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	A comprehensive 1000 Genomes–based genome-wide association meta-analysis of coronary artery disease. Nature Genetics, 2015, 47, 1121-1130.	9.4	2,054
2	Hundreds of variants clustered in genomic loci and biological pathways affect human height. Nature, 2010, 467, 832-838.	13.7	1,789
3	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
4	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
5	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
6	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
7	Genetic Associations with Valvular Calcification and Aortic Stenosis. New England Journal of Medicine, 2013, 368, 503-512.	13.9	767
8	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
9	Abdominal Aortic Calcific Deposits Are an Important Predictor of Vascular Morbidity and Mortality. Circulation, 2001, 103, 1529-1534.	1.6	546
10	Cohorts for Heart and Aging Research in Genomic Epidemiology (CHARGE) Consortium. Circulation: Cardiovascular Genetics, 2009, 2, 73-80.	5.1	519
11	Genome-wide association study identifies loci influencing concentrations of liver enzymes in plasma. Nature Genetics, 2011, 43, 1131-1138.	9.4	501
12	Genetics of blood lipids among ~300,000 multi-ethnic participants of the Million Veteran Program. Nature Genetics, 2018, 50, 1514-1523.	9.4	497
13	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
14	Mitral Annular Calcification Predicts Cardiovascular Morbidity and Mortality. Circulation, 2003, 107, 1492-1496.	1.6	397
15	Differential Control of Systolic and Diastolic Blood Pressure. Hypertension, 2000, 36, 594-599.	1.3	378
16	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
17	Trans-ethnic association study of blood pressure determinants in over 750,000 individuals. Nature Genetics, 2019, 51, 51-62.	9.4	328
18	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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19	Genetic association study of QT interval highlights role for calcium signaling pathways in myocardial repolarization. Nature Genetics, 2014, 46, 826-836.	9.4	281
20	Genome-Wide Association Study for Coronary Artery Calcification With Follow-Up in Myocardial Infarction. Circulation, 2011, 124, 2855-2864.	1.6	269
21	GRASP: analysis of genotype–phenotype results from 1390 genome-wide association studies and corresponding open access database. Bioinformatics, 2014, 30, i185-i194.	1.8	261
22	Target genes, variants, tissues and transcriptional pathways influencing human serum urate levels. Nature Genetics, 2019, 51, 1459-1474.	9.4	251
23	Exome sequencing of 20,791 cases of type 2 diabetes and 24,440 controls. Nature, 2019, 570, 71-76.	13.7	248
24	Increased Platelet Aggregability Associated With Platelet <i> GPIIIa Pl ^{<i>A2</i>} Polymorphism </i>. Arteriosclerosis, Thrombosis, and Vascular Biology, 1999, 19, 1142-1147.	1.1	241
25	Genetically Determined Height and Coronary Artery Disease. New England Journal of Medicine, 2015, 372, 1608-1618.	13.9	220
26	Best Practices and Joint Calling of the HumanExome BeadChip: The CHARGE Consortium. PLoS ONE, 2013, 8, e68095.	1.1	219
27	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
28	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
29	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
30	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
31	Genome-wide association study of peripheral artery disease in the Million Veteran Program. Nature Medicine, 2019, 25, 1274-1279.	15.2	177
32	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
33	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
34	Defining Normal Distributions of Coronary Artery Calcium in Women and Men (from the Framingham) Tj ETQq0 0 0 rgBT /Overlock 10 Tf	0.7	150
35	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
36	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

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37	Low Cardiac Index Is Associated With Incident Dementia and Alzheimer Disease. Circulation, 2015, 131, 1333-1339.	1.6	140
38	Causal Assessment of Serum Urate Levels in Cardiometabolic Diseases Through a Mendelian Randomization Study. Journal of the American College of Cardiology, 2016, 67, 407-416.	1.2	138
39	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
40	Discovery and prioritization of variants and genes for kidney function in >1.2 million individuals. Nature Communications, 2021, 12, 4350.	5.8	125
41	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
42	Actionable druggable genome-wide Mendelian randomization identifies repurposing opportunities for COVID-19. Nature Medicine, 2021, 27, 668-676.	15.2	120
43	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
44	GWAS and colocalization analyses implicate carotid intima-media thickness and carotid plaque loci in cardiovascular outcomes. Nature Communications, 2018, 9, 5141.	5.8	119
45	Genotyping Array Design and Data Quality Control in the Million Veteran Program. American Journal of Human Genetics, 2020, 106, 535-548.	2.6	118
46	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
47	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
48	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
49	Identification of common genetic variants controlling transcript isoform variation in human whole blood. Nature Genetics, 2015, 47, 345-352.	9.4	103
50	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
51	Inherited myeloproliferative neoplasm risk affects haematopoietic stem cells. Nature, 2020, 586, 769-775.	13.7	101
52	Magnesium Intake Is Inversely Associated With Coronary Artery Calcification. JACC: Cardiovascular Imaging, 2014, 7, 59-69.	2.3	97
53	Epidemiology of venous thromboembolism in the Framingham Heart Study. Thrombosis Research, 2016, 145, 27-33.	0.8	94
54	Rural-Urban Differences in Cardiovascular Mortality in the US, 1999-2017. JAMA - Journal of the American Medical Association, 2020, 323, 1852.	3.8	94

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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	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
57	Coronary Artery Calcium Distribution Is an Independent Predictor of Incident Major Coronary Heart Disease Events. Circulation: Cardiovascular Imaging, 2017, 10, .	1.3	78
58	Association of Interleukin 6 Receptor Variant With Cardiovascular Disease Effects of Interleukin 6 Receptor Blocking Therapy. JAMA Cardiology, 2018, 3, 849.	3.0	75
59	A meta-analysis of 120 246 individuals identifies 18 new loci for fibrinogen concentration. Human Molecular Genetics, 2016, 25, 358-370.	1.4	73
60	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
61	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
62	Chromosome 1q21.2 and additional loci influence risk of spontaneous coronary artery dissection and myocardial infarction. Nature Communications, 2020, 11, 4432.	5.8	60
63	Cholesteryl ester transfer protein (CETP) as a drug target for cardiovascular disease. Nature Communications, 2021, 12, 5640.	5.8	57
64	Factor VII Gene Polymorphism, Factor VII Levels, and Prevalent Cardiovascular Disease. Arteriosclerosis, Thrombosis, and Vascular Biology, 2000, 20, 593-600.	1.1	55
65	Multiethnic Exome-Wide Association Study of Subclinical Atherosclerosis. Circulation: Cardiovascular Genetics, 2016, 9, 511-520.	5.1	54
66	Astronaut Cardiovascular Health and Risk Modification (Astro-CHARM) Coronary Calcium Atherosclerotic Cardiovascular Disease Risk Calculator. Circulation, 2018, 138, 1819-1827.	1.6	54
67	Calcium Concentration of Individual Coronary Calcified Plaques as Measured by Multidetector Row Computed Tomography. Circulation, 2005, 111, 3236-3241.	1.6	53
68	Phosphodiesterase 1 regulation is a key mechanism in vascular aging. Clinical Science, 2015, 129, 1061-1075.	1.8	53
69	Genetic analysis in European ancestry individuals identifies 517 loci associated with liver enzymes. Nature Communications, 2021, 12, 2579.	5.8	51
70	Determinants of penetrance and variable expressivity in monogenic metabolic conditions across 77,184 exomes. Nature Communications, 2021, 12, 3505.	5.8	49
71	Serum Sortilin Associates With Aortic Calcification and Cardiovascular Risk in Men. Arteriosclerosis, Thrombosis, and Vascular Biology, 2017, 37, 1005-1011.	1.1	44
72	Association of *APOL1* Risk Alleles With Cardiovascular Disease in Blacks in the Million Veteran Program. Circulation, 2019, 140, 1031-1040.	1.6	31

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73	*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
74	Strengthening the reporting of genetic risk prediction studies (GRIPS): explanation and elaboration. European Journal of Clinical Investigation, 2011, 41, 1010-1035.	1.7	30
75	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
76	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
77	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
78	Renal Artery Calcium, Cardiovascular Risk Factors, and Indexes of Renal Function. American Journal of Cardiology, 2014, 113, 156-161.	0.7	23
79	Alcohol Consumption and Risk of Coronary Artery Disease (from the Million Veteran Program). American Journal of Cardiology, 2018, 121, 1162-1168.	0.7	23
80	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
81	Radiomics of Coronary Artery Calcium in the Framingham Heart Study. Radiology: Cardiothoracic Imaging, 2020, 2, e190119.	0.9	22
82	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
83	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
84	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
85	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
86	Association of Multiorgan Computed Tomographic Phenomap With Adverse Cardiovascular Health Outcomes. JAMA Cardiology, 2017, 2, 1236.	3.0	19
87	Opportunities and Challenges in Mendelian Randomization Studies to Guide Trial Design. JAMA Cardiology, 2018, 3, 967.	3.0	19
88	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
89	Plasma Protein Profile of Carotid Artery Atherosclerosis and Atherosclerotic Outcomes. Arteriosclerosis, Thrombosis, and Vascular Biology, 2021, 41, 1777-1788.	1.1	18
90	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

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91	Mendelian randomization evaluation of causal effects of fibrinogen on incident coronary heart disease. PLoS ONE, 2019, 14, e0216222.	1.1	17
92	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
93	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
94	Hypothesis-Based Analysis of Gene-Gene Interactions and Risk of Myocardial Infarction. PLoS ONE, 2012, 7, e41730.	1.1	17
95	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
96	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
97	Fried food consumption and risk of coronary artery disease: The Million Veteran Program. Clinical Nutrition, 2020, 39, 1203-1208.	2.3	15
98	Validating a non-invasive, ALT-based non-alcoholic fatty liver phenotype in the million veteran program. PLoS ONE, 2020, 15, e0237430.	1.1	15
99	Opportunities, challenges and expectations management for translating biobank research to precision medicine. European Journal of Epidemiology, 2020, 35, 1-4.	2.5	15
100	Biomarkers for the prediction of venous thromboembolism in the community. Thrombosis Research, 2016, 145, 34-39.	0.8	14
101	Risk Factor Differences in Calcified and Noncalcified Aortic Plaque. Arteriosclerosis, Thrombosis, and Vascular Biology, 2014, 34, 1580-1586.	1.1	13
102	Rapid evaluation of phenotypes, SNPs and results through the dbGaP CHARGE Summary Results site. Nature Genetics, 2016, 48, 702-703.	9.4	13
103	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
104	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
105	A multi-population phenome-wide association study of genetically-predicted height in the Million Veteran Program. PLoS Genetics, 2022, 18, e1010193.	1.5	12
106	Comparison of family health history in surveys vs electronic health record data mapped to the observational medical outcomes partnership data model in the <i>All of Us</i> Research Program. Journal of the American Medical Informatics Association: JAMIA, 2021, 28, 695-703.	2.2	11
107	A Missense Variant in the IL-6 Receptor and Protection From Peripheral Artery Disease. Circulation Research, 2021, 129, 968-970.	2.0	11
108	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

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109	Meta-analysis of epigenome-wide association studies of carotid intima-media thickness. European Journal of Epidemiology, 2021, 36, 1143-1155.	2.5	10
110	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
111	Genomic Medicine for Improved Prediction and Primordial Prevention of Cardiovascular Disease. Arteriosclerosis, Thrombosis, and Vascular Biology, 2013, 33, 2049-2050.	1.1	9
112	Reducing Cardiovascular Risk Using Genomic Information in the Era of Precision Medicine. Circulation, 2016, 133, 1155-1159.	1.6	9
113	Expressing Results From a Mendelian Randomization Analysis. JAMA Cardiology, 2020, 6, 7-8.	3.0	9
114	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
115	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
116	Large-Scale Genomic Biobanks and Cardiovascular Disease. Current Cardiology Reports, 2018, 20, 22.	1.3	8
117	Integration of rare expression outlier-associated variants improves polygenic risk prediction. American Journal of Human Genetics, 2022, 109, 1055-1064.	2.6	8
118	Distribution of Abdominal Aortic Calcium by Computed Tomography. Academic Radiology, 2013, 20, 1422-1428.	1.3	6
119	Mendelian Randomization Evidence for Cardiovascular Precision Medicine. JAMA Cardiology, 2018, 3, 627.	3.0	6
120	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
121	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
122	Genome-wide transcriptome study using deep RNA sequencing for myocardial infarction and coronary artery calcification. BMC Medical Genomics, 2021, 14, 45.	0.7	5
123	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
124	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
125	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
126	Opportunities and Challenges for Polygenic Risk Scores in Prognostication and Prevention of Cardiovascular Disease. JAMA Cardiology, 2020, 5, 399.	3.0	4

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127	Observational and Genetic Associations of Resting Heart Rate With Aortic Valve Calcium. American Journal of Cardiology, 2018, 121, 1246-1252.	0.7	3
128	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
129	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
130	Multi-Trait Genome-Wide Association Study of Atherosclerosis Detects Novel Pleiotropic Loci. Frontiers in Genetics, 2021, 12, 787545.	1.1	3
131	Lp-PLA2, scavenger receptor class B type I gene (SCARB1) rs10846744 variant, and cardiovascular disease. PLoS ONE, 2018, 13, e0204352.	1.1	2
132	Is Heart Failure Inherited?. JAMA Cardiology, 2018, 3, 710.	3.0	2
133	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
134	Genome-wide and phenome-wide analysis of ideal cardiovascular health in the VA Million Veteran Program. PLoS ONE, 2022, 17, e0267900.	1.1	2
135	Genetic Contribution to Common Heart Failureâ€"Not So Rare?. JAMA Cardiology, 2021, 6, 387.	3.0	1
136	Plasma vitamin K levels are associated with coronary calcification in older adults.. FASEB Journal, 2006, 20, A134.	0.2	1
137	Reply:. Hepatology, 2010, 52, 1519-1519.	3.6	0
138	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
139	Title is missing!. , 2020, 16, e1008684.		0
140	Title is missing!. , 2020, 16, e1008684.		0
141	Title is missing!. , 2020, 16, e1008684.		0
142	Title is missing!. , 2020, 16, e1008684.		0
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144	Title is missing!. , 2020, 16, e1008684.		0