Christopher J O'donnell

List of Publications by Year
in descending order

144 papers	20,770 citations	22099 59 h-index	11581 135 g-index
153 all docs	153 docs citations	153 times ranked	30654 citing authors

#	Article	IF	Citations
1	Matrix Gla Protein Levels Are Associated With Arterial Stiffness and Incident Heart Failure With Preserved Ejection Fraction. Arteriosclerosis, Thrombosis, and Vascular Biology, 2022, 42, ATVBAHA121316664.	1.1	10
2	APOL1 Risk Variants, Acute Kidney Injury, and Death in Participants With African Ancestry Hospitalized With COVID-19 From the Million Veteran Program. JAMA Internal Medicine, 2022, 182, 386.	2.6	31
3	Coronary Artery Disease Risk of Familial Hypercholesterolemia Genetic Variants Independent of Clinically Observed Longitudinal Cholesterol Exposure. Circulation Genomic and Precision Medicine, 2022, 15, CIRCGEN121003501.	1.6	6
4	Genetic and clinical determinants of abdominal aortic diameter: genome-wide association studies, exome array data and Mendelian randomization study. Human Molecular Genetics, 2022, 31, 3566-3579.	1.4	5
5	A Phenome-Wide Association Study of genes associated with COVID-19 severity reveals shared genetics with complex diseases in the Million Veteran Program. PLoS Genetics, 2022, 18, e1010113.	1.5	16
6	Integration of rare expression outlier-associated variants improves polygenic risk prediction. American Journal of Human Genetics, 2022, 109, 1055-1064.	2.6	8
7	Genome-wide and phenome-wide analysis of ideal cardiovascular health in the VA Million Veteran Program. PLoS ONE, 2022, 17, e0267900.	1.1	2
8	A multiancestry genome-wide association study of unexplained chronic ALT elevation as a proxy for nonalcoholic fatty liver disease with histological and radiological validation. Nature Genetics, 2022, 54, 761-771.	9.4	68
9	A multi-population phenome-wide association study of genetically-predicted height in the Million Veteran Program. PLoS Genetics, 2022, 18, e1010193.	1.5	12
10	Trends in cardiovascular procedural volumes in the setting of COVIDâ€19: Insights from the VA clinical assessment, reporting, and tracking program. Catheterization and Cardiovascular Interventions, 2021, 98, E326-E328.	0.7	3
11	A Noncoding Variant Near PPP1R3B Promotes Liver Glycogen Storage and MetS, but Protects Against Myocardial Infarction. Journal of Clinical Endocrinology and Metabolism, 2021, 106, 372-387.	1.8	12
12	Comparison of family health history in surveys vs electronic health record data mapped to the observational medical outcomes partnership data model in the All of Us Research Program. Journal of the American Medical Informatics Association: JAMIA, 2021, 28, 695-703.	2.2	11
13	Genome-wide transcriptome study using deep RNA sequencing for myocardial infarction and coronary artery calcification. BMC Medical Genomics, 2021, 14, 45.	0.7	5
14	Actionable druggable genome-wide Mendelian randomization identifies repurposing opportunities for COVID-19. Nature Medicine, 2021, 27, 668-676.	15.2	120
15	Genetic Contribution to Common Heart Failureâ€"Not So Rare?. JAMA Cardiology, 2021, 6, 387.	3.0	1
16	Rural-Urban Differences in Mortality From Ischemic Heart Disease, Heart Failure, and Stroke in the United States. Circulation: Cardiovascular Quality and Outcomes, 2021, 14, e007341.	0.9	18
17	Phenome-wide association of 1809 phenotypes and COVID-19 disease progression in the Veterans Health Administration Million Veteran Program. PLoS ONE, 2021, 16, e0251651.	1.1	17
18	Genetic analysis in European ancestry individuals identifies 517 loci associated with liver enzymes. Nature Communications, 2021, 12, 2579.	5.8	51

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19	Plasma Protein Profile of Carotid Artery Atherosclerosis and Atherosclerotic Outcomes. Arteriosclerosis, Thrombosis, and Vascular Biology, 2021, 41, 1777-1788.	1.1	18
20	Determinants of penetrance and variable expressivity in monogenic metabolic conditions across 77,184 exomes. Nature Communications, 2021, 12, 3505.	5.8	49
21	Meta-analysis of epigenome-wide association studies of carotid intima-media thickness. European Journal of Epidemiology, 2021, 36, 1143-1155.	2.5	10
22	Discovery and prioritization of variants and genes for kidney function in >1.2 million individuals. Nature Communications, 2021, 12, 4350.	5.8	125
23	Association of Apparent Treatment-Resistant Hypertension With Differential Risk of End-Stage Kidney Disease Across Racial Groups in the Million Veteran Program. Hypertension, 2021, 78, 376-386.	1.3	2
24	Multiethnic Genome-Wide Association Study of Subclinical Atherosclerosis in Individuals With Type 2 Diabetes. Circulation Genomic and Precision Medicine, 2021, 14, e003258.	1.6	4
25	A Missense Variant in the IL-6 Receptor and Protection From Peripheral Artery Disease. Circulation Research, 2021, 129, 968-970.	2.0	11
26	Cholesteryl ester transfer protein (CETP) as a drug target for cardiovascular disease. Nature Communications, 2021, 12, 5640.	5.8	57
27	Genetic Loci Associated With COVID-19 Positivity and Hospitalization in White, Black, and Hispanic Veterans of the VA Million Veteran Program. Frontiers in Genetics, 2021, 12, 777076.	1.1	9
28	Multi-Trait Genome-Wide Association Study of Atherosclerosis Detects Novel Pleiotropic Loci. Frontiers in Genetics, 2021, 12, 787545.	1.1	3
29	Fried food consumption and risk of coronary artery disease: The Million Veteran Program. Clinical Nutrition, 2020, 39, 1203-1208.	2.3	15
30	Inherited myeloproliferative neoplasm risk affects haematopoietic stem cells. Nature, 2020, 586, 769-775.	13.7	101
31	Expressing Results From a Mendelian Randomization Analysis. JAMA Cardiology, 2020, 6, 7-8.	3.0	9
32	Discovery of rare variants associated with blood pressure regulation through meta-analysis of 1.3 million individuals. Nature Genetics, 2020, 52, 1314-1332.	9.4	91
33	Chromosome 1q21.2 and additional loci influence risk of spontaneous coronary artery dissection and myocardial infarction. Nature Communications, 2020, 11, 4432.	5.8	60
34	Genetic determinants of increased body mass index mediate the effect of smoking on increased risk for type 2 diabetes but not coronary artery disease. Human Molecular Genetics, 2020, 29, 3327-3337.	1.4	6
35	Validating a non-invasive, ALT-based non-alcoholic fatty liver phenotype in the million veteran program. PLoS ONE, 2020, 15, e0237430.	1.1	15
36	Radiomics of Coronary Artery Calcium in the Framingham Heart Study. Radiology: Cardiothoracic Imaging, 2020, 2, e190119.	0.9	22

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37	Rural-Urban Differences in Cardiovascular Mortality in the US, 1999-2017. JAMA - Journal of the American Medical Association, 2020, 323, 1852.	3.8	94
38	Prescription Fill Patterns for Commonly Used Drugs During the COVID-19 Pandemic in the United States. JAMA - Journal of the American Medical Association, 2020, 323, 2524.	3.8	121
39	Discovery of 318 new risk loci for type 2 diabetes and related vascular outcomes among 1.4 million participants in a multi-ancestry meta-analysis. Nature Genetics, 2020, 52, 680-691.	9.4	445
40	Minority-centric meta-analyses of blood lipid levels identify novel loci in the Population Architecture using Genomics and Epidemiology (PAGE) study. PLoS Genetics, 2020, 16, e1008684.	1.5	17
41	Opportunities, challenges and expectations management for translating biobank research to precision medicine. European Journal of Epidemiology, 2020, 35, 1-4.	2.5	15
42	Genotyping Array Design and Data Quality Control in the Million Veteran Program. American Journal of Human Genetics, 2020, 106, 535-548.	2.6	118
43	Opportunities and Challenges for Polygenic Risk Scores in Prognostication and Prevention of Cardiovascular Disease. JAMA Cardiology, 2020, 5, 399.	3.0	4
44	Genetic loci associated with prevalent and incident myocardial infarction and coronary heart disease in the Cohorts for Heart and Aging Research in Genomic Epidemiology (CHARGE) Consortium. PLoS ONE, 2020, 15, e0230035.	1.1	5
45	PCSK9 loss of function is protective against extra-coronary atherosclerotic cardiovascular disease in a large multi-ethnic cohort. PLoS ONE, 2020, 15, e0239752.	1.1	9
46	Title is missing!. , 2020, 16, e1008684.		0
47	Title is missing!. , 2020, 16, e1008684.		0
48	Title is missing!. , 2020, 16, e1008684.		0
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50	Title is missing!. , 2020, 16, e1008684.		0
51	Title is missing!. , 2020, 16, e1008684.		0
52	Association of APOL1 Risk Alleles With Cardiovascular Disease in Blacks in the Million Veteran Program. Circulation, 2019, 140, 1031-1040.	1.6	31
53	Genome-wide association study of peripheral artery disease in the Million Veteran Program. Nature Medicine, 2019, 25, 1274-1279.	15.2	177
54	High-throughput multimodal automated phenotyping (MAP) with application to PheWAS. Journal of the American Medical Informatics Association: JAMIA, 2019, 26, 1255-1262.	2.2	69

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55	HDAC9 is implicated in atherosclerotic aortic calcification and affects vascular smooth muscle cell phenotype. Nature Genetics, 2019, 51, 1580-1587.	9.4	92
56	Target genes, variants, tissues and transcriptional pathways influencing human serum urate levels. Nature Genetics, 2019, 51, 1459-1474.	9.4	251
57	Exome sequencing of 20,791Âcases of type 2 diabetes and 24,440Âcontrols. Nature, 2019, 570, 71-76.	13.7	248
58	Mendelian randomization evaluation of causal effects of fibrinogen on incident coronary heart disease. PLoS ONE, 2019, 14, e0216222.	1.1	17
59	Trans-ethnic association study of blood pressure determinants in over 750,000 individuals. Nature Genetics, 2019, 51, 51-62.	9.4	328
60	Genome-Wide Association Transethnic Meta-Analyses Identifies Novel Associations Regulating Coagulation Factor VIII and von Willebrand Factor Plasma Levels. Circulation, 2019, 139, 620-635.	1.6	102
61	Association of the PHACTR1/EDN1 Genetic Locus With Spontaneous Coronary Artery Dissection. Journal of the American College of Cardiology, 2019, 73, 58-66.	1.2	147
62	Alcohol Consumption and Risk of Coronary Artery Disease (from the Million Veteran Program). American Journal of Cardiology, 2018, 121, 1162-1168.	0.7	23
63	Maintenance of Ideal Cardiovascular Health and Coronary Artery Calcium Progression in Low-Risk Men and Women in the Framingham Heart Study. Circulation: Cardiovascular Imaging, 2018, 11, e006209.	1.3	28
64	Large-Scale Genomic Biobanks and Cardiovascular Disease. Current Cardiology Reports, 2018, 20, 22.	1.3	8
65	Observational and Genetic Associations of Resting Heart Rate With Aortic Valve Calcium. American Journal of Cardiology, 2018, 121, 1246-1252.	0.7	3
66	Longitudinal Associations of Pericardial and Intrathoracic Fat With Progression of Coronary Artery Calcium (from the Framingham Heart Study). American Journal of Cardiology, 2018, 121, 162-167.	0.7	10
67	Effects of Genetic Variants Associated with Familial Hypercholesterolemia on Low-Density Lipoprotein-Cholesterol Levels and Cardiovascular Outcomes in the Million Veteran Program. Circulation Genomic and Precision Medicine, 2018, 11, .	1.6	15
68	GWAS and colocalization analyses implicate carotid intima-media thickness and carotid plaque loci in cardiovascular outcomes. Nature Communications, 2018, 9, 5141.	5.8	119
69	Lp-PLA2, scavenger receptor class B type I gene (SCARB1) rs10846744 variant, and cardiovascular disease. PLoS ONE, 2018, 13, e0204352.	1.1	2
70	Genetics of blood lipids among ~300,000 multi-ethnic participants of the Million Veteran Program. Nature Genetics, 2018, 50, 1514-1523.	9.4	497
71	A phenotyping algorithm to identify acute ischemic stroke accurately from a national biobank: the Million Veteran Program. Clinical Epidemiology, 2018, Volume 10, 1509-1521.	1.5	20
72	Genetic analysis of over 1 million people identifies 535 new loci associated with blood pressure traits. Nature Genetics, 2018, 50, 1412-1425.	9.4	924

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73	Opportunities and Challenges in Mendelian Randomization Studies to Guide Trial Design. JAMA Cardiology, 2018, 3, 967.	3.0	19
74	Astronaut Cardiovascular Health and Risk Modification (Astro-CHARM) Coronary Calcium Atherosclerotic Cardiovascular Disease Risk Calculator. Circulation, 2018, 138, 1819-1827.	1.6	54
75	Is Heart Failure Inherited?. JAMA Cardiology, 2018, 3, 710.	3.0	2
76	Association of Interleukin 6 Receptor Variant With Cardiovascular Disease Effects of Interleukin 6 Receptor Blocking Therapy. JAMA Cardiology, 2018, 3, 849.	3.0	75
77	Mendelian Randomization Evidence for Cardiovascular Precision Medicine. JAMA Cardiology, 2018, 3, 627.	3.0	6
78	Baseline Characterization and Annual Trends of Body Mass Index for a Mega-Biobank Cohort of US Veterans 2011-2017. Journal of Health Research and Reviews, 2018, 5, 98-107.	0.1	3
79	Novel Thrombotic Function of a Human SNP in STXBP5 Revealed by CRISPR/Cas9 Gene Editing in Mice. Arteriosclerosis, Thrombosis, and Vascular Biology, 2017, 37, 264-270.	1.1	24
80	Serum Sortilin Associates With Aortic Calcification and Cardiovascular Risk in Men. Arteriosclerosis, Thrombosis, and Vascular Biology, 2017, 37, 1005-1011.	1.1	44
81	Relation of Risk Factors and Abdominal Aortic Calcium to Progression of Coronary Artery Calcium (from the Framingham Heart Study). American Journal of Cardiology, 2017, 119, 1584-1589.	0.7	21
82	Association of Multiorgan Computed Tomographic Phenomap With Adverse Cardiovascular Health Outcomes. JAMA Cardiology, 2017, 2, 1236.	3.0	19
83	Coronary Artery Calcium Distribution Is an Independent Predictor of Incident Major Coronary Heart Disease Events. Circulation: Cardiovascular Imaging, 2017, 10, .	1.3	78
84	Association of descending thoracic aortic plaque with brain atrophy and white matter hyperintensities: The Framingham Heart Study. Atherosclerosis, 2017, 265, 305-311.	0.4	13
85	Guideline-Based Statin Eligibility, Cancer Events, and Noncardiovascular Mortality in the Framingham Heart Study. Journal of Clinical Oncology, 2017, 35, 2927-2933.	0.8	22
86	Epidemiology of venous thromboembolism in the Framingham Heart Study. Thrombosis Research, 2016, 145, 27-33.	0.8	94
87	Genetic loci associated with ideal cardiovascular health: A meta-analysis of genome-wide association studies. American Heart Journal, 2016, 175, 112-120.	1.2	25
88	Cardiovascular Event Prediction and Risk Reclassification by Coronary, Aortic, and Valvular Calcification in the Framingham Heart Study. Journal of the American Heart Association, 2016, 5, .	1.6	150
89	Biomarkers for the prediction of venous thromboembolism in the community. Thrombosis Research, 2016, 145, 34-39.	0.8	14
90	Multiethnic Exome-Wide Association Study of Subclinical Atherosclerosis. Circulation: Cardiovascular Genetics, 2016, 9, 511-520.	5.1	54

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91	The genetics of blood pressure regulation and its target organs from association studies in 342,415 individuals. Nature Genetics, 2016, 48, 1171-1184.	9.4	362
92	Prevalence and Prognostic Implications of Coronary Artery Calcification in Low-Risk Women. JAMA - Journal of the American Medical Association, 2016, 316, 2126.	3.8	107
93	Rapid evaluation of phenotypes, SNPs and results through the dbGaP CHARGE Summary Results site. Nature Genetics, 2016, 48, 702-703.	9.4	13
94	Causal Assessment of Serum Urate Levels inÂCardiometabolic Diseases Through a Mendelian Randomization Study. Journal of the American College of Cardiology, 2016, 67, 407-416.	1.2	138
95	Reducing Cardiovascular Risk Using Genomic Information in the Era of Precision Medicine. Circulation, 2016, 133, 1155-1159.	1.6	9
96	Circulating Sex Steroids and Vascular Calcification in Community-Dwelling Men: The Framingham Heart Study. Journal of Clinical Endocrinology and Metabolism, 2016, 101, 2160-2167.	1.8	20
97	A meta-analysis of 120 246 individuals identifies 18 new loci for fibrinogen concentration. Human Molecular Genetics, 2016, 25, 358-370.	1.4	73
98	Phosphodiesterase 1 regulation is a key mechanism in vascular aging. Clinical Science, 2015, 129, 1061-1075.	1.8	53
99	Left Ventricular Structure and Risk of Cardiovascular Events: A Framingham Heart Study Cardiac Magnetic Resonance Study. Journal of the American Heart Association, 2015, 4, e002188.	1.6	109
100	Identification of common genetic variants controlling transcript isoform variation in human whole blood. Nature Genetics, 2015, 47, 345-352.	9.4	103
101	Low Cardiac Index Is Associated With Incident Dementia and Alzheimer Disease. Circulation, 2015, 131, 1333-1339.	1.6	140
102	Genetically Determined Height and Coronary Artery Disease. New England Journal of Medicine, 2015, 372, 1608-1618.	13.9	220
103	A comprehensive 1000 Genomesâ€"based genome-wide association meta-analysis of coronary artery disease. Nature Genetics, 2015, 47, 1121-1130.	9.4	2,054
104	Fetuin-A and risk of coronary heart disease: A Mendelian randomization analysis and a pooled analysis of AHSG genetic variants in 7 prospective studies. Atherosclerosis, 2015, 243, 44-52.	0.4	21
105	Risk Factor Differences in Calcified and Noncalcified Aortic Plaque. Arteriosclerosis, Thrombosis, and Vascular Biology, 2014, 34, 1580-1586.	1.1	13
106	Renal Artery Calcium, Cardiovascular Risk Factors, and Indexes of Renal Function. American Journal of Cardiology, 2014, 113, 156-161.	0.7	23
107	The Systolic Blood Pressure Difference Between Arms and Cardiovascular Disease in the Framingham Heart Study. American Journal of Medicine, 2014, 127, 209-215.	0.6	112
108	Association of Low-Frequency and Rare Coding-Sequence Variants with Blood Lipids and Coronary Heart Disease in 56,000 Whites and Blacks. American Journal of Human Genetics, 2014, 94, 223-232.	2.6	287

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109	Whole-Exome Sequencing Identifies Rare and Low-Frequency Coding Variants Associated with LDL Cholesterol. American Journal of Human Genetics, 2014, 94, 233-245.	2.6	193
110	Magnesium Intake Is Inversely Associated With Coronary Artery Calcification. JACC: Cardiovascular Imaging, 2014, 7, 59-69.	2.3	97
111	Genetic association study of QT interval highlights role for calcium signaling pathways in myocardial repolarization. Nature Genetics, 2014, 46, 826-836.	9.4	281
112	GRASP: analysis of genotypeâ€“phenotype results from 1390 genome-wide association studies and corresponding open access database. Bioinformatics, 2014, 30, i185-i194.	1.8	261
113	Distribution of Abdominal Aortic Calcium by Computed Tomography. Academic Radiology, 2013, 20, 1422-1428.	1.3	6
114	Assessing the phenotypic effects in the general population of rare variants in genes for a dominant Mendelian form of diabetes. Nature Genetics, 2013, 45, 1380-1385.	9.4	129
115	Common genetic loci influencing plasma homocysteine concentrations and their effect on risk of coronary artery disease. American Journal of Clinical Nutrition, 2013, 98, 668-676.	2.2	161
116	Genetic Associations with Valvular Calcification and Aortic Stenosis. New England Journal of Medicine, 2013, 368, 503-512.	13.9	767
117	Genomic Medicine for Improved Prediction and Primordial Prevention of Cardiovascular Disease. Arteriosclerosis, Thrombosis, and Vascular Biology, 2013, 33, 2049-2050.	1.1	9
118	Best Practices and Joint Calling of the HumanExome BeadChip: The CHARGE Consortium. PLoS ONE, 2013, 8, e68095.	1.1	219
119	Using Family-Based Imputation in Genome-Wide Association Studies with Large Complex Pedigrees: The Framingham Heart Study. PLoS ONE, 2012, 7, e51589.	1.1	17
120	Hypothesis-Based Analysis of Gene-Gene Interactions and Risk of Myocardial Infarction. PLoS ONE, 2012, 7, e41730.	1.1	17
121	Genome-wide association study identifies loci influencing concentrations of liver enzymes in plasma. Nature Genetics, 2011, 43, 1131-1138.	9.4	501
122	Genome-Wide Association Analysis Identifies Variants Associated with Nonalcoholic Fatty Liver Disease That Have Distinct Effects on Metabolic Traits. PLoS Genetics, 2011, 7, e1001324.	1.5	796
123	Strengthening the reporting of genetic risk prediction studies (GRIPS): explanation and elaboration. European Journal of Clinical Investigation, 2011, 41, 1010-1035.	1.7	30
124	Genome-Wide Association Study for Coronary Artery Calcification With Follow-Up in Myocardial Infarction. Circulation, 2011, 124, 2855-2864.	1.6	269
125	Reply:. Hepatology, 2010, 52, 1519-1519.	3.6	0
126	Hundreds of variants clustered in genomic loci and biological pathways affect human height. Nature, 2010, 467, 832-838.	13.7	1,789

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127	Whole- and refined-grain intakes are differentially associated with abdominal visceral and subcutaneous adiposity in healthy adults: the Framingham Heart Study. American Journal of Clinical Nutrition, 2010, 92, 1165-1171.	2.2	119
128	Vitamin K supplementation and progression of coronary artery calcium in older men and women. American Journal of Clinical Nutrition, 2009, 89, 1799-1807.	2.2	212
129	Cohorts for Heart and Aging Research in Genomic Epidemiology (CHARGE) Consortium. Circulation: Cardiovascular Genetics, 2009, 2, 73-80.	5.1	519
130	Defining Normal Distributions of Coronary Artery Calcium in Women and Men (from the Framingham) Tj ETQq0 0 0 rgBT /Overlock 10 T	0.7	150
131	Pericardial Fat, Visceral Abdominal Fat, Cardiovascular Disease Risk Factors, and Vascular Calcification in a Community-Based Sample. Circulation, 2008, 117, 605-613.	1.6	896
132	Visceral and Subcutaneous Adipose Tissue Volumes Are Cross-Sectionally Related to Markers of Inflammation and Oxidative Stress. Circulation, 2007, 116, 1234-1241.	1.6	779
133	The Third Generation Cohort of the National Heart, Lung, and Blood Institute's Framingham Heart Study: Design, Recruitment, and Initial Examination. American Journal of Epidemiology, 2007, 165, 1328-1335.	1.6	752
134	Clinical and Genetic Correlates of Aldosterone-to-Renin Ratio and Relations to Blood Pressure in a Community Sample. Hypertension, 2007, 49, 846-856.	1.3	187
135	Abstract 291: Association of Single Nucleotide Polymorphisms in Inflammatory Candidate Genes with Circulating Inflammatory Biomarker Concentrations: The Framingham Heart Study. Circulation, 2007, 116, .	1.6	0
136	Plasma vitamin K levels are associated with coronary calcification in older adults.. FASEB Journal, 2006, 20, A134.	0.2	1
137	Calcium Concentration of Individual Coronary Calcified Plaques as Measured by Multidetector Row Computed Tomography. Circulation, 2005, 111, 3236-3241.	1.6	53
138	Usefulness of Exercise Testing in the Prediction of Coronary Disease Risk Among Asymptomatic Persons as a Function of the Framingham Risk Score. Circulation, 2004, 110, 1920-1925.	1.6	157
139	Mitral Annular Calcification Predicts Cardiovascular Morbidity and Mortality. Circulation, 2003, 107, 1492-1496.	1.6	397
140	Association of C-Reactive Protein With Carotid Atherosclerosis in Men and Women: The Framingham Heart Study. Arteriosclerosis, Thrombosis, and Vascular Biology, 2002, 22, 1662-1667.	1.1	217
141	Abdominal Aortic Calcific Deposits Are an Important Predictor of Vascular Morbidity and Mortality. Circulation, 2001, 103, 1529-1534.	1.6	546
142	Factor VII Gene Polymorphism, Factor VII Levels, and Prevalent Cardiovascular Disease. Arteriosclerosis, Thrombosis, and Vascular Biology, 2000, 20, 593-600.	1.1	55
143	Differential Control of Systolic and Diastolic Blood Pressure. Hypertension, 2000, 36, 594-599.	1.3	378
144	Increased Platelet Aggregability Associated With Platelet GPIIIa Pl ^{A2} Polymorphism . Arteriosclerosis, Thrombosis, and Vascular Biology, 1999, 19, 1142-1147.	1.1	241