List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3754479/publications.pdf Version: 2024-02-01

| 151 | 10,149 citations | 81900 39 h-index | 43889 91 g-index |
|-----------------|-----------------------|------------------------|-------------------------|
| papers | citations | n-mdex | g-mdex |
| 172 all docs | 172 docs citations | 172 times ranked | 16682 citing authors |

YAN V SUN

| # | Article | IF | CITATIONS |
|----|--|------|-----------|
| 1 | Genetic variants in novel pathways influence blood pressure and cardiovascular disease risk. Nature, 2011, 478, 103-109. | 27.8 | 1,855 |
| 2 | Genetic analysis of over 1 million people identifies 535 new loci associated with blood pressure traits. Nature Genetics, 2018, 50, 1412-1425. | 21.4 | 924 |
| 3 | Genetics of blood lipids among ~300,000 multi-ethnic participants of the Million Veteran Program. Nature Genetics, 2018, 50, 1514-1523. | 21.4 | 497 |
| 4 | 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. | 21.4 | 445 |
| 5 | The power of genetic diversity in genome-wide association studies of lipids. Nature, 2021, 600, 675-679. | 27.8 | 353 |
| 6 | Trans-ethnic association study of blood pressure determinants in over 750,000 individuals. Nature Genetics, 2019, 51, 51-62. | 21.4 | 328 |
| 7 | Meta-analysis of Correlated Traits via Summary Statistics from GWASs with an Application in Hypertension. American Journal of Human Genetics, 2015, 96, 21-36. | 6.2 | 321 |
| 8 | Integrative Analysis of Multi-omics Data for Discovery and Functional Studies of Complex Human Diseases. Advances in Genetics, 2016, 93, 147-190. | 1.8 | 306 |
| 9 | Meta-analyses identify 13 loci associated with age at menopause and highlight DNA repair and immune pathways. Nature Genetics, 2012, 44, 260-268. | 21.4 | 303 |
| 10 | Genome-Wide Association Study for Coronary Artery Calcification With Follow-Up in Myocardial Infarction. Circulation, 2011, 124, 2855-2864. | 1.6 | 269 |
| 11 | A Bivariate Genome-Wide Approach to Metabolic Syndrome. Diabetes, 2011, 60, 1329-1339. | 0.6 | 226 |
| 12 | Meta-Analysis of Genome-Wide Association Studies in African Americans Provides Insights into the Genetic Architecture of Type 2 Diabetes. PLoS Genetics, 2014, 10, e1004517. | 3.5 | 191 |
| 13 | Genome-wide Association Analysis of Blood-Pressure Traits in African-Ancestry Individuals Reveals Common Associated Genes in African and Non-African Populations. American Journal of Human Genetics, 2013, 93, 545-554. | 6.2 | 189 |
| 14 | Genome-wide association study of peripheral artery disease in the Million Veteran Program. Nature Medicine, 2019, 25, 1274-1279. | 30.7 | 177 |
| 15 | Harmonizing Genetic Ancestry and Self-identified Race/Ethnicity in Genome-wide Association Studies. American Journal of Human Genetics, 2019, 105, 763-772. | 6.2 | 169 |
| 16 | Epigenomic association analysis identifies smoking-related DNA methylation sites in African Americans. Human Genetics, 2013, 132, 1027-1037. | 3.8 | 153 |
| 17 | Genome-wide association analysis of venous thromboembolism identifies new risk loci and genetic overlap with arterial vascular disease. Nature Genetics, 2019, 51, 1574-1579. | 21.4 | 152 |
| 18 | A Genome-Wide Association Study of Depressive Symptoms. Biological Psychiatry, 2013, 73, 667-678. | 1.3 | 149 |

| # | Article | IF | CITATIONS |
|----|---|------|-----------|
| 19 | Actionable druggable genome-wide Mendelian randomization identifies repurposing opportunities for COVID-19. Nature Medicine, 2021, 27, 668-676. | 30.7 | 120 |
| 20 | Genome-wide analysis identifies novel susceptibility loci for myocardial infarction. European Heart Journal, 2021, 42, 919-933. | 2.2 | 113 |
| 21 | Pleiotropic genes for metabolic syndrome and inflammation. Molecular Genetics and Metabolism, 2014, 112, 317-338. | 1.1 | 107 |
| 22 | Brief review of regressionâ€based and machine learning methods in genetic epidemiology: the Genetic Analysis Workshop 17 experience. Genetic Epidemiology, 2011, 35, S5-11. | 1.3 | 93 |
| 23 | Discovery of rare variants associated with blood pressure regulation through meta-analysis of 1.3 million individuals. Nature Genetics, 2020, 52, 1314-1332. | 21.4 | 91 |
| 24 | Single-trait and multi-trait genome-wide association analyses identify novel loci for blood pressure in African-ancestry populations. PLoS Genetics, 2017, 13, e1006728. | 3.5 | 88 |
| 25 | Genetic Architecture of Abdominal Aortic Aneurysm in the Million Veteran Program. Circulation, 2020, 142, 1633-1646. | 1.6 | 78 |
| 26 | Association of Interleukin 6 Receptor Variant With Cardiovascular Disease Effects of Interleukin 6 Receptor Blocking Therapy. JAMA Cardiology, 2018, 3, 849. | 6.1 | 75 |
| 27 | Perceived Racial Discrimination and DNA Methylation Among African American Women in the InterGEN Study. Biological Research for Nursing, 2018, 20, 145-152. | 1.9 | 72 |
| 28 | The Mental Stress Ischemia Prognosis Study: Objectives, Study Design, and Prevalence of Inducible Ischemia. Psychosomatic Medicine, 2017, 79, 311-317. | 2.0 | 71 |
| 29 | 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. | 21.4 | 68 |
| 30 | Association Between High-Density Lipoprotein Cholesterol Levels and Adverse Cardiovascular Outcomes in High-risk Populations. JAMA Cardiology, 2022, 7, 672. | 6.1 | 66 |
| 31 | Hemodynamic, catecholamine, vasomotor and vascular responses: Determinants of myocardial ischemia during mental stress. International Journal of Cardiology, 2017, 243, 47-53. | 1.7 | 64 |
| 32 | Application of machine learning algorithms to predict coronary artery calcification with a sibshipâ€based design. Genetic Epidemiology, 2008, 32, 350-360. | 1.3 | 59 |
| 33 | Telomere Shortening, Regenerative Capacity, and Cardiovascular Outcomes. Circulation Research, 2017, 120, 1130-1138. | 4.5 | 59 |
| 34 | Comparison of the DNA methylation profiles of human peripheral blood cells and transformed B-lymphocytes. Human Genetics, 2010, 127, 651-658. | 3.8 | 53 |
| 35 | Identification of HIV infection-related DNA methylation sites and advanced epigenetic aging in HIV-positive, treatment-naive U.S. veterans. Aids, 2017, 31, 571-575. | 2.2 | 49 |
| 36 | Dystonia genes and their biological pathways. Neurobiology of Disease, 2019, 129, 159-168. | 4.4 | 49 |

| # | Article | IF | CITATIONS |
|----|--|------------------|----------------|
| 37 | Genetic Variation in NCAM1 Contributes to Left Ventricular Wall Thickness in Hypertensive Families. Circulation Research, 2011, 108, 279-283. | 4.5 | 47 |
| 38 | Inhibition of Corneal Inflammation by the Resolvin E1. , 2015, 56, 2728. | | 42 |
| 39 | Inflammatory response to mental stress and mental stress induced myocardial ischemia. Brain, Behavior, and Immunity, 2018, 68, 90-97. | 4.1 | 41 |
| 40 | Integration of biological networks and pathways with genetic association studies. Human Genetics, 2012, 131, 1677-1686. | 3.8 | 40 |
| 41 | Complexity in the genetic architecture of leukoaraiosis in hypertensive sibships from the GENOA Study. BMC Medical Genomics, 2009, 2, 16. | 1.5 | 39 |
| 42 | Novel DNA methylation sites associated with cigarette smoking among African Americans. Epigenetics, 2019, 14, 383-391. | 2.7 | 38 |
| 43 | The Intergenerational Impact of Genetic and Psychological Factors on Blood Pressure (InterGEN) Study. Biological Research for Nursing, 2016, 18, 521-530. | 1.9 | 37 |
| 44 | Gene-Specific DNA Methylation Association with Serum Levels of C-Reactive Protein in African Americans. PLoS ONE, 2013, 8, e73480. | 2.5 | 36 |
| 45 | A scan statistic for identifying chromosomal patterns of SNP association. Genetic Epidemiology, 2006, 30, 627-635. | 1.3 | 34 |
| 46 | A Genome-wide study of blood pressure in African Americans accounting for gene-smoking interaction. Scientific Reports, 2016, 6, 18812. | 3.3 | 34 |
| 47 | The Intergenerational Impact of Genetic and Psychological Factors on Blood Pressure Study (InterGEN). Nursing Research, 2016, 65, 331-338. | 1.7 | 33 |
| 48 | Identification, Heritability, and Relation With Gene Expression of Novel DNA Methylation Loci for Blood Pressure. Hypertension, 2020, 76, 195-205. | 2.7 | 33 |
| 49 | Serum Metabolomics and Incidence of Atrial Fibrillation (from the Atherosclerosis Risk in) Tj ETQq1 1 0.784314 | rgBT /Ove 1.6 | rlock 10 Tf 50 |
| 50 | Imputing missing genotypic data of single-nucleotide polymorphisms using neural networks. European Journal of Human Genetics, 2008, 16, 487-495. | 2.8 | 31 |
| 51 | Multigenic Modeling of Complex Disease by Random Forests. Advances in Genetics, 2010, 72, 73-99. | 1.8 | 31 |
| 52 | Association of <i>APOL1</i> Risk Alleles With Cardiovascular Disease in Blacks in the Million Veteran Program. Circulation, 2019, 140, 1031-1040. | 1.6 | 31 |
| 53 | <i>APOL1</i> 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. | 5.1 | 31 |
| 54 | X chromosome-wide analysis identifies DNA methylation sites influenced by cigarette smoking. Clinical Epigenetics, 2016, 8, 20. | 4.1 | 30 |

| # | Article | IF | CITATIONS |
|----|--|-----|-----------|
| 55 | Association of Obesity with DNA Methylation Age Acceleration in African American Mothers from the InterGEN Study. International Journal of Molecular Sciences, 2019, 20, 4273. | 4.1 | 28 |
| 56 | Classification of rheumatoid arthritis status with candidate gene and genome-wide single-nucleotide polymorphisms using random forests. BMC Proceedings, 2007, 1, S62. | 1.6 | 27 |
| 57 | Association Between High-Sensitivity Cardiac Troponin Levels and Myocardial Ischemia During Mental Stress and Conventional Stress. JACC: Cardiovascular Imaging, 2018, 11, 603-611. | 5.3 | 27 |
| 58 | Epigenetic Age Acceleration and Cognitive Decline: A Twin Study. Journals of Gerontology - Series A Biological Sciences and Medical Sciences, 2021, 76, 1854-1863. | 3.6 | 27 |
| 59 | Peripheral Vasoconstriction During Mental Stress and Adverse Cardiovascular Outcomes in Patients With Coronary Artery Disease. Circulation Research, 2019, 125, 874-883. | 4.5 | 24 |
| 60 | Considerations for Cardiovascular Genetic and Genomic Research With Marginalized Racial and Ethnic Groups and Indigenous Peoples: A Scientific Statement From the American Heart Association. Circulation Genomic and Precision Medicine, 2021, 14, e000084. | 3.6 | 24 |
| 61 | Identification of epistatic effects using a protein-protein interaction database. Human Molecular Genetics, 2010, 19, 4345-4352. | 2.9 | 23 |
| 62 | Parenting stress and DNA methylation among African Americans in the InterGEN Study. Journal of Clinical and Translational Science, 2017, 1, 328-333. | 0.6 | 22 |
| 63 | Association of Chromosome 9p21 With Subsequent Coronary Heart Disease Events. Circulation Genomic and Precision Medicine, 2019, 12, e002471. | 3.6 | 22 |
| 64 | Sexual Differences in Genetic Predisposition of Coronary Artery Disease. Circulation Genomic and Precision Medicine, 2021, 14, e003147. | 3.6 | 22 |
| 65 | ChromoScan: a scan statistic application for identifying chromosomal regions in genomic studies. Bioinformatics, 2006, 22, 2945-2947. | 4.1 | 21 |
| 66 | The Influences of Genetic and Environmental Factors on Methylome-Wide Association Studies for Human Diseases. Current Genetic Medicine Reports, 2014, 2, 261-270. | 1.9 | 21 |
| 67 | Epigenetic Associations With Estimated Glomerular Filtration Rate Among Men With Human Immunodeficiency Virus Infection. Clinical Infectious Diseases, 2020, 70, 667-673. | 5.8 | 21 |
| 68 | Higher Activation of the Rostromedial Prefrontal Cortex During Mental Stress Predicts Major Cardiovascular Disease Events in Individuals With Coronary Artery Disease. Circulation, 2020, 142, 455-465. | 1.6 | 21 |
| 69 | Very High High-Density Lipoprotein Cholesterol Levels and Cardiovascular Mortality. American Journal of Cardiology, 2022, 167, 43-53. | 1.6 | 21 |
| 70 | The cis and trans effects of the risk variants of coronary artery disease in the Chr9p21 region. BMC Medical Genomics, 2015, 8, 21. | 1.5 | 20 |
| 71 | DNA Methylation Markers of Type 2 Diabetes Mellitus Among Male Veterans With or Without Human Immunodeficiency Virus Infection. Journal of Infectious Diseases, 2019, 219, 1959-1962. | 4.0 | 20 |
| 72 | An investigation of racial/ethnic and sex differences in the association between experiences of everyday discrimination and leukocyte telomere length among patients with coronary artery disease. Psychoneuroendocrinology, 2019, 106, 122-128. | 2.7 | 19 |

| # | Article | IF | CITATIONS |
|----|---|-----|-----------|
| 73 | Identification of genetic association of multiple rare variants using collapsing methods. Genetic Epidemiology, 2011, 35, S101-6. | 1.3 | 18 |
| 74 | Untargeted metabolomics reveals multiple metabolites influencing smoking-related DNA methylation. Epigenomics, 2018, 10, 379-393. | 2.1 | 18 |
| 75 | Untargeted high-resolution plasma metabolomic profiling predicts outcomes in patients with coronary artery disease. PLoS ONE, 2020, 15, e0237579. | 2.5 | 18 |
| 76 | Discovery and fine-mapping of height loci via high-density imputation of GWASs in individuals of African ancestry. American Journal of Human Genetics, 2021, 108, 564-582. | 6.2 | 18 |
| 77 | Gene-environment effects of SLC4A5 and skin color on blood pressure among African American women. Ethnicity and Disease, 2012, 22, 155-61. | 2.3 | 18 |
| 78 | Subsequent Event Risk in Individuals With Established Coronary Heart Disease. Circulation Genomic and Precision Medicine, 2019, 12, e002470. | 3.6 | 17 |
| 79 | GEM: scalable and flexible gene–environment interaction analysis in millions of samples. Bioinformatics, 2021, 37, 3514-3520. | 4.1 | 17 |
| 80 | Copy Number Variations Associated With Obesityâ€Related Traits in African Americans: A Joint Analysis Between GENOA and HyperGEN. Obesity, 2012, 20, 2431-2437. | 3.0 | 15 |
| 81 | 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, . | 3.6 | 15 |
| 82 | Association of Kidney Comorbidities and Acute Kidney Failure With Unfavorable Outcomes After COVID-19 in Individuals With the Sickle Cell Trait. JAMA Internal Medicine, 0, , . | 5.1 | 15 |
| 83 | Polymorphisms in the vitamin D receptor gene are associated with reduced rate of sputum culture conversion in multidrug-resistant tuberculosis patients in South Africa. PLoS ONE, 2017, 12, e0180916. | 2.5 | 14 |
| 84 | Association Between Change in Circulating Progenitor Cells During Exercise Stress and Risk of Adverse Cardiovascular Events in Patients With Coronary Artery Disease. JAMA Cardiology, 2020, 5, 147. | 6.1 | 14 |
| 85 | Mendelian Randomization Analysis of Hemostatic Factors and Their Contribution to Peripheral Artery Disease. Arteriosclerosis, Thrombosis, and Vascular Biology, 2020, 41, 380-386. | 2.4 | 14 |
| 86 | Both metal binding sites in the homodimer are required for metalloregulation by the CadC repressor. Molecular Microbiology, 2002, 44, 1323-1329. | 2.5 | 13 |
| 87 | GWAS of longitudinal trajectories at biobank scale. American Journal of Human Genetics, 2022, 109, 433-445. | 6.2 | 13 |
| 88 | Comparative genome-wide association studies of a depressive symptom phenotype in a repeated measures setting by race/ethnicity in the multi-ethnic study of atherosclerosis. BMC Genetics, 2015, 16, 118. | 2.7 | 12 |
| 89 | Hypertension in HIV-Infected Adults Compared with Similar but Uninfected Adults in China: Body Mass Index-Dependent Effects of Nadir CD4 Count. AIDS Research and Human Retroviruses, 2017, 33, 1117-1125. | 1.1 | 12 |
| 90 | A multi-population phenome-wide association study of genetically-predicted height in the Million Veteran Program. PLoS Genetics, 2022, 18, e1010193. | 3.5 | 12 |

| # | Article | IF | CITATIONS |
|-----|---|-----|-----------|
| 91 | Change in Left Ventricular Ejection Fraction With Coronary Artery Revascularization and Subsequent Risk for Adverse Cardiovascular Outcomes. Circulation: Cardiovascular Interventions, 2022, 15, 101161CIRCINTERVENTIONS121011284. | 3.9 | 11 |
| 92 | Identification of correlated genetic variants jointly associated with rheumatoid arthritis using ridge regression. BMC Proceedings, 2009, 3, S67. | 1.6 | 10 |
| 93 | Fast implementation of a scan statistic for identifying chromosomal patterns of genome wide association studies. Computational Statistics and Data Analysis, 2009, 53, 1794-1801. | 1.2 | 10 |
| 94 | Study designs and methods post genome-wide association studies. Human Genetics, 2012, 131, 1525-1531. | 3.8 | 10 |
| 95 | Epigenomic Indicators of Age in African Americans. Hereditary Genetics: Current Research, 2014, 03, . | 0.1 | 10 |
| 96 | Association Between Stress and Coping with DNA Methylation of Blood Pressure-Related Genes Among African American Women. Chronic Stress, 2019, 3, 247054701987908. | 3.4 | 9 |
| 97 | PCSK9 loss of function is protective against extra-coronary atherosclerotic cardiovascular disease in a large multi-ethnic cohort. PLoS ONE, 2020, 15, e0239752. | 2.5 | 9 |
| 98 | 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. | 2.3 | 9 |
| 99 | Systematic Heritability and Heritability Enrichment Analysis for Diabetes Complications in UK Biobank and ACCORD Studies. Diabetes, 2022, 71, 1137-1148. | 0.6 | 9 |
| 100 | A Common Copy Number Variation on Chromosome 6 Association With the Gene Expression Level of Endothelin 1 in Transformed B Lymphocytes From Three Racial Groups. Circulation: Cardiovascular Genetics, 2009, 2, 483-488. | 5.1 | 8 |
| 101 | Integration of rare expression outlier-associated variants improves polygenic risk prediction. American Journal of Human Genetics, 2022, 109, 1055-1064. | 6.2 | 8 |
| 102 | Myocardial Ischemia and Mobilization of Circulating Progenitor Cells. Journal of the American Heart Association, 2018, 7, e007504. | 3.7 | 7 |
| 103 | Genomics of Reproductive Traits and Cardiometabolic Disease Risk in African American Women. Nursing Research, 2019, 68, 135-144. | 1.7 | 7 |
| 104 | Induced pluripotent stem cells from subjects with Lesch-Nyhan disease. Scientific Reports, 2021, 11, 8523. | 3.3 | 7 |
| 105 | Metabolomic Profiling Demonstrates Postprandial Changes in Fatty Acids and Glycerophospholipids Are Associated with Fasting Inflammation in Guatemalan Adults. Journal of Nutrition, 2021, 151, 2564-2573. | 2.9 | 7 |
| 106 | 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. | 2.9 | 6 |
| 107 | A metabolomic study of cervical dystonia. Parkinsonism and Related Disorders, 2021, 82, 98-103. | 2.2 | 6 |
| 108 | Coronary Artery Disease Risk of Familial Hypercholesterolemia Genetic Variants Independent of Clinically Observed Longitudinal Cholesterol Exposure. Circulation Genomic and Precision Medicine, 2022, 15, CIRCGEN121003501. | 3.6 | 6 |

| # | Article | IF | CITATIONS |
|-----|--|-----|-----------|
| 109 | Gene-by-Psychosocial Factor Interactions Influence Diastolic Blood Pressure in European and African Ancestry Populations: Meta-Analysis of Four Cohort Studies. International Journal of Environmental Research and Public Health, 2017, 14, 1596. | 2.6 | 5 |
| 110 | A Multiâ€center Genomeâ€wide Association Study of Cervical Dystonia. Movement Disorders, 2021, 36, 2795-2801. | 3.9 | 5 |
| 111 | Epigenome-wide epidemiologic studies of human immunodeficiency virus infection, treatment, and disease progression. Clinical Epigenetics, 2022, 14, 8. | 4.1 | 5 |
| 112 | High blood pressure in pregnancy, DNA methylation, and later blood pressure in African American women enrolled in the InterGEN Study. Birth, 2020, 47, 290-298. | 2.2 | 4 |
| 113 | Evaluation of the Host Genetic Effects of Tuberculosis-Associated Variants Among Patients With Type 1 and Type 2 Diabetes Mellitus. Open Forum Infectious Diseases, 2020, 7, ofaa106. | 0.9 | 4 |
| 114 | Cardiovascular disease risk and pathophysiology in South Asians: can longitudinal multi-omics shed light?. Wellcome Open Research, 2020, 5, 255. | 1.8 | 4 |
| 115 | Relation of High-sensitivity Cardiac Troponin I Elevation With Exercise to Major Adverse Cardiovascular Events in Patients With Coronary Artery Disease. American Journal of Cardiology, 2020, 136, 1-8. | 1.6 | 4 |
| 116 | DNA Methylation of TXNIP Independently Associated with Inflammation and Diabetes Mellitus in Twins. Twin Research and Human Genetics, 2021, , 1-8. | 0.6 | 4 |
| 117 | Cardiovascular disease risk and pathophysiology in South Asians: can longitudinal multi-omics shed light?. Wellcome Open Research, 2020, 5, 255. | 1.8 | 4 |
| 118 | Obesityâ€associated metabolites in relation to type 2 diabetes risk: A prospective nested <scp>caseâ€control</scp> study of the <scp>CARRS</scp> cohort. Diabetes, Obesity and Metabolism, 2022, 24, 2008-2016. | 4.4 | 4 |
| 119 | Rapid Collection of Biospecimens by Automated Identification of Patients Eligible for Pharmacoepigenetic Studies. Journal of Personalized Medicine, 2013, 3, 263-274. | 2.5 | 3 |
| 120 | Association analysis of whole genome sequencing data accounting for longitudinal and family designs. BMC Proceedings, 2014, 8, S89. | 1.6 | 3 |
| 121 | DNA methylation changes in African American women with a history of preterm birth from the InterGEN study. BMC Genomic Data, 2021, 22, 30. | 1.7 | 3 |
| 122 | Interaction between genetics and smoking in determining risk of coronary artery diseases. Genetic Epidemiology, 2022, 46, 199-212. | 1.3 | 3 |
| 123 | Identification of genes associated with complex traits by testing the genetic dissimilarity between individuals. BMC Proceedings, 2011, 5, S120. | 1.6 | 2 |
| 124 | Depressive Symptoms and Blood Pressure in African American Women. Journal of Cardiovascular Nursing, 2022, 37, E89-E96. | 1.1 | 2 |
| 125 | Anticipation of Precision Diabetes and Promise of Integrative Multi-Omics. Endocrinology and Metabolism Clinics of North America, 2021, 50, 559-574. | 3.2 | 2 |
| 126 | Abstract 02: Brain Regions Activation During Stress and Accelerated Biological Aging. Circulation, 2020, 141, . | 1.6 | 2 |

| # | Article | IF | CITATIONS |
|-----|--|-----|-----------|
| 127 | Neural correlates of stress and leucocyte telomere length in patients with coronary artery disease. Journal of Psychosomatic Research, 2022, 155, 110760. | 2.6 | 2 |
| 128 | Early Life Trauma Is Associated With Increased Microvolt Tâ€Wave Alternans During Mental Stress Challenge: A Substudy of Mental Stress Ischemia: Prognosis and Genetic Influences. Journal of the American Heart Association, 2022, 11, e021582. | 3.7 | 2 |
| 129 | DNA Methylation, Preterm Birth and Blood Pressure in African American Children: The DPREG Study. Journal of Immigrant and Minority Health, 2022, 24, 334-341. | 1.6 | 1 |
| 130 | Abstract MP10: A Novel Genetic Locus Influences Microvascular Reactivity To Acute Psychological Stress. Circulation, 2021, 143, . | 1.6 | 1 |
| 131 | Associations of biogeographic ancestry with hypertension traits. Journal of Hypertension, 2021, 39, 633-642. | 0.5 | 1 |
| 132 | Abstract 126: Genome Wide Association Study in the Million Veteran Program Identifies a Novel Role for Thrombosis in the Pathogenesis of Peripheral Artery Disease. Arteriosclerosis, Thrombosis, and Vascular Biology, 2018, 38, . | 2.4 | 1 |
| 133 | Rare variant association study of veteran twin whole-genomes links severe depression with a nonsynonymous change in the neuronal gene <i>BHLHE22</i> . World Journal of Biological Psychiatry, 2022, 23, 295-306. | 2.6 | 1 |
| 134 | Hemoglobin A1c (A1c) Genetics Contributes to A1c/Glucose Mismatches in the Multiethnic VA Million Veteran Program (MVP). Diabetes, 2018, 67, . | 0.6 | 1 |
| 135 | Associations Between DNA Methylation Age Acceleration, Depressive Symptoms, and Cardiometabolic Traits in African American Mothers From the InterGEN Study. Epigenetics Insights, 2022, 15, 251686572211097. | 2.0 | 1 |
| 136 | A complex system of chemokines may hold the key to optimal CD4+ T-cell recovery after antiretroviral therapy. EBioMedicine, 2020, 62, 103113. | 6.1 | 0 |
| 137 | Metabolomic Profiling Demonstrates Postprandial Changes in Saturated Fatty Acids and Glycerophospholipids Are Associated With Fasting Inflammation. Current Developments in Nutrition, 2021, 5, 1106. | 0.3 | 0 |
| 138 | 244-OR: Toward Improved Identification of Adult-Onset Type 1 Diabetes (T1D): Clinical Characteristics Associated with T1D Polygenic Risk Scores in the Million Veteran Program (MVP). Diabetes, 2021, 70, . | 0.6 | 0 |
| 139 | Abstract P327: Stress Overload and Methylation of Hypertension Related Genes. Circulation, 2018, 137, . | 1.6 | 0 |
| 140 | Abstract 050: Phenome Wide Association Study of IL6R Variants Identifies a Drug Target for Cardiovascular Disease and Inflammation. Circulation, 2018, 137, . | 1.6 | 0 |
| 141 | Abstract P134: Genome-wide Association Study of Heart Rate Response to Mental Stress. Circulation, 2018, 137, . | 1.6 | 0 |
| 142 | Abstract 024: Association of APOL1 Risk Alleles with Coronary Heart Disease in Million Veteran Program. Arteriosclerosis, Thrombosis, and Vascular Biology, 2018, 38, . | 2.4 | 0 |
| 143 | Abstract P336: Predictors of High Intensity Statin Initiation for Primary Prevention in Veterans With Familial Hypercholesterolemia Phenotype. Circulation, 2019, 139, . | 1.6 | 0 |
| 144 | 239-OR: Genome-Wide Association Study (GWAS) of Hypoglycemia in the Million Veteran Program (MVP). Diabetes, 2020, 69, 239-OR. | 0.6 | 0 |

| # | Article | IF | CITATIONS |
|-----|---|-----|-----------|
| 145 | Polygenic Risk Score Identifies Patients at Increased Risk for Abdominal Aortic Aneurysm and May Benefit from Ultrasound Screening. JVS Vascular Science, 2020, 1, 251-252. | 1.1 | 0 |
| 146 | Abstract P457: Age Dependence of Genetic Risk Scores in Relation to Coronary Artery Disease. Circulation, 2020, 141, . | 1.6 | 0 |
| 147 | Abstract P141: A Novel GWAS Locus Influences Microvascular Response to Acute Psychological Stress. Circulation, 2020, 141, . | 1.6 | 0 |
| 148 | Abstract P148: Sex Differences in the Genetic Predisposition of Coronary Artery Disease. Circulation, 2020, 141, . | 1.6 | 0 |
| 149 | Abstract 13601: Risk of Coronary Artery Disease Associated With Familial Hypercholesterolemia Genetic Variants is Independent of Historical Low-density Lipoprotein Cholesterol Exposure. Circulation, 2020, 142, . | 1.6 | 0 |
| 150 | Abstract 16761: Refining Individualized Risk Prediction for Apparent Treatment-Resistant Hypertension in a Large Multiethnic Biobank: The Million Veteran Program. Circulation, 2020, 142, . | 1.6 | 0 |
| 151 | Baseline Characteristics of Mitochondrial DNA and Mutations Associated With Short-Term Posttreatment CD4+T-Cell Recovery in Chinese People With HIV. Frontiers in Immunology, 2021, 12, 793375 | 4.8 | Ο |