Christopher J O donnell

List of Publications by Citations

139 papers	14,256 citations	50 h-index	119 g-index
153 ext. papers	18,098 ext. citations	14 avg, IF	5.57 L-index

#	Paper	IF	Citations
139	Hundreds of variants clustered in genomic loci and biological pathways affect human height. *Nature*, **2010**, 467, 832-8	50.4	1514
138	A comprehensive 1,000 Genomes-based genome-wide association meta-analysis of coronary artery disease. *Nature Genetics*, **2015**, 47, 1121-1130	36.3	1290
137	Pericardial fat, visceral abdominal fat, cardiovascular disease risk factors, and vascular calcification in a community-based sample: the Framingham Heart Study. *Circulation*, **2008**, 117, 605-13	16.7	747
136	Visceral and subcutaneous adipose tissue volumes are cross-sectionally related to markers of inflammation and oxidative stress: the Framingham Heart Study. *Circulation*, **2007**, 116, 1234-41	16.7	665
135	Genome-wide association analysis identifies variants associated with nonalcoholic fatty liver disease that have distinct effects on metabolic traits. *PLoS Genetics*, **2011**, 7, e1001324	6	629
134	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, 1328-35	3.8	605
133	Genetic associations with valvular calcification and aortic stenosis. *New England Journal of Medicine* , **2013**, 368, 503-12	59.2	556
132	Abdominal aortic calcific deposits are an important predictor of vascular morbidity and mortality. *Circulation*, **2001**, 103, 1529-34	16.7	465
131	Cohorts for Heart and Aging Research in Genomic Epidemiology (CHARGE) Consortium: Design of prospective meta-analyses of genome-wide association studies from 5 cohorts. *Circulation: Cardiovascular Genetics*, **2009**, 2, 73-80		423
130	Genome-wide association study identifies loci influencing concentrations of liver enzymes in plasma. *Nature Genetics*, **2011**, 43, 1131-8	36.3	415
129	Genetic analysis of over 1 million people identifies 535 new loci associated with blood pressure traits. *Nature Genetics*, **2018**, 50, 1412-1425	36.3	386
128	Differential control of systolic and diastolic blood pressure : factors associated with lack of blood pressure control in the community. *Hypertension*, **2000**, 36, 594-9	8.5	329
127	Mitral annular calcification predicts cardiovascular morbidity and mortality: the Framingham Heart Study. *Circulation*, **2003**, 107, 1492-6	16.7	311
126	Genetics of blood lipids among ~300,000 multi-ethnic participants of the Million Veteran Program. *Nature Genetics*, **2018**, 50, 1514-1523	36.3	260
125	The genetics of blood pressure regulation and its target organs from association studies in 342,415 individuals. *Nature Genetics*, **2016**, 48, 1171-1184	36.3	251
124	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, 223-32	11	233
123	Increased platelet aggregability associated with platelet GPIIIa PlA2 polymorphism: the Framingham Offspring Study. *Arteriosclerosis, Thrombosis, and Vascular Biology*, **1999**, 19, 1142-7	9.4	217

122	Genome-wide association study for coronary artery calcification with follow-up in myocardial infarction. *Circulation*, **2011**, 124, 2855-64	16.7	213
121	Best practices and joint calling of the HumanExome BeadChip: the CHARGE Consortium. *PLoS ONE*, **2013**, 8, e68095	3.7	203
120	Genetic association study of QT interval highlights role for calcium signaling pathways in myocardial repolarization. *Nature Genetics*, **2014**, 46, 826-36	36.3	199
119	Association of C-reactive protein with carotid atherosclerosis in men and women: the Framingham Heart Study. *Arteriosclerosis, Thrombosis, and Vascular Biology*, **2002**, 22, 1662-7	9.4	193
118	GRASP: analysis of genotype-phenotype results from 1390 genome-wide association studies and corresponding open access database. *Bioinformatics*, **2014**, 30, i185-94	7.2	181
117	Vitamin K supplementation and progression of coronary artery calcium in older men and women. *American Journal of Clinical Nutrition*, **2009**, 89, 1799-807	7	171
116	Whole-exome sequencing identifies rare and low-frequency coding variants associated with LDL cholesterol. *American Journal of Human Genetics*, **2014**, 94, 233-45	11	170
115	Clinical and genetic correlates of aldosterone-to-renin ratio and relations to blood pressure in a community sample. *Hypertension*, **2007**, 49, 846-56	8.5	163
114	Genetically determined height and coronary artery disease. *New England Journal of Medicine*, **2015**, 372, 1608-18	59.2	152
113	Trans-ethnic association study of blood pressure determinants in over 750,000 individuals. *Nature Genetics*, **2019**, 51, 51-62	36.3	152
112	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, 1920-5	16.7	141
111	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	36.3	140
110	Exome sequencing of 20,791 cases of type 2 diabetes and 24,440 controls. *Nature*, **2019**, 570, 71-76	50.4	129
109	Defining normal distributions of coronary artery calcium in women and men (from the Framingham Heart Study). *American Journal of Cardiology*, **2008**, 102, 1136-41, 1141.e1	3	124
108	Target genes, variants, tissues and transcriptional pathways influencing human serum urate levels. *Nature Genetics*, **2019**, 51, 1459-1474	36.3	122
107	Common genetic loci influencing plasma homocysteine concentrations and their effect on risk of coronary artery disease. *American Journal of Clinical Nutrition*, **2013**, 98, 668-76	7	122
106	Assessing the phenotypic effects in the general population of rare variants in genes for a dominant Mendelian form of diabetes. *Nature Genetics*, **2013**, 45, 1380-5	36.3	103
105	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, 1165-71	7	102

104	Low cardiac index is associated with incident dementia and Alzheimer disease: the Framingham Heart Study. *Circulation*, **2015**, 131, 1333-9	16.7	101
103	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	15.1	101
102	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,	6	101
101	The systolic blood pressure difference between arms and cardiovascular disease in the Framingham Heart Study. *American Journal of Medicine*, **2014**, 127, 209-15	2.4	87
100	Association of the PHACTR1/EDN1 Genetic Locus With Spontaneous Coronary Artery Dissection. *Journal of the American College of Cardiology*, **2019**, 73, 58-66	15.1	86
99	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, 2524-2526	27.4	81
98	Magnesium intake is inversely associated with coronary artery calcification: the Framingham Heart Study. *JACC: Cardiovascular Imaging*, **2014**, 7, 59-69	8.4	80
97	Prevalence and Prognostic Implications of Coronary Artery Calcification in Low-Risk Women: A Meta-analysis. *JAMA - Journal of the American Medical Association*, **2016**, 316, 2126-2134	27.4	79
96	Identification of common genetic variants controlling transcript isoform variation in human whole blood. *Nature Genetics*, **2015**, 47, 345-52	36.3	77
95	Genome-wide association study of peripheral artery disease in the Million Veteran Program. *Nature Medicine*, **2019**, 25, 1274-1279	50.5	73
94	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	6	64
93	Epidemiology of venous thromboembolism in the Framingham Heart Study. *Thrombosis Research*, **2016**, 145, 27-33	8.2	64
92	GWAS and colocalization analyses implicate carotid intima-media thickness and carotid plaque loci in cardiovascular outcomes. *Nature Communications*, **2018**, 9, 5141	17.4	64
91	A meta-analysis of 120 246 individuals identifies 18 new loci for fibrinogen concentration. *Human Molecular Genetics*, **2016**, 25, 358-70	5.6	54
90	Genome-Wide Association Transethnic Meta-Analyses Identifies Novel Associations Regulating Coagulation Factor VIII and von Willebrand Factor Plasma Levels. *Circulation*, **2019**, 139, 620-635	16.7	51
89	Association of Interleukin 6 Receptor Variant With Cardiovascular Disease Effects of Interleukin 6 Receptor Blocking Therapy: A Phenome-Wide Association Study. *JAMA Cardiology*, **2018**, 3, 849-857	16.2	48
88	Factor VII gene polymorphism, factor VII levels, and prevalent cardiovascular disease: the Framingham Heart Study. *Arteriosclerosis, Thrombosis, and Vascular Biology*, **2000**, 20, 593-600	9.4	48
87	HDAC9 is implicated in atherosclerotic aortic calcification and affects vascular smooth muscle cell phenotype. *Nature Genetics*, **2019**, 51, 1580-1587	36.3	45

86	Calcium concentration of individual coronary calcified plaques as measured by multidetector row computed tomography. *Circulation*, **2005**, 111, 3236-41	16.7	45
85	Rural-Urban Differences in Cardiovascular Mortality in the US, 1999-2017. *JAMA - Journal of the American Medical Association*, **2020**, 323, 1852-1854	27.4	39
84	Phosphodiesterase 1 regulation is a key mechanism in vascular aging. *Clinical Science*, **2015**, 129, 1061-75	6.5	39
83	Coronary Artery Calcium Distribution Is an Independent Predictor of Incident Major Coronary Heart Disease Events: Results From the Framingham Heart Study. *Circulation: Cardiovascular Imaging*, **2017**, 10,	3.9	38
82	Multiethnic Exome-Wide Association Study of Subclinical Atherosclerosis. *Circulation: Cardiovascular Genetics*, **2016**, 9, 511-520		34
81	Serum Sortilin Associates With Aortic Calcification and Cardiovascular Risk in Men. *Arteriosclerosis, Thrombosis, and Vascular Biology*, **2017**, 37, 1005-1011	9.4	33
80	Inherited myeloproliferative neoplasm risk affects haematopoietic stem cells. *Nature*, **2020**, 586, 769-775	50.4	32
79	Astronaut Cardiovascular Health and Risk Modification (Astro-CHARM) Coronary Calcium Atherosclerotic Cardiovascular Disease Risk Calculator. *Circulation*, **2018**, 138, 1819-1827	16.7	30
78	Discovery of rare variants associated with blood pressure regulation through meta-analysis of 1.3 million individuals. *Nature Genetics*, **2020**, 52, 1314-1332	36.3	26
77	High-throughput multimodal automated phenotyping (MAP) with application to PheWAS. *Journal of the American Medical Informatics Association: JAMIA*, **2019**, 26, 1255-1262	8.6	23
76	Chromosome 1q21.2 and additional loci influence risk of spontaneous coronary artery dissection and myocardial infarction. *Nature Communications*, **2020**, 11, 4432	17.4	22
75	Genotyping Array Design and Data Quality Control in the Million Veteran Program. *American Journal of Human Genetics*, **2020**, 106, 535-548	11	22
74	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	9.4	19
73	Strengthening the reporting of genetic risk prediction studies (GRIPS): explanation and elaboration. *European Journal of Clinical Investigation*, **2011**, 41, 1010-35	4.6	19
72	Actionable druggable genome-wide Mendelian randomization identifies repurposing opportunities for COVID-19. *Nature Medicine*, **2021**, 27, 668-676	50.5	19
71	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	3.9	18
70	Association of Risk Alleles With Cardiovascular Disease in Blacks in the Million Veteran Program. *Circulation*, **2019**, 140, 1031-1040	16.7	18
69	Renal artery calcium, cardiovascular risk factors, and indexes of renal function. *American Journal of Cardiology*, **2014**, 113, 156-61	3	17

68	Genetic loci associated with ideal cardiovascular health: A meta-analysis of genome-wide association studies. *American Heart Journal*, **2016**, 175, 112-20	4.9	17
67	Guideline-Based Statin Eligibility, Cancer Events, and Noncardiovascular Mortality in the Framingham Heart Study. *Journal of Clinical Oncology*, **2017**, 35, 2927-2933	2.2	16
66	Association of Multiorgan Computed Tomographic Phenomap With Adverse Cardiovascular Health Outcomes: The Framingham Heart Study. *JAMA Cardiology*, **2017**, 2, 1236-1246	16.2	15
65	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	3.1	15
64	Hypothesis-based analysis of gene-gene interactions and risk of myocardial infarction. *PLoS ONE*, **2012**, 7, e41730	3.7	15
63	Circulating Sex Steroids and Vascular Calcification in Community-Dwelling Men: The Framingham Heart Study. *Journal of Clinical Endocrinology and Metabolism*, **2016**, 101, 2160-7	5.6	14
62	A phenotyping algorithm to identify acute ischemic stroke accurately from a national biobank: the Million Veteran Program. *Clinical Epidemiology*, **2018**, 10, 1509-1521	5.9	14
61	Alcohol Consumption and Risk of Coronary Artery Disease (from the Million Veteran Program). *American Journal of Cardiology*, **2018**, 121, 1162-1168	3	13
60	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	3	11
59	Mendelian randomization evaluation of causal effects of fibrinogen on incident coronary heart disease. *PLoS ONE*, **2019**, 14, e0216222	3.7	11
58	Risk factor differences in calcified and noncalcified aortic plaque: the Framingham Heart Study. *Arteriosclerosis, Thrombosis, and Vascular Biology*, **2014**, 34, 1580-6	9.4	11
57	Discovery and prioritization of variants and genes for kidney function in >1.2 million individuals. *Nature Communications*, **2021**, 12, 4350	17.4	11
56	Radiomics of Coronary Artery Calcium in the Framingham Heart Study. *Radiology: Cardiothoracic Imaging*, **2020**, 2, e190119	8.3	10
55	Rapid evaluation of phenotypes, SNPs and results through the dbGaP CHARGE Summary Results site. *Nature Genetics*, **2016**, 48, 702-3	36.3	10
54	Using family-based imputation in genome-wide association studies with large complex pedigrees: the Framingham Heart Study. *PLoS ONE*, **2012**, 7, e51589	3.7	10
53	Opportunities, challenges and expectations management for translating biobank research to precision medicine. *European Journal of Epidemiology*, **2020**, 35, 1-4	12.1	9
52	Association of descending thoracic aortic plaque with brain atrophy and white matter hyperintensities: The Framingham Heart Study. *Atherosclerosis*, **2017**, 265, 305-311	3.1	8
51	Biomarkers for the prediction of venous thromboembolism in the community. *Thrombosis Research*, **2016**, 145, 34-9	8.2	8

50	Large-Scale Genomic Biobanks and Cardiovascular Disease. *Current Cardiology Reports*, **2018**, 20, 22	4.2	7
49	Cholesteryl ester transfer protein (CETP) as a drug target for cardiovascular disease. *Nature Communications*, **2021**, 12, 5640	17.4	7
48	Genetic analysis in European ancestry individuals identifies 517 loci associated with liver enzymes. *Nature Communications*, **2021**, 12, 2579	17.4	7
47	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	3	7
46	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,	5.2	7
45	Expressing Results From a Mendelian Randomization Analysis: Separating Results From Inferences. *JAMA Cardiology*, **2021**, 6, 7-8	16.2	6
44	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	6	5
43	Distribution of abdominal aortic calcium by computed tomography: impact of analysis method on quantitative calcium score. *Academic Radiology*, **2013**, 20, 1422-8	4.3	5
42	Validating a non-invasive, ALT-based non-alcoholic fatty liver phenotype in the million veteran program. *PLoS ONE*, **2020**, 15, e0237430	3.7	5
41	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	5.8	5
40	Determinants of penetrance and variable expressivity in monogenic metabolic conditions across 77,184 exomes. *Nature Communications*, **2021**, 12, 3505	17.4	5
39	Fried food consumption and risk of coronary artery disease: The Million Veteran Program. *Clinical Nutrition*, **2020**, 39, 1203-1208	5.9	5
38	A trans-ancestry genome-wide association study of unexplained chronic ALT elevation as a proxy for nonalcoholic fatty liver disease with histological and radiological validation		5
37	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	3.7	4
36	Phenome-wide association of 1809 phenotypes and COVID-19 disease progression in the Veterans Health Administration Million Veteran Program. *PLoS ONE*, **2021**, 16, e0251651	3.7	4
35	Plasma Protein Profile of Carotid Artery Atherosclerosis and Atherosclerotic Outcomes: Meta-Analyses and Mendelian Randomization Analyses. *Arteriosclerosis, Thrombosis, and Vascular Biology*, **2021**, 41, 1777-1788	9.4	4
34	Meta-analysis of epigenome-wide association studies of carotid intima-media thickness. *European Journal of Epidemiology*, **2021**, 36, 1143-1155	12.1	4
33	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	0.2	3

32	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	5.6	3
31	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	8.6	3
30	Observational and Genetic Associations of Resting Heart Rate With Aortic Valve Calcium. *American Journal of Cardiology*, **2018**, 121, 1246-1252	3	2
29	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**,	11.5	2
28	Coronary Artery Disease Risk of Familial Hypercholesterolemia Genetic Variants Independent of Clinically Observed Longitudinal Cholesterol Exposure.. *Circulation Genomic and Precision Medicine*, **2022**, CIRCGEN121003501	5.2	2
27	PCSK9 loss of function is protective against extra-coronary atherosclerotic cardiovascular disease in a large multi-ethnic cohort. *PLoS ONE*, **2020**, 15, e0239752	3.7	2
26	Trends in cardiovascular procedural volumes in the setting of COVID-19: Insights from the VA clinical assessment, reporting, and tracking program. *Catheterization and Cardiovascular Interventions*, **2021**, 98, E326-E328	2.7	2
25	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	5.6	2
24	Lp-PLA2, scavenger receptor class B type I gene (SCARB1) rs10846744 variant, and cardiovascular disease. *PLoS ONE*, **2018**, 13, e0204352	3.7	2
23	Opportunities and Challenges for Polygenic Risk Scores in Prognostication and Prevention of Cardiovascular Disease. *JAMA Cardiology*, **2020**, 5, 399-400	16.2	2
22	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*,	36.3	2
21	Is Heart Failure Inherited?: Beyond the Cardiomyopathies, Genetics Do Matter. *JAMA Cardiology*, **2018**, 3, 710-711	16.2	1
20	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	4.5	1
19	Plasma vitamin K levels are associated with coronary calcification in older adults.. *FASEB Journal*, **2006**, 20, A134	0.9	1
18	Genetic Contribution to Common Heart Failure-Not So Rare?. *JAMA Cardiology*, **2021**, 6, 387	16.2	1
17	Genome-wide transcriptome study using deep RNA sequencing for myocardial infarction and coronary artery calcification. *BMC Medical Genomics*, **2021**, 14, 45	3.7	1
16	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	8.5	1
15	A MUC5B gene polymorphism, rs35705950-T, confers protective effects in COVID-19 infection		1

14	Matrix Gla Protein Levels Are Associated With Arterial Stiffness and Incident Heart Failure With Preserved Ejection Fraction. *Arteriosclerosis, Thrombosis, and Vascular Biology*, **2021**, ATVBAHA121316664	9.4	0
13	Multiethnic Genome-Wide Association Study of Subclinical Atherosclerosis in Individuals With Type 2 Diabetes. *Circulation Genomic and Precision Medicine*, **2021**, 14, e003258	5.2	0
12	A Missense Variant in the IL-6 Receptor and Protection From Peripheral Artery Disease. *Circulation Research*, **2021**, 129, 968-970	15.7	0
11	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	6	0
10	A multi-population phenome-wide association study of genetically-predicted height in the Million Veteran Program. *PLoS Genetics*, **2022**, 18, e1010193	6	0
9	Reply:. *Hepatology*, **2010**, 52, 1519-1519	11.2	
8	Multi-Trait Genome-Wide Association Study of Atherosclerosis Detects Novel Pleiotropic Loci.. *Frontiers in Genetics*, **2021**, 12, 787545	4.5	
7	Minority-centric meta-analyses of blood lipid levels identify novel loci in the Population Architecture using Genomics and Epidemiology (PAGE) study **2020**, 16, e1008684		
6	Minority-centric meta-analyses of blood lipid levels identify novel loci in the Population Architecture using Genomics and Epidemiology (PAGE) study **2020**, 16, e1008684		
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2	Minority-centric meta-analyses of blood lipid levels identify novel loci in the Population Architecture using Genomics and Epidemiology (PAGE) study **2020**, 16, e1008684		
1	Genome-wide and phenome-wide analysis of ideal cardiovascular health in the VA Million Veteran Program. *PLoS ONE*, **2022**, 17, e0267900	3.7	