

Donna K Arnett

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8243767/publications.pdf>

Version: 2024-02-01

515
papers

61,932
citations

3325

91
h-index

1082

232
g-index

529
all docs

529
docs citations

529
times ranked

73745
citing authors

#	ARTICLE	IF	CITATIONS
1	Dietary patterns associated with stroke among West Africans: A case-control study. <i>International Journal of Stroke</i> , 2023, 18, 193-200.	2.9	8
2	Tree nut consumption and prevalence of carotid artery plaques: The National Heart, Lung, and Blood Institute Family Heart Study. <i>European Journal of Nutrition</i> , 2022, 61, 211-218.	1.8	0
3	Risk Factor Characterization of Ischemic Stroke Subtypes Among West Africans. <i>Stroke</i> , 2022, 53, 134-144.	1.0	10
4	Genetic determinants of telomere length from 109,122 ancestrally diverse whole-genome sequences in TOPMed. <i>Cell Genomics</i> , 2022, 2, 100084.	3.0	29
5	Rare coding variants in 35 genes associate with circulating lipid levels—A multi-ancestry analysis of 170,000 exomes. <i>American Journal of Human Genetics</i> , 2022, 109, 81-96.	2.6	24
6	Rare coding variants in RCN3 are associated with blood pressure. <i>BMC Genomics</i> , 2022, 23, 148.	1.2	2
7	Assessing the contribution of rare variants to complex trait heritability from whole-genome sequence data. <i>Nature Genetics</i> , 2022, 54, 263-273.	9.4	156
8	Preliminary Research on a COVID-19 Test Strategy to Guide Quarantine Interval in University Students. <i>Covid</i> , 2022, 2, 254-260.	0.7	1
9	An Amish founder population reveals rare-population genetic determinants of the human lipidome. <i>Communications Biology</i> , 2022, 5, 334.	2.0	7
10	Mendelian randomization supports bidirectional causality between telomere length and clonal hematopoiesis of indeterminate potential. <i>Science Advances</i> , 2022, 8, eabl6579.	4.7	36
11	The Value of Rare Genetic Variation in the Prediction of Common Obesity in European Ancestry Populations. <i>Frontiers in Endocrinology</i> , 2022, 13, 863893.	1.5	7
12	A multi-ethnic polygenic risk score is associated with hypertension prevalence and progression throughout adulthood. <i>Nature Communications</i> , 2022, 13, .	5.8	27
13	Insights From a Large-Scale Whole-Genome Sequencing Study of Systolic Blood Pressure, Diastolic Blood Pressure, and Hypertension. <i>Hypertension</i> , 2022, 79, 1656-1667.	1.3	12
14	Epigenome-wide association study identifies DNA methylation sites associated with target organ damage in older African Americans. <i>Epigenetics</i> , 2021, 16, 862-875.	1.3	10
15	Multi-ancestry genome-wide association study accounting for gene-psychosocial factor interactions identifies novel loci for blood pressure traits. <i>Human Genetics and Genomics Advances</i> , 2021, 2, 100013.	1.0	2
16	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.	2.7	14
17	Genome-wide association study of circulating interleukin 6 levels identifies novel loci. <i>Human Molecular Genetics</i> , 2021, 30, 393-409.	1.4	32
18	Bending the Curve in Cardiovascular Disease Mortality. <i>Circulation</i> , 2021, 143, 837-851.	1.6	35

#	ARTICLE	IF	CITATIONS
19	Whole-Exome Sequencing and hiPSC Cardiomyocyte Models Identify MYRIP, TRAPPC11, and SLC27A6 of Potential Importance to Left Ventricular Hypertrophy in an African Ancestry Population. <i>Frontiers in Genetics</i> , 2021, 12, 588452.	1.1	3
20	Sequencing of 53,831 diverse genomes from the NHLBI TOPMed Program. <i>Nature</i> , 2021, 590, 290-299.	13.7	1,069
21	Donor-specific phenotypic variation in hiPSC cardiomyocyte-derived exosomes impacts endothelial cell function. <i>American Journal of Physiology - Heart and Circulatory Physiology</i> , 2021, 320, H954-H968.	1.5	8
22	DNA Methylation and Blood Pressure Phenotypes: A Review of the Literature. <i>American Journal of Hypertension</i> , 2021, 34, 267-273.	1.0	9
23	Epigenome-wide association study of kidney function identifies trans-ethnic and ethnic-specific loci. <i>Genome Medicine</i> , 2021, 13, 74.	3.6	20
24	Discovery and fine-mapping of height loci via high-density imputation of GWASs in individuals of African ancestry. <i>American Journal of Human Genetics</i> , 2021, 108, 564-582.	2.6	18
25	Chromosome Xq23 is associated with lower atherogenic lipid concentrations and favorable cardiometabolic indices. <i>Nature Communications</i> , 2021, 12, 2182.	5.8	17
26	Age and sex are associated with the plasma lipidome: findings from the GOLDN study. <i>Lipids in Health and Disease</i> , 2021, 20, 30.	1.2	36
27	A System for Phenotype Harmonization in the National Heart, Lung, and Blood Institute Trans-Omics for Precision Medicine (TOPMed) Program. <i>American Journal of Epidemiology</i> , 2021, 190, 1977-1992.	1.6	29
28	Sugar-Sweetened Beverage Consumption and Calcified Atherosclerotic Plaques in the Coronary Arteries: The NHLBI Family Heart Study. <i>Nutrients</i> , 2021, 13, 1775.	1.7	2
29	Genome-wide association studies identify 137 genetic loci for DNA methylation biomarkers of aging. <i>Genome Biology</i> , 2021, 22, 194.	3.8	90
30	A multi-ethnic epigenome-wide association study of leukocyte DNA methylation and blood lipids. <i>Nature Communications</i> , 2021, 12, 3987.	5.8	18
31	Association of high-sensitivity C-reactive protein and odds of breast cancer by molecular subtype: analysis of the MEND study. <i>Oncotarget</i> , 2021, 12, 1230-1242.	0.8	5
32	Widespread diabetes screening for cardiovascular disease risk estimation. <i>Lancet, The</i> , 2021, 397, 2228-2230.	6.3	4
33	Association of Life-Course Educational Attainment and Breast Cancer Grade in the MEND Study. <i>Annals of Global Health</i> , 2021, 87, 59.	0.8	2
34	Identification of novel and rare variants associated with handgrip strength using whole genome sequence data from the NHLBI Trans-Omics in Precision Medicine (TOPMed) Program. <i>PLoS ONE</i> , 2021, 16, e0253611.	1.1	4
35	Adverse Cardiovascular Outcomes and Antihypertensive Treatment: A Genome-Wide Interaction Meta-Analysis in the International Consortium for Antihypertensive Pharmacogenomics Studies. <i>Clinical Pharmacology and Therapeutics</i> , 2021, 110, 723-732.	2.3	6
36	Population sequencing data reveal a compendium of mutational processes in the human germ line. <i>Science</i> , 2021, 373, 1030-1035.	6.0	43

#	ARTICLE	IF	CITATIONS
37	Influence of age on links between major modifiable risk factors and stroke occurrence in West Africa. <i>Journal of the Neurological Sciences</i> , 2021, 428, 117573.	0.3	6
38	Lipid Phenotypes and DNA Methylation: a Review of the Literature. <i>Current Atherosclerosis Reports</i> , 2021, 23, 71.	2.0	17
39	A Novel Afrocentric Stroke Risk Assessment Score: Models from the Siren Study. <i>Journal of Stroke and Cerebrovascular Diseases</i> , 2021, 30, 106003.	0.7	6
40	Association of Sickle Cell Trait With Incidence of Coronary Heart Disease Among African American Individuals. <i>JAMA Network Open</i> , 2021, 4, e2030435.	2.8	5
41	Genetic correlations between traits associated with hyperuricemia, gout, and comorbidities. <i>European Journal of Human Genetics</i> , 2021, 29, 1438-1445.	1.4	11
42	Genomics of Postprandial Lipidomics in the Genetics of Lipid-Lowering Drugs and Diet Network Study. <i>Nutrients</i> , 2021, 13, 4000.	1.7	2
43	A 6-CpG validated methylation risk score model for metabolic syndrome: The HyperGEN and GOLDN studies. <i>PLoS ONE</i> , 2021, 16, e0259836.	1.1	7
44	Metabolic Syndrome and Risk of Breast Cancer by Molecular Subtype: analysis of the MEND study. <i>Clinical Breast Cancer</i> , 2021, , .	1.1	7
45	Proximal and distal effects of genetic susceptibility to multiple sclerosis on the T cell epigenome. <i>Nature Communications</i> , 2021, 12, 7078.	5.8	15
46	The power of genetic diversity in genome-wide association studies of lipids. <i>Nature</i> , 2021, 600, 675-679.	13.7	353
47	SNPs Filtered by Allele Frequency Improve the Prediction of Hypertension Subtypes. , 2021, , .		0
48	Genetic Contributors of Incident Stroke in 10,700 African Americans With Hypertension: A Meta-Analysis From the Genetics of Hypertension Associated Treatments and Reasons for Geographic and Racial Differences in Stroke Studies. <i>Frontiers in Genetics</i> , 2021, 12, 781451.	1.1	7
49	Inherited causes of clonal haematopoiesis in 97,691 whole genomes. <i>Nature</i> , 2020, 586, 763-768.	13.7	376
50	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.	1.0	26
51	Metabolomics, Lipid Pathways, and Blood Pressure Change. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2020, 40, 1801-1803.	1.1	3
52	Carbohydrate and fat intake associated with risk of metabolic diseases through epigenetics of CPT1A. <i>American Journal of Clinical Nutrition</i> , 2020, 112, 1200-1211.	2.2	48
53	Genome-Wide Association Meta-Analysis of Individuals of European Ancestry Identifies Suggestive Loci for Sodium Intake, Potassium Intake, and Their Ratio Measured from 24-Hour or Half-Day Urine Samples. <i>Journal of Nutrition</i> , 2020, 150, 2635-2645.	1.3	4
54	Dynamic incorporation of multiple in silico functional annotations empowers rare variant association analysis of large whole-genome sequencing studies at scale. <i>Nature Genetics</i> , 2020, 52, 969-983.	9.4	146

#	ARTICLE	IF	CITATIONS
55	Loss-of-function genomic variants highlight potential therapeutic targets for cardiovascular disease. Nature Communications, 2020, 11, 6417.	5.8	39
56	Whole genome sequence association analyses of brain volumes in the TOPMed program. Alzheimer's and Dementia, 2020, 16, e040627.	0.4	0
57	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
58	Mendelian randomization analysis does not support causal associations of birth weight with hypertension risk and blood pressure in adulthood. European Journal of Epidemiology, 2020, 35, 685-697.	2.5	9
59	Smoking-by-genotype interaction in type 2 diabetes risk and fasting glucose. PLoS ONE, 2020, 15, e0230815.	1.1	10
60	A lipidome-wide association study of the lipoprotein insulin resistance index. Lipids in Health and Disease, 2020, 19, 153.	1.2	6
61	Report of the National Heart, Lung, and Blood Institute Working Group on Hypertension. Hypertension, 2020, 75, 902-917.	1.3	24
62	Salivary AMY1 Copy Number Variation Modifies Age-Related Type 2 Diabetes Risk. Clinical Chemistry, 2020, 66, 718-726.	1.5	7
63	Unraveling the risk factors for spontaneous intracerebral hemorrhage among West Africans. Neurology, 2020, 94, e998-e1012.	1.5	31
64	Smoking-by-genotype interaction in type 2 diabetes risk and fasting glucose. , 2020, 15, e0230815.		0
65	Smoking-by-genotype interaction in type 2 diabetes risk and fasting glucose. , 2020, 15, e0230815.		0
66	Smoking-by-genotype interaction in type 2 diabetes risk and fasting glucose. , 2020, 15, e0230815.		0
67	Smoking-by-genotype interaction in type 2 diabetes risk and fasting glucose. , 2020, 15, e0230815.		0
68	Genome-wide meta-analysis of SNP-by-ACEI/ARB and SNP-by-thiazide diuretic and effect on serum potassium in cohorts of European and African ancestry. Pharmacogenomics Journal, 2019, 19, 97-108.	0.9	3
69	Clinical correlates and heritability of cardiac mechanics: The HyperGEN study. International Journal of Cardiology, 2019, 274, 208-213.	0.8	5
70	Genome-wide meta-analysis of SNP and antihypertensive medication interactions on left ventricular traits in African Americans. Molecular Genetics & Genomic Medicine, 2019, 7, e00788.	0.6	4
71	2019 ACC/AHA Guideline on the Primary Prevention of Cardiovascular Disease: Part 1, Lifestyle and Behavioral Factors. JAMA Cardiology, 2019, 4, 1043.	3.0	100
72	A Clinician's Guide to Healthy Eating for Cardiovascular Disease Prevention. Mayo Clinic Proceedings Innovations, Quality & Outcomes, 2019, 3, 251-267.	1.2	72

#	ARTICLE	IF	CITATIONS
73	Gene Variants at Loci Related to Blood Pressure Account for Variation in Response to Antihypertensive Drugs Between Black and White Individuals. <i>Hypertension</i> , 2019, 74, 614-622.	1.3	14
74	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.	5.8	62
75	Association of Birth Weight With Type 2 Diabetes and Glycemic Traits. <i>JAMA Network Open</i> , 2019, 2, e1910915.	2.8	41
76	Tracing and Assessing the Evolution of Clinical Guidelines. <i>Journal of the American Heart Association</i> , 2019, 8, e014060.	1.6	3
77	Whole Genome Sequencing Identifies CRISPLD2 as a Lung Function Gene in Children With Asthma. <i>Chest</i> , 2019, 156, 1068-1079.	0.4	5
78	Comparison of smoking-related DNA methylation between newborns from prenatal exposure and adults from personal smoking. <i>Epigenomics</i> , 2019, 11, 1487-1500.	1.0	64
79	Genome-Wide Association Study of Apparent Treatment-Resistant Hypertension in the CHARGE Consortium: The CHARGE Pharmacogenetics Working Group. <i>American Journal of Hypertension</i> , 2019, 32, 1146-1153.	1.0	17
80	Genetic influences on susceptibility to rheumatoid arthritis in African-Americans. <i>Human Molecular Genetics</i> , 2019, 28, 858-874.	1.4	55
81	Multi-ancestry study of blood lipid levels identifies four loci interacting with physical activity. <i>Nature Communications</i> , 2019, 10, 376.	5.8	64
82	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.	1.8	29
83	A PheWAS study of a large observational epidemiological cohort of African Americans from the REGARDS study. <i>BMC Medical Genomics</i> , 2019, 12, 26.	0.7	9
84	Association of dietary folate and vitamin B-12 intake with genome-wide DNA methylation in blood: a large-scale epigenome-wide association analysis in 5841 individuals. <i>American Journal of Clinical Nutrition</i> , 2019, 110, 437-450.	2.2	46
85	Echocardiographic Abnormalities and Determinants of 1â€Month Outcome of Stroke Among West Africans in the SIREN Study. <i>Journal of the American Heart Association</i> , 2019, 8, e010814.	1.6	3
86	Collaborative Molecular Epidemiology Study of Metabolic Dysregulation, DNA Methylation, and Breast Cancer Risk Among Nigerian Women: MEND Study Objectives and Design. <i>Journal of Global Oncology</i> , 2019, 5, 1-9.	0.5	6
87	An integrative cross-omics analysis of DNA methylation sites of glucose and insulin homeostasis. <i>Nature Communications</i> , 2019, 10, 2581.	5.8	62
88	2019 ACC/AHA Guideline on the Primary Prevention of Cardiovascular Disease: A Report of the American College of Cardiology/American Heart Association Task Force on Clinical Practice Guidelines. <i>Circulation</i> , 2019, 140, e596-e646.	1.6	1,789
89	Differential Impact of Risk Factors on Stroke Occurrence Among Men Versus Women in West Africa. <i>Stroke</i> , 2019, 50, 820-827.	1.0	26
90	An Exome-Wide Sequencing Study of the GOLDN Cohort Reveals Novel Associations of Coding Variants and Fasting Plasma Lipids. <i>Frontiers in Genetics</i> , 2019, 10, 158.	1.1	2

#	ARTICLE	IF	CITATIONS
91	2019 ACC/AHA Guideline on the Primary Prevention of Cardiovascular Disease: Executive Summary. <i>Journal of the American College of Cardiology</i> , 2019, 74, 1376-1414.	1.2	820
92	2019 ACC/AHA Guideline on the Primary Prevention of Cardiovascular Disease: Executive Summary: A Report of the American College of Cardiology/American Heart Association Task Force on Clinical Practice Guidelines. <i>Circulation</i> , 2019, 140, e563-e595.	1.6	1,676
93	A multi-ancestry genome-wide study incorporating gene-smoking interactions identifies multiple new loci for pulse pressure and mean arterial pressure. <i>Human Molecular Genetics</i> , 2019, 28, 2615-2633.	1.4	31
94	Associations between SLC16A11 variants and diabetes in the Hispanic Community Health Study/Study of Latinos (HCHS/SOL). <i>Scientific Reports</i> , 2019, 9, 843.	1.6	9
95	Systematic Error Removal Using Random Forest for Normalizing Large-Scale Untargeted Lipidomics Data. <i>Analytical Chemistry</i> , 2019, 91, 3590-3596.	3.2	163
96	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. <i>PLoS Genetics</i> , 2019, 15, e1008500.	1.5	203
97	Integrating hypertension phenotype and genotype with hybrid non-negative matrix factorization. <i>Bioinformatics</i> , 2019, 35, 1395-1403.	1.8	12
98	Associations of Mitochondrial and Nuclear Mitochondrial Variants and Genes with Seven Metabolic Traits. <i>American Journal of Human Genetics</i> , 2019, 104, 112-138.	2.6	106
99	The Interaction of a Diabetes Gene Risk Score With 3 Different Antihypertensive Medications for Incident Glucose-level Elevation. <i>American Journal of Hypertension</i> , 2019, 32, 343-349.	1.0	0
100	Combined linkage and association analysis identifies rare and low frequency variants for blood pressure at 1q31. <i>European Journal of Human Genetics</i> , 2019, 27, 269-277.	1.4	5
101	An exome-wide sequencing study of lipid response to high-fat meal and fenofibrate in Caucasians from the GOLDN cohort. <i>Journal of Lipid Research</i> , 2018, 59, 722-729.	2.0	10
102	Dominant modifiable risk factors for stroke in Ghana and Nigeria (SIREN): a case-control study. <i>The Lancet Global Health</i> , 2018, 6, e436-e446.	2.9	183
103	Association of Sickle Cell Trait With Ischemic Stroke Among African Americans. <i>JAMA Neurology</i> , 2018, 75, 802.	4.5	25
104	Association of Methylation Signals With Incident Coronary Heart Disease in an Epigenome-Wide Assessment of Circulating Tumor Necrosis Factor \pm . <i>JAMA Cardiology</i> , 2018, 3, 463.	3.0	33
105	Epigenome-wide association study of metabolic syndrome in African-American adults. <i>Clinical Epigenetics</i> , 2018, 10, 49.	1.8	49
106	Genome-Wide Interactions with Dairy Intake for Body Mass Index in Adults of European Descent. <i>Molecular Nutrition and Food Research</i> , 2018, 62, 1700347.	1.5	9
107	Dairy Consumption and Body Mass Index Among Adults: Mendelian Randomization Analysis of 184802 Individuals from 25 Studies. <i>Clinical Chemistry</i> , 2018, 64, 183-191.	1.5	34
108	Hypermethylation of <i>MIR21</i> in CD4+ T cells from patients with relapsing-remitting multiple sclerosis associates with lower miRNA-21 levels and concomitant up-regulation of its target genes. <i>Multiple Sclerosis Journal</i> , 2018, 24, 1288-1300.	1.4	33

#	ARTICLE	IF	CITATIONS
109	Data for GAW20: genome-wide DNA sequence variation and epigenome-wide DNA methylation before and after fenofibrate treatment in a family study of metabolic phenotypes. BMC Proceedings, 2018, 12, 35.	1.8	11
110	Genome-wide association meta-analysis of circulating odd-numbered chain saturated fatty acids: Results from the CHARGE Consortium. PLoS ONE, 2018, 13, e0196951.	1.1	14
111	Genome-wide association study of response to methotrexate in early rheumatoid arthritis patients. Pharmacogenomics Journal, 2018, 18, 528-538.	0.9	42
112	Metabolic and inflammatory biomarkers are associated with epigenetic aging acceleration estimates in the GOLDN study. Clinical Epigenetics, 2018, 10, 56.	1.8	68
113	Omics of Blood Pressure and Hypertension. Circulation Research, 2018, 122, 1409-1419.	2.0	74
114	Epigenomics and metabolomics reveal the mechanism of the APOA2-saturated fat intake interaction affecting obesity. American Journal of Clinical Nutrition, 2018, 108, 188-200.	2.2	54
115	Epigenetic Patterns in Blood Associated With Lipid Traits Predict Incident Coronary Heart Disease Events and Are Enriched for Results From Genome-Wide Association Studies. Circulation: Cardiovascular Genetics, 2017, 10, .	5.1	104
116	Sickle Cell Trait and the Risk of ESRD in Blacks. Journal of the American Society of Nephrology: JASN, 2017, 28, 2180-2187.	3.0	79
117	Phenomapping for the Identification of Hypertensive Patients with the Myocardial Substrate for Heart Failure with Preserved Ejection Fraction. Journal of Cardiovascular Translational Research, 2017, 10, 275-284.	1.1	61
118	Interleukin-6 (IL-6) rs1800796 and cyclin dependent kinase inhibitor (CDKN2A/CDKN2B) rs2383207 are associated with ischemic stroke in indigenous West African Men. Journal of the Neurological Sciences, 2017, 379, 229-235.	0.3	31
119	Genome- and CD4 + T-cell methylome-wide association study of circulating trimethylamine-N-oxide in the Genetics of Lipid Lowering Drugs and Diet Network (GOLDN). Journal of Nutrition & Intermediary Metabolism, 2017, 8, 1-7.	1.7	11
120	Non-linear patterns in age-related DNA methylation may reflect CD4 ⁺ T cell differentiation. Epigenetics, 2017, 12, 492-503.	1.3	24
121	Stroke in Indigenous Africans, African Americans, and European Americans. Stroke, 2017, 48, 1169-1175.	1.0	44
122	Genetic associations with lipoprotein subfraction measures differ by ethnicity in the multi-ethnic study of atherosclerosis (MESA). Human Genetics, 2017, 136, 715-726.	1.8	12
123	Discovery and fine-mapping of loci associated with MUFAs through trans-ethnic meta-analysis in Chinese and European populations. Journal of Lipid Research, 2017, 58, 974-981.	2.0	18
124	Coffee consumption and calcified atherosclerotic plaques in the coronary arteries: The NHLBI Family Heart Study. Clinical Nutrition ESPEN, 2017, 17, 18-21.	0.5	10
125	Genetic variation at 16q24.2 is associated with small vessel stroke. Annals of Neurology, 2017, 81, 383-394.	2.8	73
126	Detection of gene-environment interactions in a family-based population using SCAD. Statistics in Medicine, 2017, 36, 3547-3559.	0.8	4

#	ARTICLE	IF	CITATIONS
127	Advancing stroke genomic research in the age of Trans-Omics big data science: Emerging priorities and opportunities. <i>Journal of the Neurological Sciences</i> , 2017, 382, 18-28.	0.3	15
128	Testing Two Evolutionary Theories of Human Aging with DNA Methylation Data. <i>Genetics</i> , 2017, 207, 1547-1560.	1.2	12
129	An epigenome-wide association study of inflammatory response to fenofibrate in the Genetics of Lipid Lowering Drugs and Diet Network. <i>Pharmacogenomics</i> , 2017, 18, 1333-1341.	0.6	16
130	Association of Estimated Sodium Intake With Adverse Cardiac Structure and Function. <i>Journal of the American College of Cardiology</i> , 2017, 70, 715-724.	1.2	21
131	DNA Methylation Analysis Identifies Loci for Blood Pressure Regulation. <i>American Journal of Human Genetics</i> , 2017, 101, 888-902.	2.6	154
132	Sex Differences in Blood HDL, the Total Cholesterol/HDL Ratio, and Palmitoleic Acid are Not Associated with Variants in Common Candidate Genes. <i>Lipids</i> , 2017, 52, 969-980.	0.7	19
133	Association of Body Mass Index with DNA Methylation and Gene Expression in Blood Cells and Relations to Cardiometabolic Disease: A Mendelian Randomization Approach. <i>PLoS Medicine</i> , 2017, 14, e1002215.	3.9	246
134	Dense Genotyping of Immune-Related Regions Identifies Loci for Rheumatoid Arthritis Risk and Damage in African Americans. <i>Molecular Medicine</i> , 2017, 23, 177-187.	1.9	18
135	Exploring Overlaps Between the Genomic and Environmental Determinants of LVH and Stroke: A Multicenter Study in West Africa. <i>Global Heart</i> , 2017, 12, 107.	0.9	10
136	Prevalence and Prognostic Features of ECG Abnormalities in Acute Stroke: Findings From the SIREN Study Among Africans. <i>Global Heart</i> , 2017, 12, 99.	0.9	26
137	Large-scale genome-wide analysis identifies genetic variants associated with cardiac structure and function. <i>Journal of Clinical Investigation</i> , 2017, 127, 1798-1812.	3.9	106
138	Discovery and fine-mapping of adiposity loci using high density imputation of genome-wide association studies in individuals of African ancestry: African Ancestry Anthropometry Genetics Consortium. <i>PLoS Genetics</i> , 2017, 13, e1006719.	1.5	98
139	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.	1.5	88
140	Behavior related genes, dietary preferences and anthropometric traits. <i>FASEB Journal</i> , 2017, 31, .	0.2	1
141	Whole Exome Analyses to Examine the Impact of Rare Variants on Left Ventricular Traits in African American Participants from the HyperGEN and GENOA Studies. <i>Journal of Hypertension and Management</i> , 2017, 3, .	0.1	0
142	An Empirical Comparison of Joint and Stratified Frameworks for Studying G × E Interactions: Systolic Blood Pressure and Smoking in the CHARGE Gene-Lifestyle Interactions Working Group. <i>Genetic Epidemiology</i> , 2016, 40, 404-415.	0.6	18
143	A genome-wide study of lipid response to fenofibrate in Caucasians. <i>Pharmacogenetics and Genomics</i> , 2016, 26, 324-333.	0.7	12
144	Higher chylomicron remnants and LDL particle numbers associate with CD36 SNPs and DNA methylation sites that reduce CD36. <i>Journal of Lipid Research</i> , 2016, 57, 2176-2184.	2.0	26

#	ARTICLE	IF	CITATIONS
145	DNA methylation signatures of chronic low-grade inflammation are associated with complex diseases. <i>Genome Biology</i> , 2016, 17, 255.	3.8	251
146	A Genome-wide study of blood pressure in African Americans accounting for gene-smoking interaction. <i>Scientific Reports</i> , 2016, 6, 18812.	1.6	34
147	The US Cancer Moonshot initiative. <i>Lancet Oncology</i> , The, 2016, 17, e178-e180.	5.1	15
148	The Heart of 25 by 25: Achieving the Goal of Reducing Global and Regional Premature Deaths From Cardiovascular Diseases and Stroke. <i>Circulation</i> , 2016, 133, e674-90.	1.6	155
149	Walking and Calcified Atherosclerotic Plaque in the Coronary Arteries. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2016, 36, 1272-1277.	1.1	12
150	Identification of additional risk loci for stroke and small vessel disease: a meta-analysis of genome-wide association studies. <i>Lancet Neurology</i> , The, 2016, 15, 695-707.	4.9	130
151	The Role of Healthy Lifestyle in the Primordial Prevention of Cardiovascular Disease. <i>Current Cardiology Reports</i> , 2016, 18, 56.	1.3	61
152	Evidence-Based Policy Making: Assessment of the American Heart Association's Strategic Policy Portfolio. <i>Circulation</i> , 2016, 133, e615-53.	1.6	36
153	Neurogenomics in Africa: Perspectives, progress, possibilities and priorities. <i>Journal of the Neurological Sciences</i> , 2016, 366, 213-223.	0.3	30
154	Assessment of postprandial triglycerides in clinical practice: Validation in a general population and coronary heart disease patients. <i>Journal of Clinical Lipidology</i> , 2016, 10, 1163-1171.	0.6	22
155	Editorial: A Novel Genetic Association With Systemic Sclerosis: The Utility of Whole-Exome Sequencing in Autoimmune Disease. <i>Arthritis and Rheumatology</i> , 2016, 68, 27-30.	2.9	0
156	Mediterranean diet score and left ventricular structure and function: the Multi-Ethnic Study of Atherosclerosis. <i>American Journal of Clinical Nutrition</i> , 2016, 104, 595-602.	2.2	22
157	Epigenetics of Lipid Phenotypes. <i>Current Cardiovascular Risk Reports</i> , 2016, 10, 1.	0.8	20
158	Research Needs to Improve Hypertension Treatment and Control in African Americans. <i>Hypertension</i> , 2016, 68, 1066-1072.	1.3	78
159	Epigenetic Signatures of Cigarette Smoking. <i>Circulation: Cardiovascular Genetics</i> , 2016, 9, 436-447.	5.1	678
160	Interaction of an S100A9 gene variant with saturated fat and carbohydrates to modulate insulin resistance in 3 populations of different ancestries. <i>American Journal of Clinical Nutrition</i> , 2016, 104, 508-517.	2.2	11
161	Epigenome-wide association study of triglyceride postprandial responses to a high-fat dietary challenge. <i>Journal of Lipid Research</i> , 2016, 57, 2200-2207.	2.0	40
162	Associations of the MCM6-rs3754686 proxy for milk intake in Mediterranean and American populations with cardiovascular biomarkers, disease and mortality: Mendelian randomization. <i>Scientific Reports</i> , 2016, 6, 33188.	1.6	18

#	ARTICLE	IF	CITATIONS
163	Association of Central Adiposity With Adverse Cardiac Mechanics. <i>Circulation: Cardiovascular Imaging</i> , 2016, 9, .	1.3	65
164	Meta-analysis of genome-wide association studies discovers multiple loci for chronic lymphocytic leukemia. <i>Nature Communications</i> , 2016, 7, 10933.	5.8	94
165	Precision Medicine, Genomics, and Public Health. <i>Diabetes Care</i> , 2016, 39, 1870-1873.	4.3	16
166	The effects of omega-3 polyunsaturated fatty acids and genetic variants on methylation levels of the interleukin-6 gene promoter. <i>Molecular Nutrition and Food Research</i> , 2016, 60, 410-419.	1.5	41
167	Archeological Echocardiography: Digitization and Speckle Tracking Analysis of Archival Echocardiograms in the HyperGEN Study. <i>Echocardiography</i> , 2016, 33, 386-397.	0.3	24
168	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.	2.6	55
169	Clock Genes Explain a Large Proportion of Phenotypic Variance in Systolic Blood Pressure and This Control Is Not Modified by Environmental Temperature. <i>American Journal of Hypertension</i> , 2016, 29, 132-140.	1.0	20
170	Heart Disease and Stroke Statistics—2016 Update. <i>Circulation</i> , 2016, 133, e38-360.	1.6	5,447
171	Opportunities for the Cardiovascular Community in the Precision Medicine Initiative. <i>Circulation</i> , 2016, 133, 226-231.	1.6	50
172	Using genomic approaches to identify CVD-causing variants. <i>Nature Reviews Cardiology</i> , 2016, 13, 72-74.	6.1	4
173	Interaction of methylation-related genetic variants with circulating fatty acids on plasma lipids: a meta-analysis of 7 studies and methylation analysis of 3 studies in the Cohorts for Heart and Aging Research in Genomic Epidemiology consortium. <i>American Journal of Clinical Nutrition</i> , 2016, 103, 567-578.	2.2	24
174	Loci associated with ischaemic stroke and its subtypes (SiGN): a genome-wide association study. <i>Lancet Neurology</i> , The, 2016, 15, 174-184.	4.9	217
175	Executive Summary: Heart Disease and Stroke Statistics—2016 Update. <i>Circulation</i> , 2016, 133, 447-454.	1.6	2,093
176	Multilingual Validation of the Questionnaire for Verifying Stroke-Free Status in West Africa. <i>Stroke</i> , 2016, 47, 167-172.	1.0	45
177	Association of Sickle Cell Trait with Risk of Coronary Heart Disease in African Americans. <i>Blood</i> , 2016, 128, 11-11.	0.6	3
178	Association of DNA Methylation at CPT1A Locus with Metabolic Syndrome in the Genetics of Lipid Lowering Drugs and Diet Network (GOLDN) Study. <i>PLoS ONE</i> , 2016, 11, e0145789.	1.1	54
179	Heritable DNA Methylation in CD4+ Cells among Complex Families Displays Genetic and Non-Genetic Effects. <i>PLoS ONE</i> , 2016, 11, e0165488.	1.1	19
180	Genetic variants modify the effect of age on APOE methylation in the genetics of lipid lowering drugs and diet network study. <i>Aging Cell</i> , 2015, 14, 49-59.	3.0	34

#	ARTICLE	IF	CITATIONS
181	Comprehensive evaluation of AmpliSeq transcriptome, a novel targeted whole transcriptome RNA sequencing methodology for global gene expression analysis. <i>BMC Genomics</i> , 2015, 16, 1069.	1.2	73
182	Lack of association of apolipoprotein E (Apo E) polymorphism with the prevalence of metabolic syndrome: the National Heart, Lung and Blood Institute Family Heart Study. <i>Diabetes/Metabolism Research and Reviews</i> , 2015, 31, 582-587.	1.7	5
183	Epigenome-wide study identifies novel methylation loci associated with body mass index and waist circumference. <i>Obesity</i> , 2015, 23, 1493-1501.	1.5	152
184	Lipid changes due to fenofibrate treatment are not associated with changes in DNA methylation patterns in the GOLDN study. <i>Frontiers in Genetics</i> , 2015, 6, 304.	1.1	20
185	The burden of stroke in Africa: a glance at the present and a glimpse into the future: review article. <i>Cardiovascular Journal of Africa</i> , 2015, 26, S27-S38.	0.2	286
186	Drug-Gene Interactions of Antihypertensive Medications and Risk of Incident Cardiovascular Disease: A Pharmacogenomics Study from the CHARGE Consortium. <i>PLoS ONE</i> , 2015, 10, e0140496.	1.1	15
187	PCSK9 variation and association with blood pressure in African Americans: preliminary findings from the HyperGEN and REGARDS studies. <i>Frontiers in Genetics</i> , 2015, 6, 136.	1.1	25
188	The Role of Rare Variants in Systolic Blood Pressure: Analysis of ExomeChip Data in HyperGEN African Americans. <i>Human Heredity</i> , 2015, 79, 20-27.	0.4	13
189	Phenotyping Stroke in Sub-Saharan Africa: Stroke Investigative Research and Education Network (SIREN) Phenomics Protocol. <i>Neuroepidemiology</i> , 2015, 45, 73-82.	1.1	73
190	Healthy Habits, Healthy Women —. <i>Journal of the American College of Cardiology</i> , 2015, 65, 52-54.	1.2	2
191	Genetic loci associated with circulating phospholipid trans fatty acids: a meta-analysis of genome-wide association studies from the CHARGE Consortium. <i>American Journal of Clinical Nutrition</i> , 2015, 101, 398-406.	2.2	49
192	Executive Summary: Heart Disease and Stroke Statistics—2015 Update. <i>Circulation</i> , 2015, 131, 434-441.	1.6	509
193	Dietary fatty acids modulate associations between genetic variants and circulating fatty acids in plasma and erythrocyte membranes: Meta-analysis of nine studies in the CHARGE consortium. <i>Molecular Nutrition and Food Research</i> , 2015, 59, 1373-1383.	1.5	37
194	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.	2.6	321
195	Genetic loci associated with circulating levels of very long-chain saturated fatty acids. <i>Journal of Lipid Research</i> , 2015, 56, 176-184.	2.0	38
196	Heart Disease and Stroke Statistics—2015 Update. <i>Circulation</i> , 2015, 131, e29-322.	1.6	5,963
197	Deep sequencing of RYR3 gene identifies rare and common variants associated with increased carotid intima-media thickness (cIMT) in HIV-infected individuals. <i>Journal of Human Genetics</i> , 2015, 60, 63-67.	1.1	3
198	Directional dominance on stature and cognition in diverse human populations. <i>Nature</i> , 2015, 523, 459-462.	13.7	173

#	ARTICLE	IF	CITATIONS
199	Genome-wide association study of triglyceride response to a high-fat meal among participants of the NHLBI Genetics of Lipid Lowering Drugs and Diet Network (GOLDN). <i>Metabolism: Clinical and Experimental</i> , 2015, 64, 1359-1371.	1.5	33
200	Gene-Environment Interactions of Circadian-Related Genes for Cardiometabolic Traits. <i>Diabetes Care</i> , 2015, 38, 1456-1466.	4.3	52
201	Future Translational Applications From the Contemporary Genomics Era. <i>Circulation</i> , 2015, 131, 1715-1736.	1.6	38
202	Variants for HDL-C, LDL-C, and Triglycerides Identified from Admixture Mapping and Fine-Mapping Analysis in African American Families. <i>Circulation: Cardiovascular Genetics</i> , 2015, 8, 106-113.	5.1	18
203	Epigenome-wide association study (EWAS) of BMI, BMI change and waist circumference in African American adults identifies multiple replicated loci. <i>Human Molecular Genetics</i> , 2015, 24, 4464-4479.	1.4	289
204	Plugging the Leaking Pipeline. <i>Circulation: Cardiovascular Quality and Outcomes</i> , 2015, 8, S63-4.	0.9	9
205	Consumption of meat is associated with higher fasting glucose and insulin concentrations regardless of glucose and insulin genetic risk scores: a meta-analysis of 50,345 Caucasians. <i>American Journal of Clinical Nutrition</i> , 2015, 102, 1266-1278.	2.2	69
206	Analysis of Large Electronic Health Record Databases Supports Blood Pressure-Independent Diabetes Association. <i>Journal of the American College of Cardiology</i> , 2015, 66, 1563-1565.	1.2	1
207	Clinical applications of epigenetics in cardiovascular disease: the long road ahead. <i>Translational Research</i> , 2015, 165, 143-153.	2.2	23
208	Association of Hemoglobin S and C Traits with Kidney Disease in African Americans in the Reasons for Geographic and Racial Differences in Stroke (REGARDS) Study. <i>Blood</i> , 2015, 126, 70-70.	0.6	0
209	Genetic Risk Scores Associated with Baseline Lipoprotein Subfraction Concentrations Do Not Associate with Their Responses to Fenofibrate. <i>Biology</i> , 2014, 3, 536-550.	1.3	1
210	Sex-Specific Associations between Screen Time and Lipoprotein Subfractions. <i>International Journal of Sport Nutrition and Exercise Metabolism</i> , 2014, 24, 59-69.	1.0	2
211	Methylation at CPT1A locus is associated with lipoprotein subfraction profiles. <i>Journal of Lipid Research</i> , 2014, 55, 1324-1330.	2.0	65
212	The effects of angiotensinogen gene polymorphisms on cardiovascular disease outcomes during antihypertensive treatment in the GenHAT study. <i>Frontiers in Pharmacology</i> , 2014, 5, 210.	1.6	14
213	<i>CRY1</i> circadian gene variant interacts with carbohydrate intake for insulin resistance in two independent populations: Mediterranean and North American. <i>Chronobiology International</i> , 2014, 31, 660-667.	0.9	56
214	AHA/ACC/HHS Strategies to Enhance Application of Clinical Practice Guidelines in Patients With Cardiovascular Disease and Comorbid Conditions. <i>Circulation</i> , 2014, 130, 1662-1667.	1.6	132
215	Hemodynamic and Echocardiographic Profiles in African American Compared With White Offspring of Hypertensive Parents: The HyperGEN Study. <i>American Journal of Hypertension</i> , 2014, 27, 21-26.	1.0	7
216	Epigenome-Wide Association Study of Fasting Measures of Glucose, Insulin, and HOMA-IR in the Genetics of Lipid Lowering Drugs and Diet Network Study. <i>Diabetes</i> , 2014, 63, 801-807.	0.3	149

#	ARTICLE	IF	CITATIONS
217	An Empirical Comparison of Meta-analysis and Mega-analysis of Individual Participant Data for Identifying Gene-Environment Interactions. <i>Genetic Epidemiology</i> , 2014, 38, 369-378.	0.6	48
218	Association of Comorbidity Burden With Abnormal Cardiac Mechanics: Findings From the HyperGEN Study. <i>Journal of the American Heart Association</i> , 2014, 3, e000631.	1.6	19
219	<i>PRKCZ</i> methylation is associated with sunlight exposure in a North American but not a Mediterranean population. <i>Chronobiology International</i> , 2014, 31, 1034-1040.	0.9	12
220	The Role of Worksite Health Screening. <i>Circulation</i> , 2014, 130, 719-734.	1.6	31
221	CPT1A: the future of heart disease detection and personalized medicine?. <i>Clinical Lipidology</i> , 2014, 9, 9-12.	0.4	8
222	Evolution of "The Guideline Advantage". <i>Circulation: Cardiovascular Quality and Outcomes</i> , 2014, 7, 493-498.	0.9	14
223	Genome-wide association studies identified novel loci for non-high-density lipoprotein cholesterol and its postprandial lipemic response. <i>Human Genetics</i> , 2014, 133, 919-930.	1.8	10
224	HIV, Inflammation, and Calcium in Atherosclerosis. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2014, 34, 244-250.	1.1	51
225	Association of Low-Grade Albuminuria With Adverse Cardiac Mechanics. <i>Circulation</i> , 2014, 129, 42-50.	1.6	70
226	Epigenome-Wide Association Study of Fasting Blood Lipids in the Genetics of Lipid-Lowering Drugs and Diet Network Study. <i>Circulation</i> , 2014, 130, 565-572.	1.6	190
227	Adult Height and Prevalence of Coronary Artery Calcium. <i>Circulation: Cardiovascular Imaging</i> , 2014, 7, 52-57.	1.3	9
228	Association of egg consumption and calcified atherosclerotic plaque in the coronary arteries: The NHLBI Family Heart Study. <i>E-SPEN Journal</i> , 2014, 9, e131-e135.	0.5	7
229	Saturated Fat Intake Modulates the Association between an Obesity Genetic Risk Score and Body Mass Index in Two US Populations. <i>Journal of the Academy of Nutrition and Dietetics</i> , 2014, 114, 1954-1966.	0.4	60
230	The 50th Anniversary of the US Surgeon General's Report on Tobacco: What We've Accomplished and Where We Go From Here. <i>Journal of the American Heart Association</i> , 2014, 3, e000740.	1.6	19
231	Genomics of Post-Prandial Lipidomic Phenotypes in the Genetics of Lipid Lowering Drugs and Diet Network (GOLDN) Study. <i>PLoS ONE</i> , 2014, 9, e99509.	1.1	21
232	RNA Expression Profiling of Human iPSC-Derived Cardiomyocytes in a Cardiac Hypertrophy Model. <i>PLoS ONE</i> , 2014, 9, e108051.	1.1	51
233	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, 545-554.	2.6	189
234	Transforming Cardiovascular Health Through Genes and Environment. <i>Circulation</i> , 2013, 127, 2066-2070.	1.6	6

#	ARTICLE	IF	CITATIONS
235	Genome-Wide Association Study of Cardiac Structure and Systolic Function in African Americans. <i>Circulation: Cardiovascular Genetics</i> , 2013, 6, 37-46.	5.1	46
236	Genetics and Genomics for the Prevention and Treatment of Cardiovascular Disease: Update. <i>Circulation</i> , 2013, 128, 2813-2851.	1.6	100
237	Gain-of-Function Lipoprotein Lipase Variant rs13702 Modulates Lipid Traits through Disruption of a MicroRNA-410 Seed Site. <i>American Journal of Human Genetics</i> , 2013, 92, 5-14.	2.6	67
238	Association of dietary omega-3 fatty acids with prevalence of metabolic syndrome: The National Heart, Lung, and Blood Institute Family Heart Study. <i>Clinical Nutrition</i> , 2013, 32, 966-969.	2.3	39
239	Genetic Analysis of 16 NMR Lipoprotein Fractions in Humans, the GOLDN Study. <i>Lipids</i> , 2013, 48, 155-165.	0.7	34
240	Associations between NOS1AP Single Nucleotide Polymorphisms (SNPs) and QT Interval Duration in Four Racial/Ethnic Groups in the Multi-Ethnic Study of Atherosclerosis (MESA). <i>Annals of Noninvasive Electrocardiology</i> , 2013, 18, 29-40.	0.5	10
241	Pharmacogenomics of high-density lipoprotein-cholesterol-raising therapies. <i>Expert Review of Cardiovascular Therapy</i> , 2013, 11, 355-364.	0.6	8
242	Modulation by Dietary Fat and Carbohydrate of <i>IRS1</i> Association With Type 2 Diabetes Traits in Two Populations of Different Ancestries. <i>Diabetes Care</i> , 2013, 36, 2621-2627.	4.3	25
243	A meta-analysis identifies new loci associated with body mass index in individuals of African ancestry. <i>Nature Genetics</i> , 2013, 45, 690-696.	9.4	232
244	Genetic variants associated with VLDL, LDL and HDL particle size differ with race/ethnicity. <i>Human Genetics</i> , 2013, 132, 405-413.	1.8	30
245	Stroke Genetics Network (SiGN) Study. <i>Stroke</i> , 2013, 44, 2694-2702.	1.0	62
246	SORCS1 contributes to the development of renal disease in rats and humans. <i>Physiological Genomics</i> , 2013, 45, 720-728.	1.0	17
247	Genome-wide meta-analysis of observational studies shows common genetic variants associated with macronutrient intake. <i>American Journal of Clinical Nutrition</i> , 2013, 97, 1395-1402.	2.2	210
248	Trans-Ethnic Fine-Mapping of Lipid Loci Identifies Population-Specific Signals and Allelic Heterogeneity That Increases the Trait Variance Explained. <i>PLoS Genetics</i> , 2013, 9, e1003379.	1.5	112
249	ACCF/AHA Clinical Practice Guideline Methodology Summit Report. <i>Circulation</i> , 2013, 127, 268-310.	1.6	101
250	Genetic Variants at PSMD3 Interact with Dietary Fat and Carbohydrate to Modulate Insulin Resistance. <i>Journal of Nutrition</i> , 2013, 143, 354-361.	1.3	17
251	Apolipoprotein A2 Polymorphism Interacts with Intakes of Dairy Foods to Influence Body Weight in 2 U.S. Populations. <i>Journal of Nutrition</i> , 2013, 143, 1865-1871.	1.3	27
252	To Replicate or Not to Replicate: The Case of Pharmacogenetic Studies. <i>Circulation: Cardiovascular Genetics</i> , 2013, 6, 409-412.	5.1	25

#	ARTICLE	IF	CITATIONS
253	SNPs located at CpG sites modulate genome-epigenome interaction. <i>Epigenetics</i> , 2013, 8, 802-806.	1.3	131
254	Polyunsaturated Fatty Acids Modulate the Association between PIK3CA-KCNMB3 Genetic Variants and Insulin Resistance. <i>PLoS ONE</i> , 2013, 8, e67394.	1.1	10
255	The role of SNP-loop diuretic interactions in hypertension across ethnic groups in HyperGEN. <i>Frontiers in Genetics</i> , 2013, 4, 304.	1.1	11
256	Genomic Copy Number Variants: Evidence for Association with Antibody Response to Anthrax Vaccine Adsorbed. <i>PLoS ONE</i> , 2013, 8, e64813.	1.1	8
257	Genome-Wide Contribution of Genotype by Environment Interaction to Variation of Diabetes-Related Traits. <i>PLoS ONE</i> , 2013, 8, e77442.	1.1	41
258	Sociedade Brasileira de Cardiologia: carta do Rio de Janeiro - III Brasil Prevent / I Am�rica Latina Prevent. <i>Arquivos Brasileiros De Cardiologia</i> , 2013, 100, 3-5.	0.3	7
259	Genome-wide contribution of genotype by environment interaction to blood lipid variation. <i>FASEB Journal</i> , 2013, 27, 222.4.	0.2	0
260	Preventing and Controlling Hypertension in the Era of Genomic Innovation and Environmental Transformation. <i>JAMA - Journal of the American Medical Association</i> , 2012, 308, 1745.	3.8	5
261	Divided States of America: Regional Variation in Cardiovascular Health. <i>Journal of the American Heart Association</i> , 2012, 1, e006114.	1.6	3
262	Genome-wide meta-analysis points to CTC1 and ZNF676 as genes regulating telomere homeostasis in humans. <i>Human Molecular Genetics</i> , 2012, 21, 5385-5394.	1.4	210
263	Association Between Family Risk of Stroke and Myocardial Infarction With Prevalent Risk Factors and Coexisting Diseases. <i>Stroke</i> , 2012, 43, 974-979.	1.0	11
264	Obesity-insulin targeted genes in the 3p26-25 region in human studies and LG/J and SM/J mice. <i>Metabolism: Clinical and Experimental</i> , 2012, 61, 1129-1141.	1.5	9
265	Chocolate consumption and prevalence of metabolic syndrome in the NHLBI Family Heart Study. <i>E-SPEN Journal</i> , 2012, 7, e139-e143.	0.5	5
266	Renin-Angiotensin Inhibition in Systolic Heart Failure and Chronic Kidney Disease. <i>American Journal of Medicine</i> , 2012, 125, 399-410.	0.6	69
267	Normal Limits in Relation to Age, Body Size and Gender of Two-Dimensional Echocardiographic Aortic Root Dimensions in Persons �15 Years of Age. <i>American Journal of Cardiology</i> , 2012, 110, 1189-1194.	0.7	303
268	Preliminary Evidence for an Association between LRP-1 Genotype and Body Mass Index in Humans. <i>PLoS ONE</i> , 2012, 7, e30732.	1.1	8
269	Pharmacogenetic Association of NOS3 Variants with Cardiovascular Disease in Patients with Hypertension: The GenHAT Study. <i>PLoS ONE</i> , 2012, 7, e34217.	1.1	36
270	Clustering by Plasma Lipoprotein Profile Reveals Two Distinct Subgroups with Positive Lipid Response to Fenofibrate Therapy. <i>PLoS ONE</i> , 2012, 7, e38072.	1.1	30

#	ARTICLE	IF	CITATIONS
271	Variants Identified in a GWAS Meta-Analysis for Blood Lipids Are Associated with the Lipid Response to Fenofibrate. PLoS ONE, 2012, 7, e48663.	1.1	39
272	Wicked Problems and Worthy Pursuits. Circulation, 2012, 125, 2554-2556.	1.6	6
273	Genotype Imputation for African Americans Using Data From Hispanic Multi-Ethnic Phase III Versus 1000 Genomes Projects. Genetic Epidemiology, 2012, 36, 508-516.	0.6	13
274	Subclinical, hemodynamic, and echocardiographic abnormalities of high pulse pressure in hypertensive and non-hypertensive adults. American Journal of Cardiovascular Disease, 2012, 2, 309-17.	0.5	5
275	The Relationship Between Visit-to-Visit Variability in Systolic Blood Pressure and All-Cause Mortality in the General Population. Hypertension, 2011, 57, 160-166.	1.3	397
276	Vascular Contributions to Cognitive Impairment and Dementia. Stroke, 2011, 42, 2672-2713.	1.0	2,989
277	Effects of fenofibrate on plasma oxidized LDL and 8-isoprostane in a sub-cohort of GOLDN participants. Atherosclerosis, 2011, 214, 422-425.	0.4	18
278	Association of gene variants with lipid levels in response to fenofibrate is influenced by metabolic syndrome status. Atherosclerosis, 2011, 215, 435-439.	0.4	19
279	A Powerful Test of Parent-of-Origin Effects for Quantitative Traits Using Haplotypes. PLoS ONE, 2011, 6, e28909.	1.1	25
280	Hospice Use in Alabama, 2002-2005. Journal of Pain and Symptom Management, 2011, 41, 374-382.	0.6	5
281	Chocolate consumption is inversely associated with calcified atherosclerotic plaque in the coronary arteries: The NHLBI Family Heart Study†. Clinical Nutrition, 2011, 30, 38-43.	2.3	39
282	Chocolate consumption is inversely associated with prevalent coronary heart disease: The National Heart, Lung, and Blood Institute Family Heart Study. Clinical Nutrition, 2011, 30, 182-187.	2.3	67
283	Genetics of Hypertension and Cardiovascular Disease and Their Interconnected Pathways: Lessons from Large Studies. Current Hypertension Reports, 2011, 13, 46-54.	1.5	33
284	A data-driven method for identifying rare variants with heterogeneous trait effects. Genetic Epidemiology, 2011, 35, 679-685.	0.6	14
285	Variants on Chromosome 6p22.3 Associated With Blood Pressure in the HyperGEN Study: Follow-Up of FBPP Quantitative Trait Loci. American Journal of Hypertension, 2011, 24, 1227-1233.	1.0	16
286	Genetic Variation in NCAM1 Contributes to Left Ventricular Wall Thickness in Hypertensive Families. Circulation Research, 2011, 108, 279-283.	2.0	47
287	Inaccurate Information in Capitol Health Call Article. JAMA - Journal of the American Medical Association, 2011, 305, 256.	3.8	0
288	A High Intake of Saturated Fatty Acids Strengthens the Association between the Fat Mass and Obesity-Associated Gene and BMI. Journal of Nutrition, 2011, 141, 2219-2225.	1.3	111

#	ARTICLE	IF	CITATIONS
289	Pharmacogenetic Effect of the Stromelysin (MMP3) Polymorphism on Stroke Risk in Relation to Antihypertensive Treatment. <i>Stroke</i> , 2011, 42, 330-335.	1.0	26
290	Association of Adiponectin With Left Ventricular Mass in Blacks. <i>Circulation: Heart Failure</i> , 2011, 4, 747-753.	1.6	35
291	Characterization of autosomal copy-number variation in African Americans: the HyperGEN Study. <i>European Journal of Human Genetics</i> , 2011, 19, 1271-1275.	1.4	7
292	Genetic Loci Associated with Plasma Phospholipid n-3 Fatty Acids: A Meta-Analysis of Genome-Wide Association Studies from the CHARGE Consortium. <i>PLoS Genetics</i> , 2011, 7, e1002193.	1.5	324
293	The PLIN4 Variant rs8887 Modulates Obesity Related Phenotypes in Humans through Creation of a Novel miR-522 Seed Site. <i>PLoS ONE</i> , 2011, 6, e17944.	1.1	51
294	Pharmacogenetic Associations of MMP9 and MMP12 Variants with Cardiovascular Disease in Patients with Hypertension. <i>PLoS ONE</i> , 2011, 6, e23609.	1.1	38
295	Genome-Wide Detection of Allele Specific Copy Number Variation Associated with Insulin Resistance in African Americans from the HyperGEN Study. <i>PLoS ONE</i> , 2011, 6, e24052.	1.1	45
296	Biomarkers of inflammation and hemostasis associated with left ventricular mass: The Multiethnic Study of Atherosclerosis (MESA). <i>International Journal of Molecular Epidemiology and Genetics</i> , 2011, 2, 391-400.	0.4	22
297	Gene Expression Algorithm for Prevalent Coronary Artery Disease: A First Step in a Long Journey. <i>Annals of Internal Medicine</i> , 2010, 153, 473.	2.0	3
298	Measurement of Erythrocyte Methotrexate Polyglutamate Levels: Ready for Clinical Use in Rheumatoid Arthritis?. <i>Current Rheumatology Reports</i> , 2010, 12, 342-347.	2.1	28
299	Generalized bone loss as a predictor of three-year radiographic damage in African American patients with recent-onset rheumatoid arthritis. <i>Arthritis and Rheumatism</i> , 2010, 62, 2219-2226.	6.7	28
300	Genetic variants in human CLOCK associate with total energy intake and cytokine sleep factors in overweight subjects (GOLDN population). <i>European Journal of Human Genetics</i> , 2010, 18, 364-369.	1.4	81
301	Mixed linear model approach adapted for genome-wide association studies. <i>Nature Genetics</i> , 2010, 42, 355-360.	9.4	2,022
302	Comparison of the Framingham Heart Study Hypertension Model With Blood Pressure Alone in the Prediction of Risk of Hypertension. <i>Hypertension</i> , 2010, 55, 1339-1345.	1.3	61
303	Defining and Setting National Goals for Cardiovascular Health Promotion and Disease Reduction. <i>Circulation</i> , 2010, 121, 586-613.	1.6	3,508
304	Apolipoprotein E Polymorphisms and Postprandial Triglyceridemia Before and After Fenofibrate Treatment in the Genetics of Lipid Lowering and Diet Network (GOLDN) Study. <i>Circulation: Cardiovascular Genetics</i> , 2010, 3, 462-467.	5.1	39
305	Heritability and Genetic Linkage of Left Ventricular Mass, Systolic and Diastolic Function in Hypertensive African Americans (From the GENOA Study). <i>American Journal of Hypertension</i> , 2010, 23, 870-875.	1.0	20
306	Genetic ancestry and lower extremity peripheral artery disease in the Multi-Ethnic Study of Atherosclerosis. <i>Vascular Medicine</i> , 2010, 15, 351-359.	0.8	36

#	ARTICLE	IF	CITATIONS
307	Differences in Albuminuria Between Hispanics and Whites: An Evaluation by Genetic Ancestry and Country of Origin. <i>Circulation: Cardiovascular Genetics</i> , 2010, 3, 240-247.	5.1	34
308	Relationship of interleukin-6 with regional and global left-ventricular function in asymptomatic individuals without clinical cardiovascular disease: insights from the Multi-Ethnic Study of Atherosclerosis. <i>European Heart Journal</i> , 2010, 31, 875-882.	1.0	66
309	Apolipoprotein B genetic variants modify the response to fenofibrate: a GOLDN study. <i>Journal of Lipid Research</i> , 2010, 51, 3316-3323.	2.0	31
310	Association between hyperuricemia and incident heart failure among older adults: A propensity-matched study. <i>International Journal of Cardiology</i> , 2010, 142, 279-287.	0.8	92
311	Effect of fenofibrate therapy and ABCA1 polymorphisms on high-density lipoprotein subclasses in the Genetics of Lipid Lowering Drugs and Diet Network. <i>Molecular Genetics and Metabolism</i> , 2010, 100, 118-122.	0.5	22
312	Comparison of Postprandial Responses to a High-Fat Meal in Hypertriglyceridemic Men and Women before and after Treatment with Fenofibrate in the Genetics and Lipid Lowering Drugs and Diet Network (GOLDN) Study. <i>SRX Pharmacology</i> , 2010, 2010, 1-8.	0.2	3
313	Fasting Triglyceride Concentrations are Associated with Early Mortality Following Antiretroviral Therapy in Zambia. <i>North American Journal of Medicine & Science</i> , 2010, 3, 079.	3.8	4
314	The launch of international journal of molecular epidemiology and genetics. <i>International Journal of Molecular Epidemiology and Genetics</i> , 2010, 1, i-ii.	0.4	0
315	Editorial board of international journal of molecular epidemiology and genetics (as of march 31,) Tj ETQq1 1 0.784314 rgBT /Overlock	0.4	0
316	Adducin 1 (alpha) Gly460Trp variant is associated with left ventricular geometry in Caucasians and African Americans: The HyperGEN Study. <i>International Journal of Molecular Epidemiology and Genetics</i> , 2010, 1, 367-76.	0.4	1
317	Kidney Function Influences Warfarin Responsiveness and Hemorrhagic Complications. <i>Journal of the American Society of Nephrology: JASN</i> , 2009, 20, 912-921.	3.0	256
318	Genetic Ancestry Is Associated With Subclinical Cardiovascular Disease in African-Americans and Hispanics From the Multi-Ethnic Study of Atherosclerosis. <i>Circulation: Cardiovascular Genetics</i> , 2009, 2, 629-636.	5.1	71
319	Associations of Genetic Variants in ATP-Binding Cassette A1 and Cholesteryl Ester Transfer Protein and Differences in Lipoprotein Subclasses in the Multi-Ethnic Study of Atherosclerosis. <i>Clinical Chemistry</i> , 2009, 55, 481-488.	1.5	18
320	Genetic Variants at the PDZ-Interacting Domain of the Scavenger Receptor Class B Type I Interact with Diet to Influence the Risk of Metabolic Syndrome in Obese Men and Women. <i>Journal of Nutrition</i> , 2009, 139, 842-848.	1.3	19
321	Genetic Variants Associated With Cardiac Structure and Function. <i>JAMA - Journal of the American Medical Association</i> , 2009, 302, 168.	3.8	202
322	Genome-Wide Association Study of Plasma Polyunsaturated Fatty Acids in the InCHIANTI Study. <i>PLoS Genetics</i> , 2009, 5, e1000338.	1.5	351
323	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, 552-558.	1.0	1
324	CLOCK genetic variation and metabolic syndrome risk: modulation by monounsaturated fatty acids. <i>American Journal of Clinical Nutrition</i> , 2009, 90, 1466-1475.	2.2	144

#	ARTICLE	IF	CITATIONS
325	Novel variants at KCTD10, MVK, and MMAB genes interact with dietary carbohydrates to modulate HDL-cholesterol concentrations in the Genetics of Lipid Lowering Drugs and Diet Network Study. <i>American Journal of Clinical Nutrition</i> , 2009, 90, 686-694.	2.2	25
326	Isolated Systolic Hypertension and Incident Heart Failure in Older Adults. <i>Hypertension</i> , 2009, 53, 458-465.	1.3	65
327	Criteria for Evaluation of Novel Markers of Cardiovascular Risk. <i>Circulation</i> , 2009, 119, 2408-2416.	1.6	998
328	Polyunsaturated Fatty Acids Modulate the Effect of TCF7L2 Gene Variants on Postprandial Lipemia. <i>Journal of Nutrition</i> , 2009, 139, 439-446.	1.3	45
329	Association between glucokinase regulatory protein (GCKR) and apolipoprotein A5 (APOA5) gene polymorphisms and triacylglycerol concentrations in fasting, postprandial, and fenofibrate-treated states. <i>American Journal of Clinical Nutrition</i> , 2009, 89, 391-399.	2.2	52
330	Preparing Effective Grant Applications. <i>Circulation</i> , 2009, 120, 2607-2612.	1.6	2
331	Genetic Effect on Blood Pressure Is Modulated by Age. <i>Hypertension</i> , 2009, 53, 35-41.	1.3	56
332	Pharmacogenetics of antihypertensive treatment: detailing disciplinary dissonance. <i>Pharmacogenomics</i> , 2009, 10, 1295-1307.	0.6	29
333	Value of Orthopnea, Paroxysmal Nocturnal Dyspnea, and Medications in Prospective Population Studies of Incident Heart Failure. <i>American Journal of Cardiology</i> , 2009, 104, 259-264.	0.7	27
334	Pharmacogenetics of the response to antihypertensive drugs. <i>Current Cardiovascular Risk Reports</i> , 2009, 3, 441-451.	0.8	0
335	A review of the role of atrial natriuretic peptide gene polymorphisms in hypertension and its sequelae. <i>Current Hypertension Reports</i> , 2009, 11, 35-42.	1.5	39
336	Genome-wide association study identifies single-nucleotide polymorphism in KCNB1 associated with left ventricular mass in humans: The HyperGEN Study. <i>BMC Medical Genetics</i> , 2009, 10, 43.	2.1	43
337	Adverse association between diabetic retinopathy and cardiac structure and function. <i>American Heart Journal</i> , 2009, 157, 563-568.	1.2	24
338	Smoking, inflammatory patterns and postprandial hypertriglyceridemia. <i>Atherosclerosis</i> , 2009, 203, 633-639.	0.4	33
339	Physical inactivity interacts with an endothelial lipase polymorphism to modulate high density lipoprotein cholesterol in the GOLDN study. <i>Atherosclerosis</i> , 2009, 206, 500-504.	0.4	33
340	Has pharmacogenetics brought us closer to "personalized medicine"™ for initial drug treatment of hypertension?. <i>Current Opinion in Cardiology</i> , 2009, 24, 333-339.	0.8	24
341	Pharmacogenetic association of the APOA1/C3/A4/A5 gene cluster and lipid responses to fenofibrate: the Genetics of Lipid-Lowering Drugs and Diet Network study. <i>Pharmacogenetics and Genomics</i> , 2009, 19, 161-169.	0.7	45
342	Novel genetic variants contributing to left ventricular hypertrophy: the HyperGEN study. <i>Journal of Hypertension</i> , 2009, 27, 1585-1593.	0.3	31

#	ARTICLE	IF	CITATIONS
343	The SCARB1 gene is associated with lipid response to dietary and pharmacological interventions. <i>Journal of Human Genetics</i> , 2008, 53, 709-717.	1.1	32
344	The genetic architecture of fasting plasma triglyceride response to fenofibrate treatment. <i>European Journal of Human Genetics</i> , 2008, 16, 603-613.	1.4	35
345	Relation of Albuminuria to Left Ventricular Mass (from the HyperGEN Study). <i>American Journal of Cardiology</i> , 2008, 101, 212-216.	0.7	17
346	Circulating soluble ICAM-1 levels shows linkage to ICAM gene cluster region on chromosome 19: The NHLBI Family Heart Study follow-up examination. <i>Atherosclerosis</i> , 2008, 199, 172-178.	0.4	16
347	Cholesteryl ester transfer protein genetic polymorphisms, HDL cholesterol, and subclinical cardiovascular disease in the Multi-Ethnic Study of Atherosclerosis. <i>Atherosclerosis</i> , 2008, 200, 359-367.	0.4	73
348	ICAM1 and VCAM1 polymorphisms, coronary artery calcium, and circulating levels of soluble ICAM-1: The multi-ethnic study of atherosclerosis (MESA). <i>Atherosclerosis</i> , 2008, 201, 339-344.	0.4	54
349	Association between endothelial biomarkers and arterial elasticity in young adults: the CARDIA Study. <i>Journal of the American Society of Hypertension</i> , 2008, 2, 70-79.	2.3	14
350	Retinal Arteriolar Narrowing and Left Ventricular Hypertrophy in African Americans. The Atherosclerosis Risk in Communities (ARIC) Study. <i>American Journal of Hypertension</i> , 2008, 21, 352-359.	1.0	48
351	Association of Common C-Reactive Protein (<i>CRP</i>) Gene Polymorphisms With Baseline Plasma CRP Levels and Fenofibrate Response. <i>Diabetes Care</i> , 2008, 31, 910-915.	4.3	44
352	Erythrocyte Fatty Acid Composition and the Metabolic Syndrome: A National Heart, Lung, and Blood Institute GOLDN Study. <i>Clinical Chemistry</i> , 2008, 54, 154-162.	1.5	59
353	A Whole-Genome Scan for Stroke or Myocardial Infarction in Family Blood Pressure Program Families. <i>Stroke</i> , 2008, 39, 1115-1120.	1.0	9
354	The effect of IL6-174C/G polymorphism on postprandial triglyceride metabolism in the GOLDN study*. <i>Journal of Lipid Research</i> , 2008, 49, 1839-1845.	2.0	22
355	Trends in cigarette smoking: The Minnesota Heart Survey, 1980-1982 through 2000-2002. <i>Nicotine and Tobacco Research</i> , 2008, 10, 827-832.	1.4	12
356	Pharmacogenetic Association of the NPPA T2238C Genetic Variant With Cardiovascular Disease Outcomes in Patients With Hypertension. <i>JAMA - Journal of the American Medical Association</i> , 2008, 299, 296-307.	3.8	97
357	Common Missense Variant in the Glucokinase Regulatory Protein Gene Is Associated With Increased Plasma Triglyceride and C-Reactive Protein but Lower Fasting Glucose Concentrations. <i>Diabetes</i> , 2008, 57, 3112-3121.	0.3	264
358	Vascular Stiffness and the "Chicken-or-the-Egg" Question. <i>Hypertension</i> , 2008, 51, 177-178.	1.3	8
359	Postprandial triacylglycerol metabolism is modified by the presence of genetic variation at the perilipin (PLIN) locus in 2 white populations. <i>American Journal of Clinical Nutrition</i> , 2008, 87, 744-752.	2.2	27
360	Ethnicity and sex modify the association of serum c-reactive protein with microalbuminuria. <i>Ethnicity and Disease</i> , 2008, 18, 324-9.	1.0	6

#	ARTICLE	IF	CITATIONS
361	The ϵ 256T>C Polymorphism in the Apolipoprotein A-II Gene Promoter Is Associated with Body Mass Index and Food Intake in the Genetics of Lipid Lowering Drugs and Diet Network Study. <i>Clinical Chemistry</i> , 2007, 53, 1144-1152.	1.5	113
362	Genome scan of glomerular filtration rate and albuminuria: the HyperGEN study. <i>Nephrology Dialysis Transplantation</i> , 2007, 22, 763-771.	0.4	34
363	Genetic Variations Associated With Echocardiographic Left Ventricular Traits in Hypertensive Blacks. <i>Hypertension</i> , 2007, 49, 992-999.	1.3	27
364	Summary of the American Heart Association's Scientific Statement on the Relevance of Genetics and Genomics for Prevention and Treatment of Cardiovascular Disease. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2007, 27, 1682-1686.	1.1	7
365	Genotype-by-Sex Interaction on Fasting Insulin Concentration: The HyperGEN Study. <i>Diabetes</i> , 2007, 56, 137-142.	0.3	19
366	Family History of Premature Coronary Heart Disease and Coronary Artery Calcification. <i>Circulation</i> , 2007, 116, 619-626.	1.6	160
367	Genome-Wide Linkage Mapping for Valve Calcification Susceptibility Loci in Hypertensive Sibships. <i>Hypertension</i> , 2007, 49, 453-460.	1.3	36
368	Fenofibrate Effect on Triglyceride and Postprandial Response of Apolipoprotein A5 Variants. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2007, 27, 1417-1425.	1.1	113
369	Relevance of Genetics and Genomics for Prevention and Treatment of Cardiovascular Disease. <i>Circulation</i> , 2007, 115, 2878-2901.	1.6	180
370	A New Era of Cardiovascular Disease Epidemiology. <i>JAMA - Journal of the American Medical Association</i> , 2007, 298, 2060.	3.8	5
371	Left ventricular geometric patterns in the Jackson cohort of the atherosclerotic risk in communities (ARIC) study: Clinical correlates and influences on systolic and diastolic dysfunction. <i>American Heart Journal</i> , 2007, 153, 238-244.	1.2	68
372	Absence of an interaction between the angiotensin-converting enzyme insertion-deletion polymorphism and pravastatin on cardiovascular disease in high-risk hypertensive patients: The Genetics of Hypertension-Associated Treatment (GenHAT) study. <i>American Heart Journal</i> , 2007, 153, 54-58.	1.2	25
373	A Whole Genome Scan for Pulse Pressure/Stroke Volume Ratio in African Americans: The HyperGEN Study. <i>American Journal of Hypertension</i> , 2007, 20, 398-402.	1.0	25
374	Evidence of QTL on 15q21 for high-density lipoprotein cholesterol: The National Heart, Lung, and Blood Institute Family Heart Study (NHLBI FHS). <i>Atherosclerosis</i> , 2007, 190, 232-237.	0.4	8
375	Implications of C-reactive protein or coronary artery calcium score as an adjunct to global risk assessment for primary prevention of CHD. <i>Atherosclerosis</i> , 2007, 193, 401-407.	0.4	22
376	Plasma adiponectin concentrations and correlates in African Americans in the Hypertension Genetic Epidemiology Network (HyperGEN) study. <i>Metabolism: Clinical and Experimental</i> , 2007, 56, 1011-1016.	1.5	10
377	Interleukin 1 β Genetic Polymorphisms Interact with Polyunsaturated Fatty Acids to Modulate Risk of the Metabolic Syndrome, 3. <i>Journal of Nutrition</i> , 2007, 137, 1846-1851.	1.3	59
378	Trends in Diet Quality for Coronary Heart Disease Prevention between 1980-1982 and 2000-2002: The Minnesota Heart Survey. <i>Journal of the American Dietetic Association</i> , 2007, 107, 213-222.	1.3	42

#	ARTICLE	IF	CITATIONS
379	Left Ventricular Architecture and Survival in African-Americans Free of Coronary Heart Disease (from) Tj ETQq1 1 0.784314 rgBT /Overlo 1413-1420.	0.7	40
380	Sex-specific effects of ACE I/D and AGT-M235T on pulse pressure: the HyperGEN Study. <i>Human Genetics</i> , 2007, 122, 33-40.	1.8	6
381	Familial Clustering for Features of the Metabolic Syndrome: The National Heart, Lung, and Blood Institute (NHLBI) Family Heart Study. <i>Diabetes Care</i> , 2006, 29, 631-636.	4.3	67
382	Contemporary Definitions and Classification of the Cardiomyopathies. <i>Circulation</i> , 2006, 113, 1807-1816.	1.6	2,935
383	Hypertension and Smoking Are Associated With Reduced Regional Left Ventricular Function in Asymptomatic Individuals. <i>Journal of the American College of Cardiology</i> , 2006, 47, 1150-1158.	1.2	118
384	Traditional Cardiovascular Risk Factors in Relation to Left Ventricular Mass, Volume, and Systolic Function by Cardiac Magnetic Resonance Imaging. <i>Journal of the American College of Cardiology</i> , 2006, 48, 2285-2292.	1.2	262
385	Use of Nonprescription Medications for Perceived Cardiovascular Health. <i>American Journal of Preventive Medicine</i> , 2006, 30, 78-81.	1.6	24
386	Trends in Blood Pressure, Hypertension Control, and Stroke Mortality: The Minnesota Heart Survey. <i>American Journal of Medicine</i> , 2006, 119, 42-49.	0.6	83
387	The prognostic value of the mitral diastolic filling velocity ratio for all-cause mortality and cardiovascular morbidity in African Americans: the atherosclerotic Risks in Communities (ARIC) study. <i>American Heart Journal</i> , 2006, 152, 749-755.	1.2	13
388	Population Trends in Leisure-Time Physical Activity. <i>Medicine and Science in Sports and Exercise</i> , 2006, 38, 1716-1723.	0.2	67
389	Genome-wide linkage scans for loci affecting total cholesterol, HDL-C, and triglycerides: the Family Blood Pressure Program. <i>Human Genetics</i> , 2006, 120, 371-380.	1.8	19
390	Pharmacogenetics of antihypertensive treatment. <i>Vascular Pharmacology</i> , 2006, 44, 107-118.	1.0	78
391	Blood Pressure Stress Reactivity and Left Ventricular Mass in a Random Community Sample of African-American and Caucasian Men and Women. <i>American Journal of Cardiology</i> , 2006, 97, 240-244.	0.7	23
392	Racial Differences in the Association of Coronary Calcified Plaque With Left Ventricular Hypertrophy: The National Heart, Lung, and Blood Institute Family Heart Study and Hypertension Genetic Epidemiology Network. <i>American Journal of Cardiology</i> , 2006, 97, 1441-1448.	0.7	17
393	Association of Coronary Artery Calcified Plaque With Clinical Coronary Heart Disease in the National Heart, Lung, and Blood Institute's Family Heart Study. <i>American Journal of Cardiology</i> , 2006, 97, 1564-1569.	0.7	22
394	Alcohol Consumption and Plasma Atrial Natriuretic Peptide (from The HyperGEN Study). <i>American Journal of Cardiology</i> , 2006, 98, 628-632.	0.7	21
395	Quantitative Trait Loci on Chromosome 8q24 for Pancreatic Î-Cell Function and 7q11 for Insulin Sensitivity in Obese Nondiabetic White and Black Families: Evidence From Genome-Wide Linkage Scans in the NHLBI Hypertension Genetic Epidemiology Network (HyperGEN) Study. <i>Diabetes</i> , 2006, 55, 551-558.	0.3	17
396	Obtaining Informed Consent for Genetic Studies. <i>American Journal of Epidemiology</i> , 2006, 164, 845-851.	1.6	26

#	ARTICLE	IF	CITATIONS
397	Interarm differences in seated systolic and diastolic blood pressure: the Hypertension Genetic Epidemiology Network study. <i>Journal of Hypertension</i> , 2005, 23, 1141-1147.	0.3	37
398	Relation of aortic valve sclerosis to risk of coronary heart disease in African-Americans. <i>American Journal of Cardiology</i> , 2005, 95, 401-404.	0.7	60
399	Effect of influenza vaccine on markers of inflammation and lipid profile. <i>Translational Research</i> , 2005, 145, 323-327.	2.4	89
400	Body composition and fat distribution influence systemic hemodynamics in the absence of obesity: the HyperGEN Study. <i>American Journal of Clinical Nutrition</i> , 2005, 81, 757-761.	2.2	43
401	Dietary Linolenic Acid Is Associated With a Lower Prevalence of Hypertension in the NHLBI Family Heart Study. <i>Hypertension</i> , 2005, 45, 368-373.	1.3	60
402	Metabolic Syndrome and Echocardiographic Left Ventricular Mass in Blacks. <i>Circulation</i> , 2005, 112, 819-827.	1.6	96
403	Genome-Wide Linkage Analysis for Loci Affecting Pulse Pressure. <i>Hypertension</i> , 2005, 46, 1286-1293.	1.3	42
404	A Population Association Study of Angiotensinogen Polymorphisms and Haplotypes With Left Ventricular Phenotypes. <i>Hypertension</i> , 2005, 46, 1294-1299.	1.3	16
405	Twenty-Year Trends in Serum Cholesterol, Hypercholesterolemia, and Cholesterol Medication Use. <i>Circulation</i> , 2005, 112, 3884-3891.	1.6	153
406	Optimal Threshold Value for Left Ventricular Hypertrophy in Blacks. <i>Hypertension</i> , 2005, 45, 58-63.	1.3	48
407	Pharmacogenetic Association of the Angiotensin-Converting Enzyme Insertion/Deletion Polymorphism on Blood Pressure and Cardiovascular Risk in Relation to Antihypertensive Treatment. <i>Circulation</i> , 2005, 111, 3374-3383.	1.6	133
408	Overweight in Children and Adolescents. <i>Circulation</i> , 2005, 111, 1999-2012.	1.6	1,234
409	Dietary Linolenic Acid Is Inversely Associated With Calcified Atherosclerotic Plaque in the Coronary Arteries. <i>Circulation</i> , 2005, 111, 2921-2926.	1.6	109
410	The Impact of Pedigree Structure on Heritability Estimates for Pulse Pressure in Three Studies. <i>Human Heredity</i> , 2005, 60, 63-72.	0.4	17
411	Haplotype Association Analysis of AGT Variants with Hypertension-Related Traits: The HyperGEN Study. <i>Human Heredity</i> , 2005, 60, 164-176.	0.4	22
412	Left Ventricular Mass Indexed to Height and Prevalent MRI Cerebrovascular Disease in an African American Cohort. <i>Stroke</i> , 2005, 36, 546-550.	1.0	23
413	Left ventricular concentric geometry is associated with impaired relaxation in hypertension: the HyperGEN study. <i>European Heart Journal</i> , 2005, 26, 1039-1045.	1.0	97
414	Evidence for a gene influencing fasting LDL cholesterol and triglyceride levels on chromosome 21q. <i>Atherosclerosis</i> , 2005, 179, 119-125.	0.4	21

#	ARTICLE	IF	CITATIONS
415	Identification of a novel 5â€“base pair deletion in calcineurin B (PPP3R1) promoter region and its association with left ventricular hypertrophy. <i>American Heart Journal</i> , 2005, 150, 845-851.	1.2	31
416	A Genome Scan for Linkage With Aortic Root Diameter in Hypertensive African Americans and Whites in the Hypertension Genetic Epidemiology Network (HyperGEN) Study. <i>American Journal of Hypertension</i> , 2005, 18, 627-632.	1.0	4
417	Left Atrial Systolic Force and Cardiac Markers of Preclinical Disease in Hypertensive PatientsThe Hypertension Genetic Epidemiology Network (HyperGEN) Study. <i>American Journal of Hypertension</i> , 2005, 18, 899-905.	1.0	25
418	A Summary of the Effects of Antihypertensive Medications on Measured Blood Pressure. <i>American Journal of Hypertension</i> , 2005, 18, 935-942.	1.0	102
419	Dietary Linolenic Acid and Adjusted QT and JT Intervals in the National Heart, Lung, and Blood Institute Family Heart Study. <i>Journal of the American College of Cardiology</i> , 2005, 45, 1716-1722.	1.2	32
420	Left Ventricular Concentric Remodeling Is Associated With Decreased Global and Regional Systolic Function: The Multi-Ethnic Study of Atherosclerosis. <i>Circulation</i> , 2005, 112, 984-991.	1.6	129
421	Apolipoprotein E polymorphism modifies the alcohol-HDL association observed in the National Heart, Lung, and Blood Institute Family Heart Study. <i>American Journal of Clinical Nutrition</i> , 2004, 80, 1639-1644.	2.2	35
422	Atherosclerotic Vascular Disease Conference. <i>Circulation</i> , 2004, 109, 2613-2616.	1.6	85
423	Differences in Left Ventricular Structure Between Black and White Hypertensive Adults. <i>Hypertension</i> , 2004, 43, 1182-1188.	1.3	187
424	Linkage Analysis of Diabetes Status Among Hypertensive Families: The Hypertension Genetic Epidemiology Network Study. <i>Diabetes</i> , 2004, 53, 3307-3312.	0.3	19
425	Comparison of Ultracentrifugation and Nuclear Magnetic Resonance Spectroscopy in the Quantification of Triglyceride-Rich Lipoproteins after an Oral Fat Load. <i>Clinical Chemistry</i> , 2004, 50, 1201-1204.	1.5	49
426	Alcohol Consumption and Metabolic Syndrome: Does the Type of Beverage Matter?. <i>Obesity</i> , 2004, 12, 1375-1385.	4.0	119
427	Association of kidney function and hemoglobin with left ventricular morphology among African Americans: The Atherosclerosis Risk in Communities (ARIC) study. <i>American Journal of Kidney Diseases</i> , 2004, 43, 836-845.	2.1	68
428	Accuracy of estimation of large food portions. <i>Journal of the American Dietetic Association</i> , 2004, 104, 804-806.	1.3	54
429	Comparison of the prognostic value of left ventricular hypertrophy in African-American men versus women. <i>American Journal of Cardiology</i> , 2004, 94, 1383-1390.	0.7	21
430	Genome-wide linkage analysis replicates susceptibility locus for fasting plasma triglycerides: NHLBI Family Heart Study. <i>Human Genetics</i> , 2004, 115, 468-474.	1.8	13
431	Genes for left ventricular hypertrophy. <i>Current Hypertension Reports</i> , 2004, 6, 36-41.	1.5	40
432	Imputing geneâ€“treatment interactions when the genotype distribution is unknown using case-only and putative placebo analysesâ€“a new method for the Genetics of Hypertension Associated Treatment(GenHAT) study. <i>Statistics in Medicine</i> , 2004, 23, 2413-2427.	0.8	10

#	ARTICLE	IF	CITATIONS
433	Pharmacogenetics of antihypertensive treatment. Drug Development Research, 2004, 62, 191-199.	1.4	2
434	Lifestyle determinants of high-density lipoprotein cholesterol: the National Heart, Lung, and Blood Institute Family Heart Study. American Heart Journal, 2004, 147, 529-535.	1.2	153
435	Arterial stiffness is greater in African Americans than in whites evidence from the Forsyth County, North Carolina, ARIC cohort. American Journal of Hypertension, 2004, 17, 304-313.	1.0	105
436	Association of inappropriate left ventricular mass with systolic and diastolic dysfunction: the HyperGEN study. American Journal of Hypertension, 2004, 17, 828-833.	1.0	24
437	Genome-wide linkage analyses for age at diagnosis of hypertension and early-onset hypertension in the HyperGEN study. American Journal of Hypertension, 2004, 17, 839-844.	1.0	30
438	Fruit and vegetable consumption and LDL cholesterol: the National Heart, Lung, and Blood Institute Family Heart Study. American Journal of Clinical Nutrition, 2004, 79, 213-217.	2.2	144
439	Left ventricular structure and systolic function in African Americans: the Atherosclerosis Risk in Communities (ARIC) study. Ethnicity and Disease, 2004, 14, 483-8.	1.0	5
440	Trends in the trans-fatty acid composition of the diet in a metropolitan area: The Minnesota Heart Survey. Journal of the American Dietetic Association, 2003, 103, 1160-1166.	1.3	45
441	Echocardiographic Left Ventricular Mass in African-Americans. The Jackson Cohort of the Atherosclerosis Risk in Communities Study. Echocardiography, 2003, 20, 111-120.	0.3	49
442	Associations of Aging and Birth Cohort with Body Mass Index in a Biethnic Cohort. Obesity, 2003, 11, 426-433.	4.0	28
443	Associations of aortic and mitral regurgitation with body composition and myocardial energy expenditure in adults with hypertension: the Hypertension Genetic Epidemiology Network study. American Heart Journal, 2003, 145, 1071-1077.	1.2	25
444	Associations of weight loss and changes in fat distribution with the remission of hypertension in a bi-ethnic cohort: the Atherosclerosis Risk in Communities Study. Preventive Medicine, 2003, 36, 330-339.	1.6	17
445	A genome-wide affected sibpair linkage analysis of hypertension: the HyperGEN network. American Journal of Hypertension, 2003, 16, 148-150.	1.0	65
446	Prevalence and correlates of aortic valve sclerosis in hypertensive adults: the hypergen study. American Journal of Hypertension, 2003, 16, A5.	1.0	0
447	The Arg16Gly polymorphism of the β_2 -adrenergic receptor and left ventricular systolic function. American Journal of Hypertension, 2003, 16, 945-951.	1.0	35
448	Linkage Analysis of a Composite Factor for the Multiple Metabolic Syndrome: The National Heart, Lung, and Blood Institute Family Heart Study. Diabetes, 2003, 52, 2840-2847.	0.3	89
449	A Genome-Wide Scan for Urinary Albumin Excretion in Hypertensive Families. Hypertension, 2003, 42, 291-296.	1.3	67
450	A Genome-Wide Scan of Pulmonary Function Measures in the National Heart, Lung, and Blood Institute Family Heart Study. American Journal of Respiratory and Critical Care Medicine, 2003, 167, 1528-1533.	2.5	43

#	ARTICLE	IF	CITATIONS
451	Dietary linolenic acid is inversely associated with plasma triacylglycerol: the National Heart, Lung, and Blood Institute Family Heart Study. <i>American Journal of Clinical Nutrition</i> , 2003, 78, 1098-1102.	2.2	71
452	Relation Between Serum Albumin and Carotid Atherosclerosis. <i>Stroke</i> , 2003, 34, 53-57.	1.0	20
453	Task Force on Strategic Research Direction. <i>Circulation</i> , 2002, 106, e167-72.	1.6	5
454	Socioeconomic Disadvantage and Change in Blood Pressure Associated With Aging. <i>Circulation</i> , 2002, 106, 703-710.	1.6	85
455	Sibling Resemblance for Left Ventricular Structure, Contractility, and Diastolic Filling. <i>Hypertension</i> , 2002, 40, 233-238.	1.3	16
456	Fifteen-Year Trends in Cardiovascular Risk Factors (1980-1982 through 1995-1997): The Minnesota Heart Survey. <i>American Journal of Epidemiology</i> , 2002, 156, 929-935.	1.6	109
457	Appetite suppressants and valvular heart disease in a population-based sample: the HyperGEN study. <i>American Journal of Medicine</i> , 2002, 112, 710-715.	0.6	32
458	Associations between angiotensinogen gene variants and left ventricular mass and function in the HyperGEN study. <i>American Heart Journal</i> , 2002, 143, 854-860.	1.2	38
459	Quantitative-Trait Loci Influencing Body-Mass Index Reside on Chromosomes 7 and 13: The National Heart, Lung, and Blood Institute Family Heart Study. <i>American Journal of Human Genetics</i> , 2002, 70, 72-82.	2.6	138
460	Refined Mapping of Suggestive Linkage to Renal Function in African Americans: The HyperGEN Study. <i>American Journal of Human Genetics</i> , 2002, 71, 204-205.	2.6	14
461	Regarding "Testing for Population Subdivision and Association in Four Case-Control Studies". <i>American Journal of Human Genetics</i> , 2002, 71, 1478-1480.	2.6	28
462	Linkage of left ventricular early diastolic peak filling velocity to chromosome 5 in hypertensive African Americans: the HyperGEN Echocardiography Study ¹ . <i>American Journal of Hypertension</i> , 2002, 15, 621-627.	1.0	14
463	Relation of insulin to left ventricular geometry and function in African American and white hypertensive adults: the HyperGEN study. <i>American Journal of Hypertension</i> , 2002, 15, 1029-1035.	1.0	25
464	Blood pressure responses to acute stress and left ventricular mass (The Hypertension Genetic) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 222	0.7	19
465	Gender difference in diastolic function in hypertension (the HyperGEN study). <i>American Journal of Cardiology</i> , 2002, 89, 1052-1056.	0.7	59
466	A genome-wide screen reveals evidence for a locus on chromosome 11 influencing variation in LDL cholesterol in the NHLBI Family Heart Study. <i>Human Genetics</i> , 2002, 111, 263-269.	1.8	26
467	Segregation analysis of HDL cholesterol in the NHLBI Family Heart Study and in Utah pedigrees. <i>European Journal of Human Genetics</i> , 2002, 10, 367-374.	1.4	15
468	A Genome Scan for Renal Function among Hypertensives: the HyperGEN Study. <i>American Journal of Human Genetics</i> , 2001, 68, 136-144.	2.6	68

#	ARTICLE	IF	CITATIONS
469	Neighborhood of Residence and Incidence of Coronary Heart Disease. <i>New England Journal of Medicine</i> , 2001, 345, 99-106.	13.9	1,529
470	Arterial distensibility and physical activity in the ARIC study. <i>Medicine and Science in Sports and Exercise</i> , 2001, 33, 2065-2071.	0.2	35
471	Association of C-reactive protein with markers of prevalent atherosclerotic disease. <i>American Journal of Cardiology</i> , 2001, 88, 112-117.	0.7	221
472	Relation of various degrees of body mass index in patients with systemic hypertension to left ventricular mass, cardiac output, and peripheral resistance (The Hypertension Genetic Epidemiology) Tj ETQq0 0 0 0 BT /Overlook 10 Tf		
473	Left Ventricular Systolic Dysfunction in a Biracial Sample of Hypertensive Adults. <i>Hypertension</i> , 2001, 38, 417-423.	1.3	65
474	Linkage of Left Ventricular Contractility to Chromosome 11 in Humans. <i>Hypertension</i> , 2001, 38, 767-772.	1.3	29
475	Genome Scan for Quantitative Trait Loci Linked to High-Density Lipoprotein Cholesterol. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2001, 21, 1823-1828.	1.1	59
476	Aortic Root Dilatation at Sinuses of Valsalva and Aortic Regurgitation in Hypertensive and Normotensive Subjects. <i>Hypertension</i> , 2001, 37, 1229-1235.	1.3	128
477	Trends in Acute Coronary Heart Disease Mortality, Morbidity, and Medical Care From 1985 Through 1997. <i>Circulation</i> , 2001, 104, 19-24.	1.6	309
478	Relationship Between Left Ventricular Diastolic Relaxation and Systolic Function in Hypertension. <i>Hypertension</i> , 2001, 38, 424-428.	1.3	49
479	Effect of Type 2 Diabetes Mellitus on Left Ventricular Geometry and Systolic Function in Hypertensive Subjects. <i>Circulation</i> , 2001, 103, 102-107.	1.6	285
480	Pulse Pressure and Prevalence of Peripheral Arterial Disease: The Atherosclerosis Risk in Communities (ARIC) Study. <i>Circulation</i> , 2001, 103, 1361-1361.	1.6	0
481	Inheritance of the Waist-to-Hip Ratio in the National Heart, Lung, and Blood Institute Family Heart Study. <i>Obesity</i> , 2000, 8, 294-301.	4.0	15
482	Hostility, social support, and carotid artery atherosclerosis in The National Heart, Lung, and Blood Institute Family Heart Study. <i>American Journal of Cardiology</i> , 2000, 86, 1086-1089.	0.7	68
483	Smoking influences the association between apolipoprotein E and lipids: The national heart, lung, and blood institute family heart study. <i>Lipids</i> , 2000, 35, 827-831.	0.7	17
484	Genetic contributions to left ventricular hypertrophy. <i>Current Hypertension Reports</i> , 2000, 2, 50-55.	1.5	20
485	Skip Patterns in DINAMAP-Measured Blood Pressure in 3 Epidemiological Studies. <i>Hypertension</i> , 2000, 35, 1032-1036.	1.3	28
486	Replication of Linkage of Familial Combined Hyperlipidemia to Chromosome 1q With Additional Heterogeneous Effect of Apolipoprotein A-I/C-III/A-IV Locus. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2000, 20, 2275-2280.	1.1	82

#	ARTICLE	IF	CITATIONS
487	Alcohol consumption and plasminogen activator inhibitor type 1: The national heart, lung, and blood institute family heart study. <i>American Heart Journal</i> , 2000, 139, 704-709.	1.2	39
488	Hypertension and arterial stiffness: the atherosclerosis risk in communities study*1. <i>American Journal of Hypertension</i> , 2000, 13, 317-323.	1.0	86
489	Orthostatic hypotension and the incidence of coronary heart disease: the atherosclerosis risk in communities study. <i>American Journal of Hypertension</i> , 2000, 13, 571-578.	1.0	220
490	NHLBI Family Blood Pressure Program. <i>Annals of Epidemiology</i> , 2000, 10, 389-400.	0.9	160
491	Differences and Trends in Antioxidant Dietary Intake in Smokers and Non-smokers, 1980-1992. <i>Annals of Epidemiology</i> , 2000, 10, 417-423.	0.9	34
492	No evidence of linkage between the very low-density lipoprotein receptor gene and fasting serum insulin or homeostasis model assessment insulin resistance index: The National Heart, Lung, and Blood Institute Family Heart Study. <i>Metabolism: Clinical and Experimental</i> , 2000, 49, 293-297.	1.5	1
493	Orthostatic Hypotension as a Risk Factor for Stroke. <i>Stroke</i> , 2000, 31, 2307-2313.	1.0	304
494	Influence of leisure time physical activity and television watching on atherosclerosis risk factors in the NHLBI Family Heart Study. <i>Atherosclerosis</i> , 2000, 153, 433-443.	0.4	162
495	Plasma homocysteine and its association with carotid intimal-medial wall thickness and prevalent coronary heart disease: NHLBI Family Heart Study. <i>Atherosclerosis</i> , 2000, 151, 519-524.	0.4	42
496	Arterial Stiffness and the Development of Hypertension. <i>Hypertension</i> , 1999, 34, 201-206.	1.3	479
497	Variability in ultrasonic measurements of arterial stiffness in the atherosclerosis risk in communities study. <i>Ultrasound in Medicine and Biology</i> , 1999, 25, 175-180.	0.7	44
498	Relationship of Serum and Dietary Magnesium to Incident Hypertension. <i>Annals of Epidemiology</i> , 1999, 9, 159-165.	0.9	103
499	Women, Employment Status, and Hypertension. <i>Annals of Epidemiology</i> , 1999, 9, 374-382.	0.9	14
500	Physical Activity and Incident Hypertension in Black and White Adults: The Atherosclerosis Risk in Communities Study. <i>Preventive Medicine</i> , 1999, 28, 304-312.	1.6	149
501	Trends in medical care of hospitalized stroke patients between 1980 and 1990: The minnesota stroke survey. <i>Journal of Stroke and Cerebrovascular Diseases</i> , 1998, 7, 76-84.	0.7	5
502	Lack of Association between Platelet Glycoprotein IIb/IIIa Receptor PIA Polymorphism and Coronary Artery Disease or Carotid Intima-Media Thickness. <i>Thrombosis Research</i> , 1998, 89, 85-89.	0.8	23
503	No Association between Factor V Leiden Mutation and Coronary Heart Disease or Carotid Intima Media Thickness. <i>Thrombosis Research</i> , 1998, 89, 289-293.	0.8	7
504	Evidence for Multiple Determinants of the Body Mass Index: The National Heart, Lung, and Blood Institute Family Heart Study. <i>Obesity</i> , 1998, 6, 107-114.	4.0	64

#	ARTICLE	IF	CITATIONS
505	Angiotensinogen and angiotensin converting enzyme genotypes and carotid atherosclerosis: The atherosclerosis risk in communities and the NHLBI family heart studies. <i>Atherosclerosis</i> , 1998, 138, 111-116.	0.4	33
506	The Importance of Arterial Compliance in Cardiovascular Drug Therapy. <i>Journal of Clinical Pharmacology</i> , 1998, 38, 202-212.	1.0	30
507	Vascular Compliance and Cardiovascular Disease A Risk Factor or a Marker?. <i>American Journal of Hypertension</i> , 1997, 10, 1175-1189.	1.0	245
508	Validity of Electrocardiographic Estimates of Left Ventricular Hypertrophy and Mass in African Americans (The Charleston Heart Study). <i>American Journal of Cardiology</i> , 1997, 79, 1289-1292.	0.7	20
509	Impact of adjustments for intermediate phenotypes on the power to detect linkage. <i>Genetic Epidemiology</i> , 1997, 14, 749-754.	0.6	6
510	Physical activity and incidence of coronary heart disease in middle-aged women and men. <i>Medicine and Science in Sports and Exercise</i> , 1997, 29, 901-909.	0.2	190
511	Variability in B-mode ultrasound measurements in the Atherosclerosis Risk in Communities (ARIC) study. <i>Ultrasound in Medicine and Biology</i> , 1996, 22, 545-554.	0.7	81
512	Gender and Ethnic Differences in Survival in a Cohort of HIV Positive Clients. <i>Ethnicity and Health</i> , 1996, 1, 77-85.	1.5	11
513	Black-white differences in electrocardiographic left ventricular mass and its association with blood pressure (the ARIC study). <i>American Journal of Cardiology</i> , 1994, 74, 247-252.	0.7	59
514	Arterial Stiffness: A New Cardiovascular Risk Factor?. <i>American Journal of Epidemiology</i> , 1994, 140, 669-682.	1.6	436
515	Update on Nitrate Therapy. <i>Journal of Clinical Pharmacology</i> , 1991, 31, 697-701.	1.0	5