2016 , 10, 321-327	2.3	3
63	Comparison of 2 models for gene-environment interactions: an example of simulated gene-medication interactions on systolic blood pressure in family-based data. <i>BMC Proceedings</i> , 2016 , 10, 371-377	2.3	0
62	Trans-ethnic Meta-analysis and Functional Annotation Illuminates the Genetic Architecture of Fasting Glucose and Insulin. <i>American Journal of Human Genetics</i> , 2016 , 99, 56-75	11	41
61	Rare variant associations with waist-to-hip ratio in European-American and African-American women from the NHLBI-Exome Sequencing Project. <i>European Journal of Human Genetics</i> , 2016 , 24, 1181-1187	5.3	2
60	Rare Exome Sequence Variants in CLCN6 Reduce Blood Pressure Levels and Hypertension Risk. <i>Circulation: Cardiovascular Genetics</i> , 2016 , 9, 64-70		35
59	Genome-wide Association Study of Platelet Count Identifies Ancestry-Specific Loci in Hispanic/Latino Americans. <i>American Journal of Human Genetics</i> , 2016 , 98, 229-42	11	54
58	Genetic Diversity and Association Studies in US Hispanic/Latino Populations: Applications in the Hispanic Community Health Study/Study of Latinos. <i>American Journal of Human Genetics</i> , 2016 , 98, 165-84	11	181
57	Investigating the Causal Relationship of C-Reactive Protein with 32 Complex Somatic and Psychiatric Outcomes: A Large-Scale Cross-Consortium Mendelian Randomization Study. <i>PLoS Medicine</i> , 2016 , 13, e1001976	11.6	100

56	Strategies for Enriching Variant Coverage in Candidate Disease Loci on a Multiethnic Genotyping Array. <i>PLoS ONE</i> , 2016 , 11, e0167758	3.7	39
55	Genome-Wide Association Study for Incident Myocardial Infarction and Coronary Heart Disease in Prospective Cohort Studies: The CHARGE Consortium. <i>PLoS ONE</i> , 2016 , 11, e0144997	3.7	53
54	Variant Discovery and Fine Mapping of Genetic Loci Associated with Blood Pressure Traits in Hispanics and African Americans. <i>PLoS ONE</i> , 2016 , 11, e0164132	3.7	19
53	Trans-ethnic Fine Mapping Highlights Kidney-Function Genes Linked to Salt Sensitivity. <i>American Journal of Human Genetics</i> , 2016 , 99, 636-646	11	44
52	Cystatin C and Cardiovascular Disease: A Mendelian Randomization Study. <i>Journal of the American College of Cardiology</i> , 2016 , 68, 934-45	15.1	65
51	A genome-wide association study meta-analysis of clinical fracture in 10,012 African American women. <i>Bone Reports</i> , 2016 , 5, 233-242	2.6	12
50	Smoking patterns and chronic kidney disease in US Hispanics: Hispanic Community Health Study/Study of Latinos. <i>Nephrology Dialysis Transplantation</i> , 2016 , 31, 1670-6	4.3	8
49	Molecular mechanisms underlying variations in lung function: a systems genetics analysis. <i>Lancet Respiratory Medicine</i> , 2015 , 3, 782-95	35.1	52
48	Large-scale genomic analyses link reproductive aging to hypothalamic signaling, breast cancer susceptibility and BRCA1-mediated DNA repair. <i>Nature Genetics</i> , 2015 , 47, 1294-1303	36.3	226
47	Linkage Analysis of Urine Arsenic Species Patterns in the Strong Heart Family Study. <i>Toxicological Sciences</i> , 2015 , 148, 89-100	4.4	14
46	Analysis of Sequence Data Under Multivariate Trait-Dependent Sampling. <i>Journal of the American Statistical Association</i> , 2015 , 110, 560-572	2.8	6
45	Sequence variation in telomerase reverse transcriptase (TERT) as a determinant of risk of cardiovascular disease: the Atherosclerosis Risk in Communities (ARIC) study. <i>BMC Medical Genetics</i> , 2015 , 16, 52	2.1	23
44	Prevalence and Correlates of CKD in Hispanics/Latinos in the United States. <i>Clinical Journal of the American Society of Nephrology: CJASN</i> , 2015 , 10, 1757-66	6.9	34
43	A comprehensive 1,000 Genomes-based genome-wide association meta-analysis of coronary artery disease. <i>Nature Genetics</i> , 2015 , 47, 1121-1130	36.3	1290
42	Phosphodiesterase 1 regulation is a key mechanism in vascular aging. <i>Clinical Science</i> , 2015 , 129, 1061-75	5.5	39
41	Rare coding variants and X-linked loci associated with age at menarche. <i>Nature Communications</i> , 2015 , 6, 7756	17.4	23
40	Adiposity patterns and the risk for ESRD in postmenopausal women. <i>Clinical Journal of the American Society of Nephrology: CJASN</i> , 2015 , 10, 241-50	6.9	21
39	Meta-analysis of correlated traits via summary statistics from GWASs with an application in hypertension. <i>American Journal of Human Genetics</i> , 2015 , 96, 21-36	11	186

38	Whole-exome sequencing identifies rare and low-frequency coding variants associated with LDL cholesterol. <i>American Journal of Human Genetics</i> , 2014 , 94, 233-45	11	170
37	Leveraging population admixture to characterize the heritability of complex traits. <i>Nature Genetics</i> , 2014 , 46, 1356-62	36.3	45
36	Genome-wide association analysis identifies six new loci associated with forced vital capacity. <i>Nature Genetics</i> , 2014 , 46, 669-77	36.3	104
35	Genetics of hypertension: discoveries from the bench to human populations. <i>American Journal of Physiology - Renal Physiology</i> , 2014 , 306, F1-F11	4.3	24
34	Parent-of-origin-specific allelic associations among 106 genomic loci for age at menarche. <i>Nature</i> , 2014 , 514, 92-97	50.4	401
33	Genetics, ancestry, and hypertension: implications for targeted antihypertensive therapies. <i>Current Hypertension Reports</i> , 2014 , 16, 461	4.7	24
32	Meta-analysis of loci associated with age at natural menopause in African-American women. <i>Human Molecular Genetics</i> , 2014 , 23, 3327-42	5.6	44
31	Loss-of-function mutations in APOC3, triglycerides, and coronary disease. <i>New England Journal of Medicine</i> , 2014 , 371, 22-31	59.2	721
30	Large-scale genome-wide association studies and meta-analyses of longitudinal change in adult lung function. <i>PLoS ONE</i> , 2014 , 9, e100776	3.7	42
29	Mapping of a blood pressure QTL on chromosome 17 in American Indians of the strong heart family study. <i>BMC Cardiovascular Disorders</i> , 2014 , 14, 158	2.3	3
28	Replication of the effect of SLC2A9 genetic variation on serum uric acid levels in American Indians. <i>European Journal of Human Genetics</i> , 2014 , 22, 938-43	5.3	22
27	The association of the vanin-1 N131S variant with blood pressure is mediated by endoplasmic reticulum-associated degradation and loss of function. <i>PLoS Genetics</i> , 2014 , 10, e1004641	6	14
26	Genetic evidence for a normal-weight "metabolically obese" phenotype linking insulin resistance, hypertension, coronary artery disease, and type 2 diabetes. <i>Diabetes</i> , 2014 , 63, 4369-77	0.9	131
25	Genetic diversity is a predictor of mortality in humans. <i>BMC Genetics</i> , 2014 , 15, 159	2.6	6
24	Generalization of associations of kidney-related genetic loci to American Indians. <i>Clinical Journal of the American Society of Nephrology: CJASN</i> , 2014 , 9, 150-8	6.9	15
23	Gene-centric meta-analysis in 87,736 individuals of European ancestry identifies multiple blood-pressure-related loci. <i>American Journal of Human Genetics</i> , 2014 , 94, 349-60	11	131
22	Prospective associations of coronary heart disease loci in African Americans using the MetaboChip: the PAGE study. <i>PLoS ONE</i> , 2014 , 9, e113203	3.7	17
21	Genome-wide association analysis of blood-pressure traits in African-ancestry individuals reveals common associated genes in African and non-African populations. <i>American Journal of Human Genetics</i> , 2013 , 93, 545-54	11	145

20	Discovery and fine mapping of serum protein loci through transethnic meta-analysis. <i>American Journal of Human Genetics</i> , 2012 , 91, 744-53	11	58
19	The association of genetic variants of type 2 diabetes with kidney function. <i>Kidney International</i> , 2012 , 82, 220-5	9.9	36
18	Association of genetic variants and incident coronary heart disease in multiethnic cohorts: the PAGE study. <i>Circulation: Cardiovascular Genetics</i> , 2011 , 4, 661-72		48
17	Combined admixture mapping and association analysis identifies a novel blood pressure genetic locus on 5p13: contributions from the CARE consortium. <i>Human Molecular Genetics</i> , 2011 , 20, 2285-95	5.6	70
16	Recent findings in the genetics of blood pressure and hypertension traits. <i>American Journal of Hypertension</i> , 2011 , 24, 392-400	2.3	35
15	Orthostatic hypotension and incident chronic kidney disease: the atherosclerosis risk in communities study. <i>Hypertension</i> , 2010 , 56, 1054-9	8.5	46
14	A whole genome association study of mother-to-child transmission of HIV in Malawi. <i>Genome Medicine</i> , 2010 , 2, 17	14.4	40
13	The association of the MYH9 gene and kidney outcomes in American Indians: the Strong Heart Family Study. <i>Human Genetics</i> , 2010 , 127, 295-301	6.3	19
12	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. <i>American Journal of Hypertension</i> , 2009 , 22, 552-8	2.3	1
11	Social- and behavioral-specific genetic effects on blood pressure traits: the Strong Heart Family Study. <i>Circulation: Cardiovascular Genetics</i> , 2009 , 2, 396-401		6
10	The effects of HIV type-1 viral suppression and non-viral factors on quantitative proteinuria in the highly active antiretroviral therapy era. <i>Antiviral Therapy</i> , 2009 , 14, 543-549	1.6	23
9	Cystatin C and preeclampsia: a case control study. <i>Renal Failure</i> , 2008 , 30, 89-95	2.9	22
8	Genome-wide linkage analysis of pulse pressure in American Indians: the Strong Heart Study. <i>American Journal of Hypertension</i> , 2008 , 21, 194-9	2.3	14
7	Diabetes-specific genetic effects on obesity traits in American Indian populations: the Strong Heart Family Study. <i>BMC Medical Genetics</i> , 2008 , 9, 90	2.1	11
6	Comparison of strategies for identification of regulatory quantitative trait loci of transcript expression traits. <i>BMC Proceedings</i> , 2007 , 1 Suppl 1, S85	2.3	0
5	Prevention and treatment strategies for glucocorticoid-induced osteoporotic fractures. <i>Clinical Rheumatology</i> , 2007 , 26, 144-53	3.9	32
4	NPHS2 gene, nephrotic syndrome and focal segmental glomerulosclerosis: a HuGE review. <i>Genetics in Medicine</i> , 2006 , 8, 63-75	8.1	71
3	A quantitative trait loci-specific gene-by-sex interaction on systolic blood pressure among American Indians: the Strong Heart Family Study. <i>Hypertension</i> , 2006 , 48, 266-70	8.5	16

2	Immunosuppression, hepatitis C infection, and acute renal failure in HIV-infected patients. <i>Journal of Acquired Immune Deficiency Syndromes (1999)</i> , 2006 , 42, 368-72	3.1	31
1	Interleukin-6 signaling effects on ischemic stroke and other cardiovascular outcomes: a Mendelian Randomization study		7